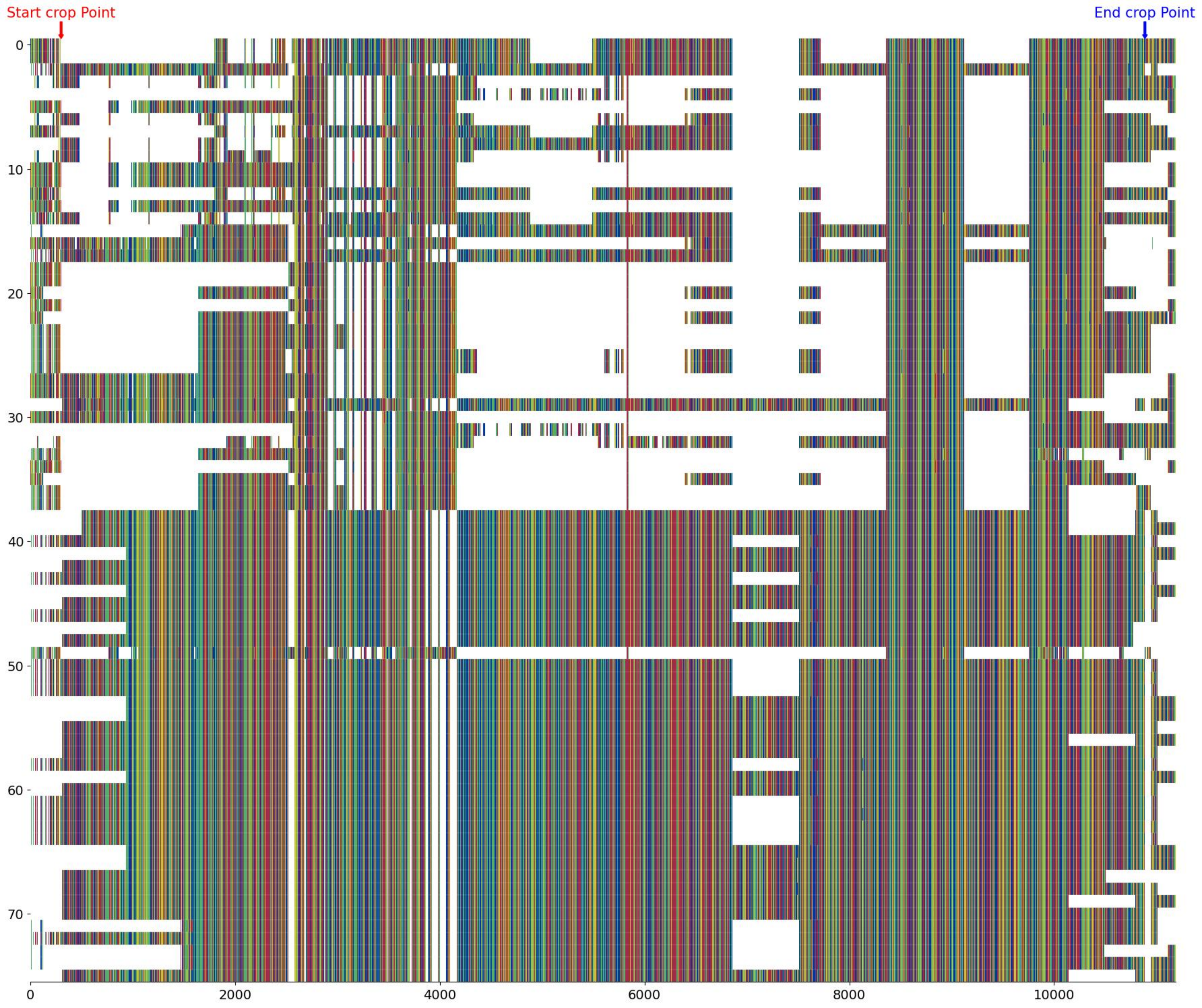
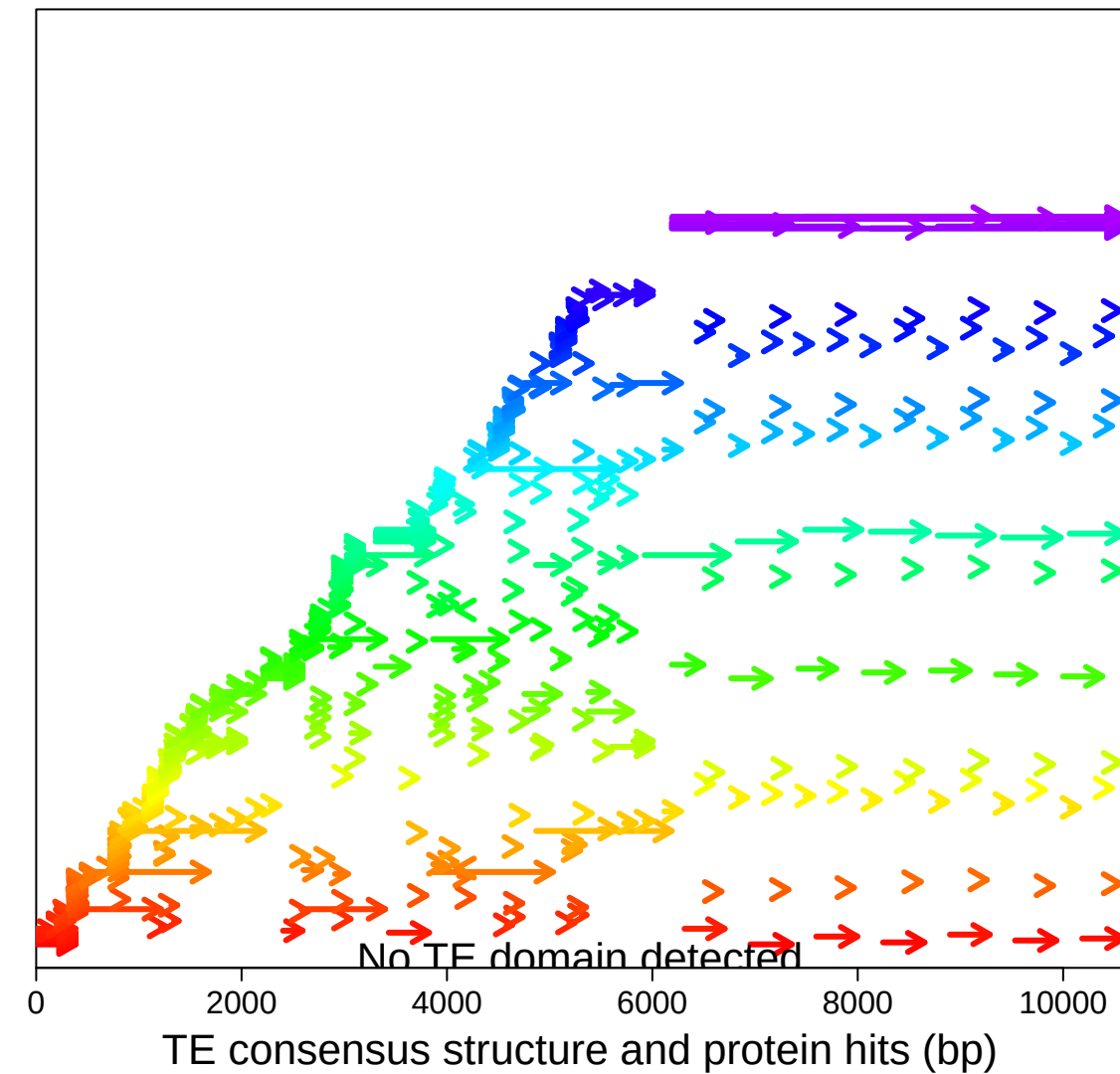
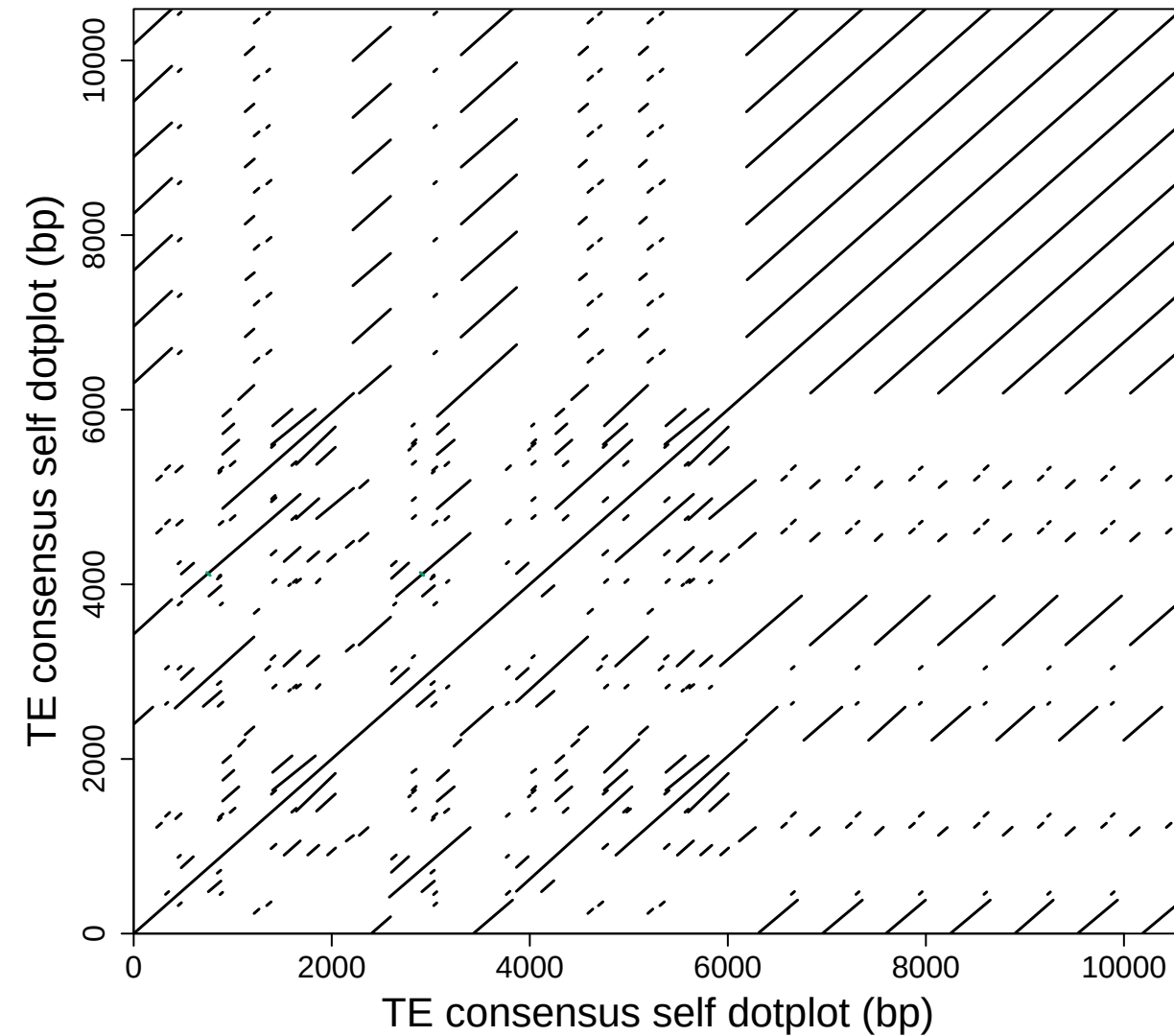
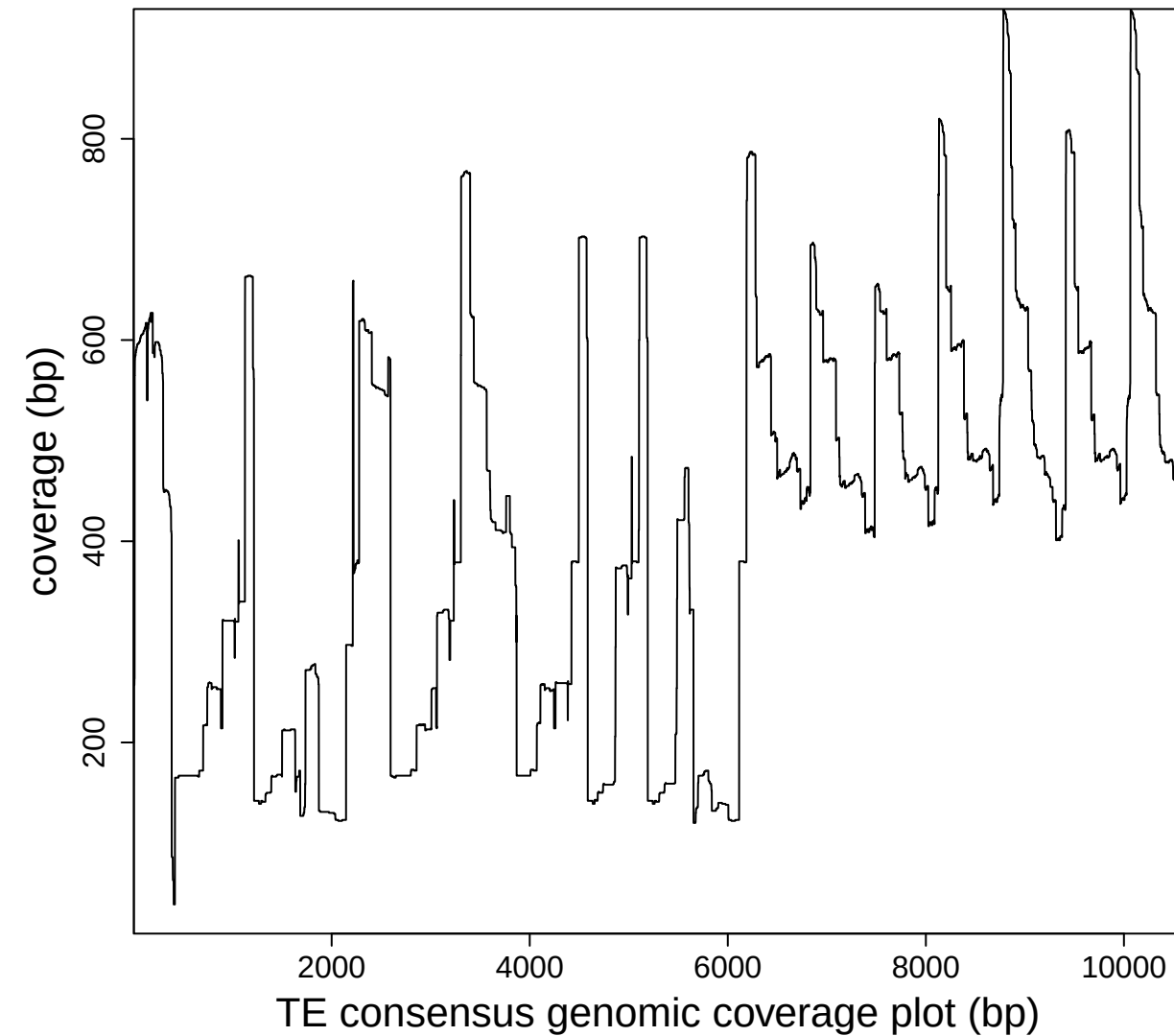
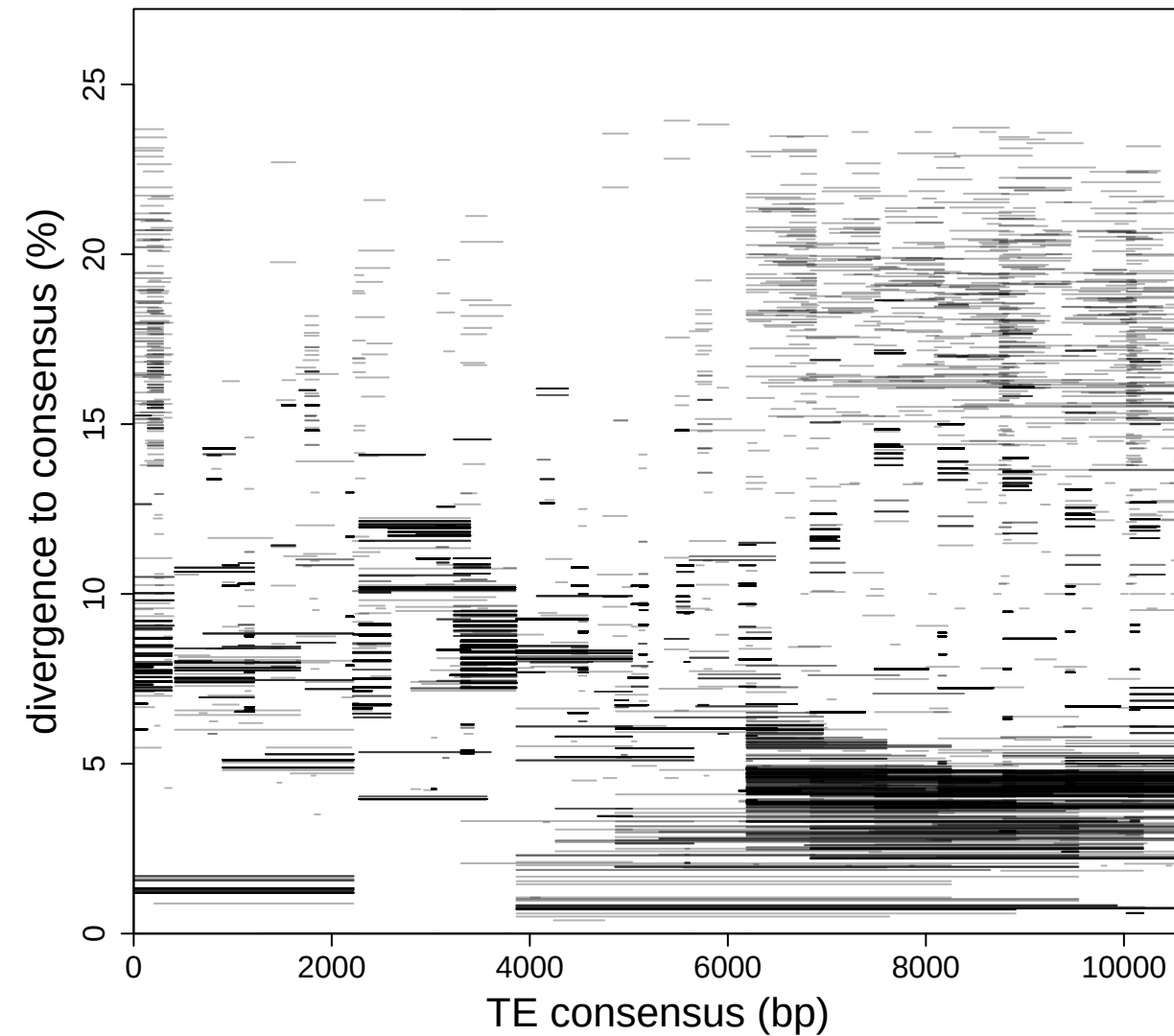






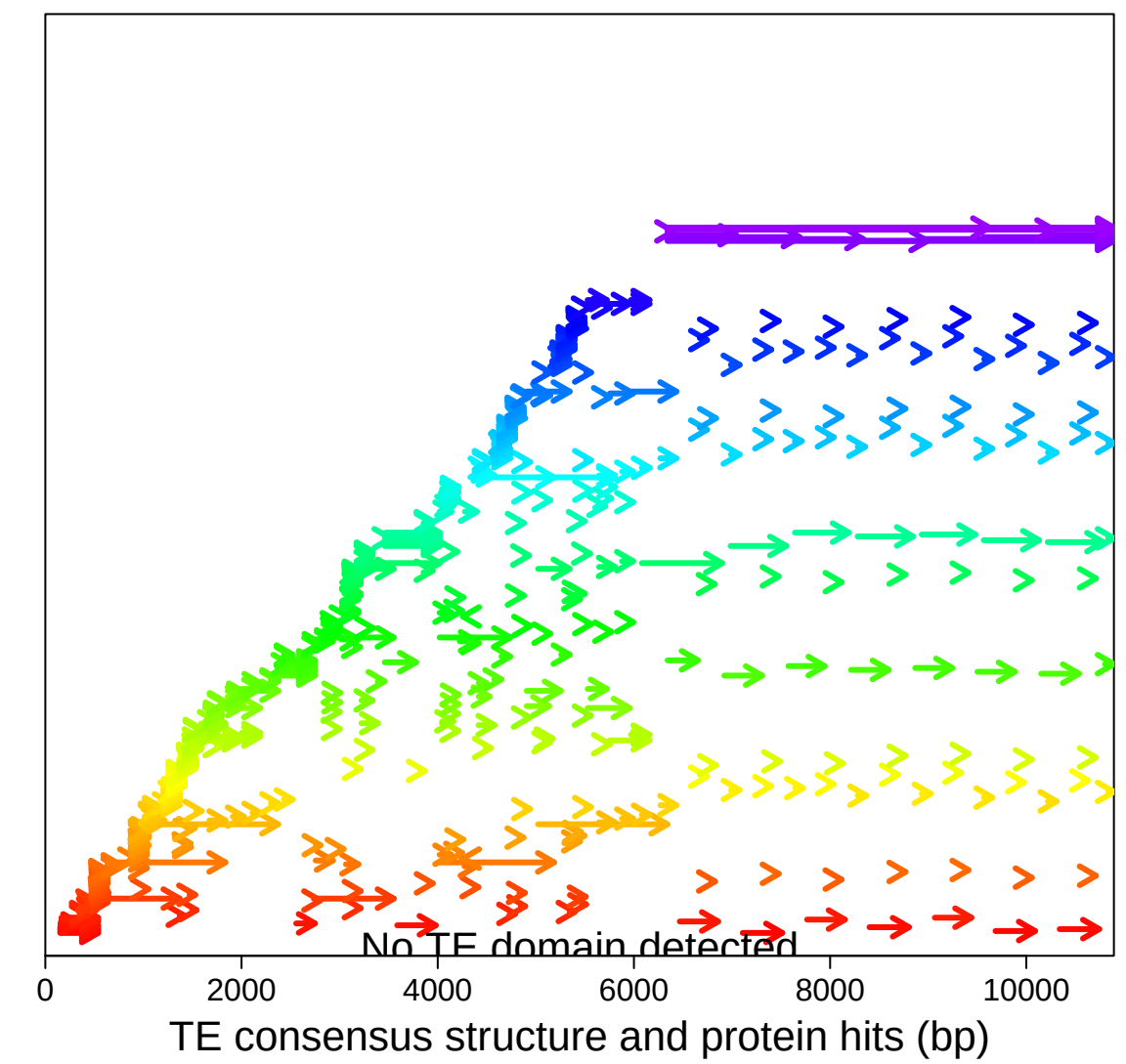
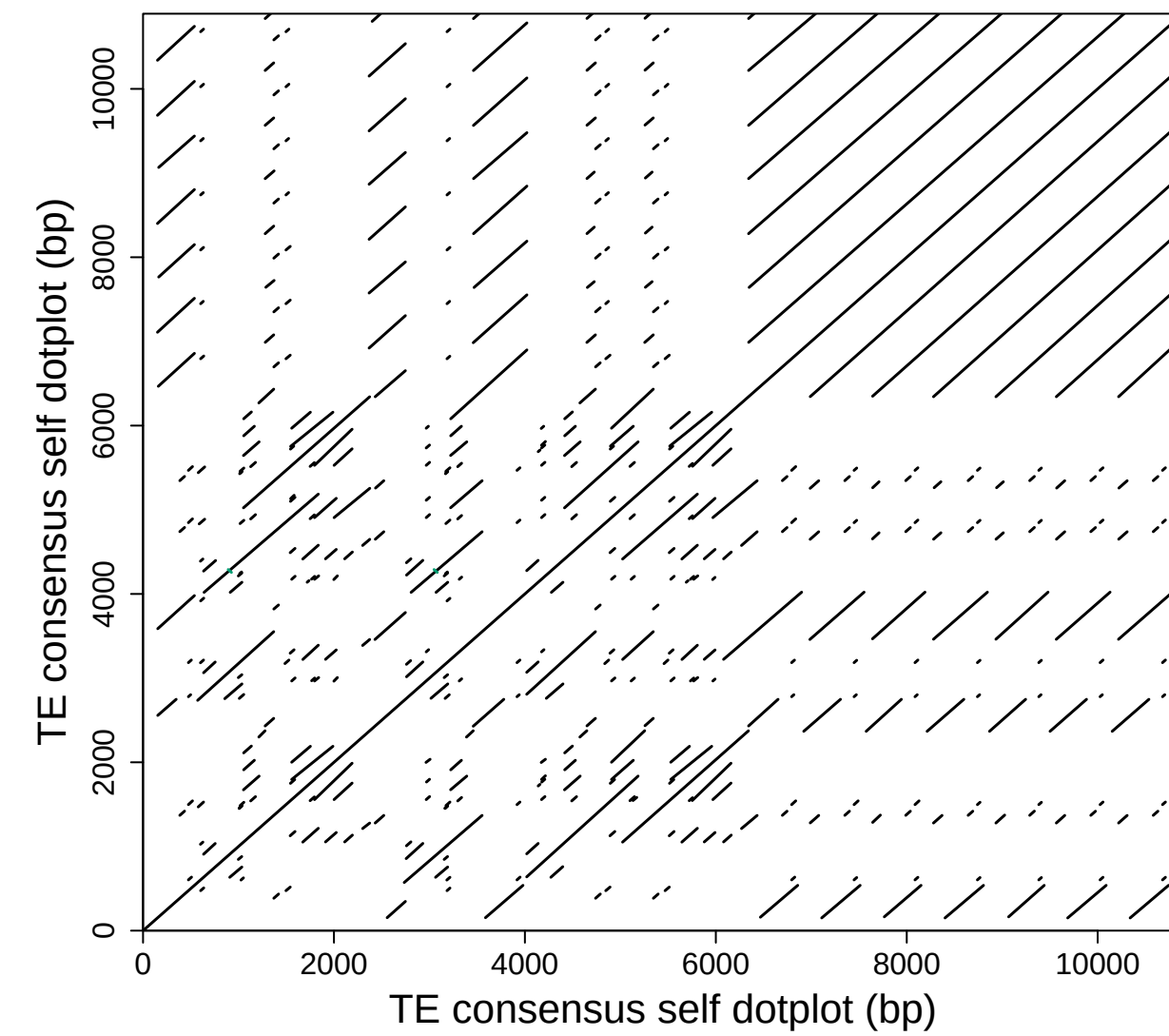
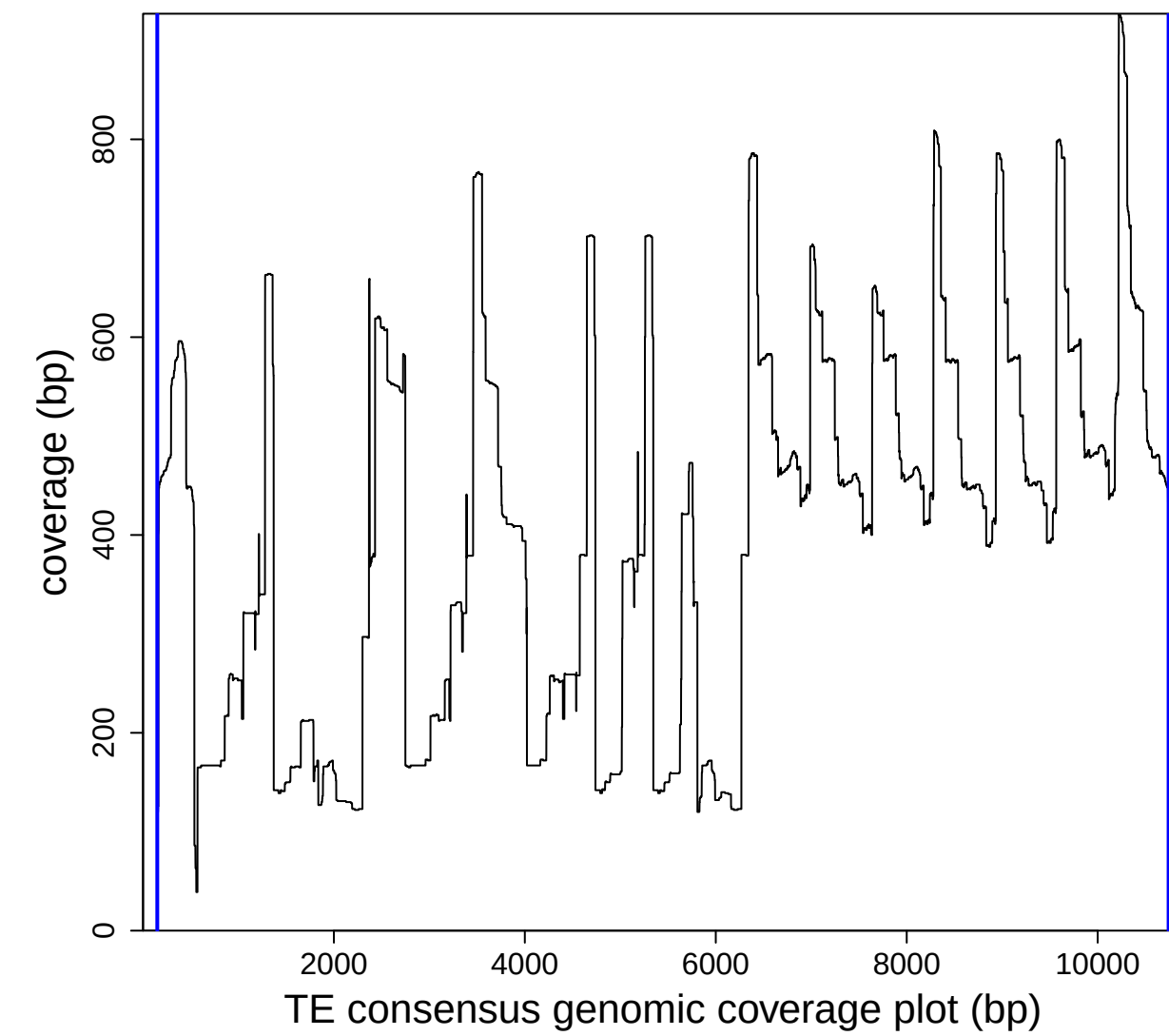
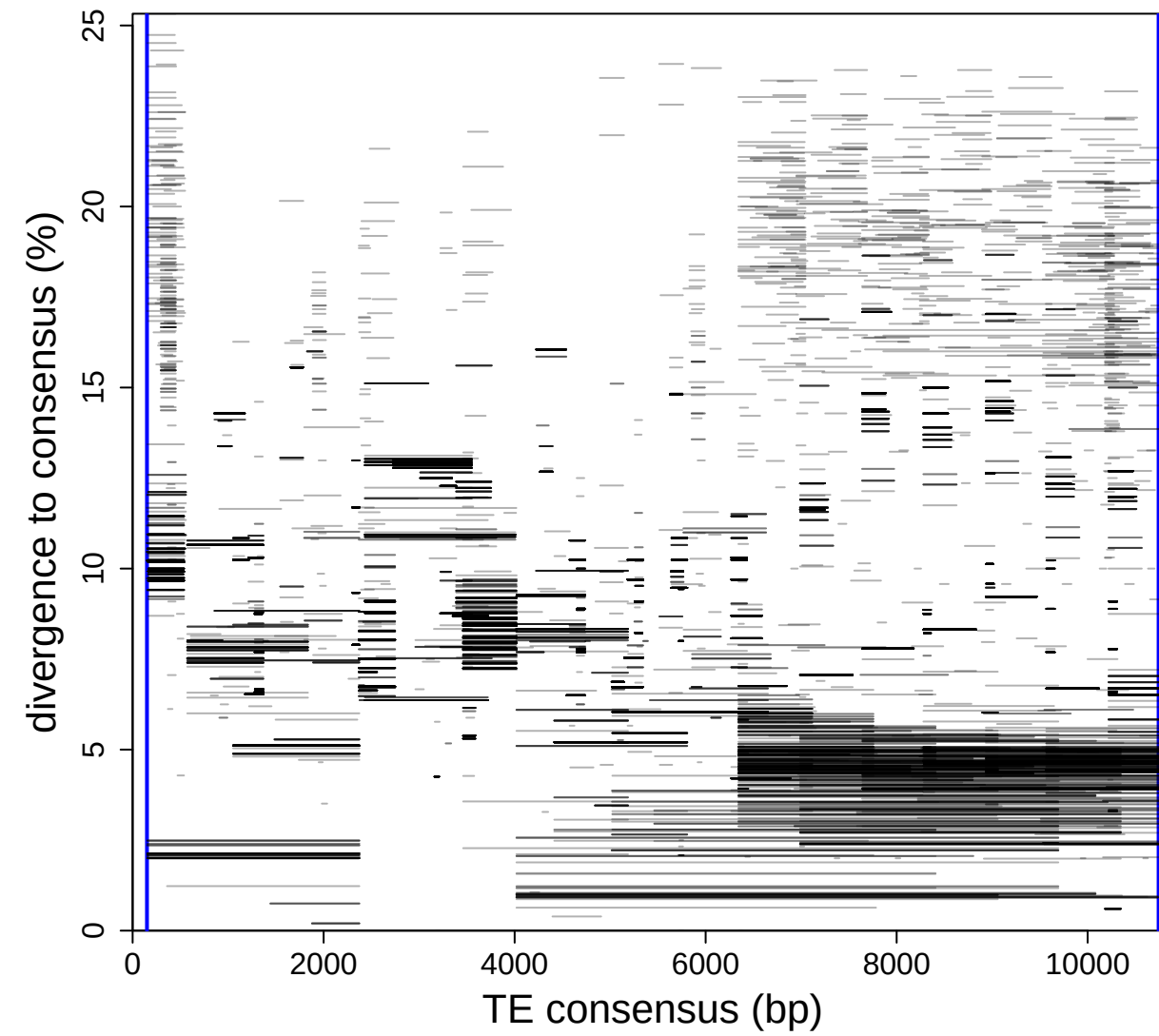
bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.fa_ce
size: 10589bp; fragments: 8864; full length: 0 (>=9530.1bp)

After TEtrimmer 10589 bp



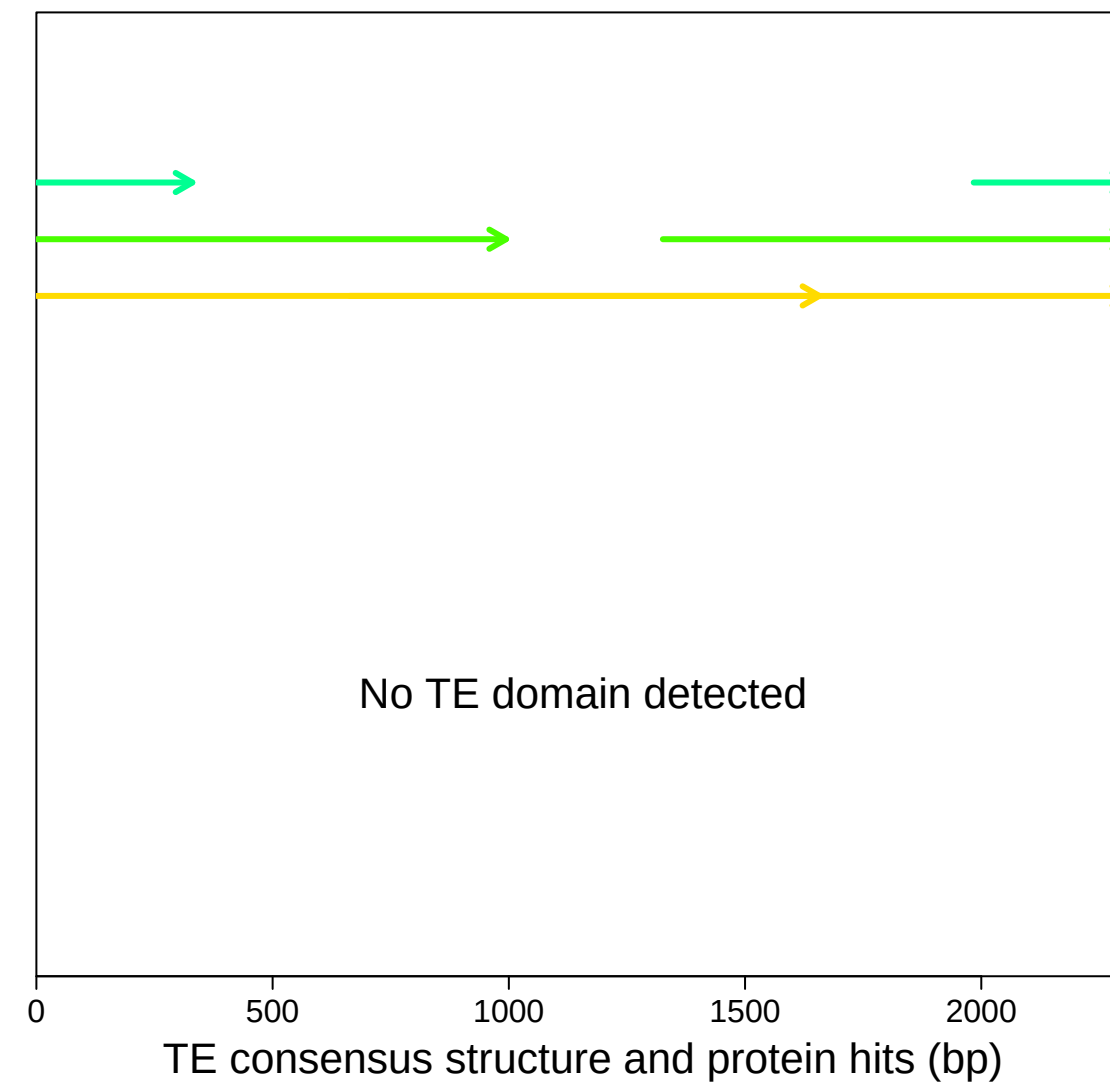
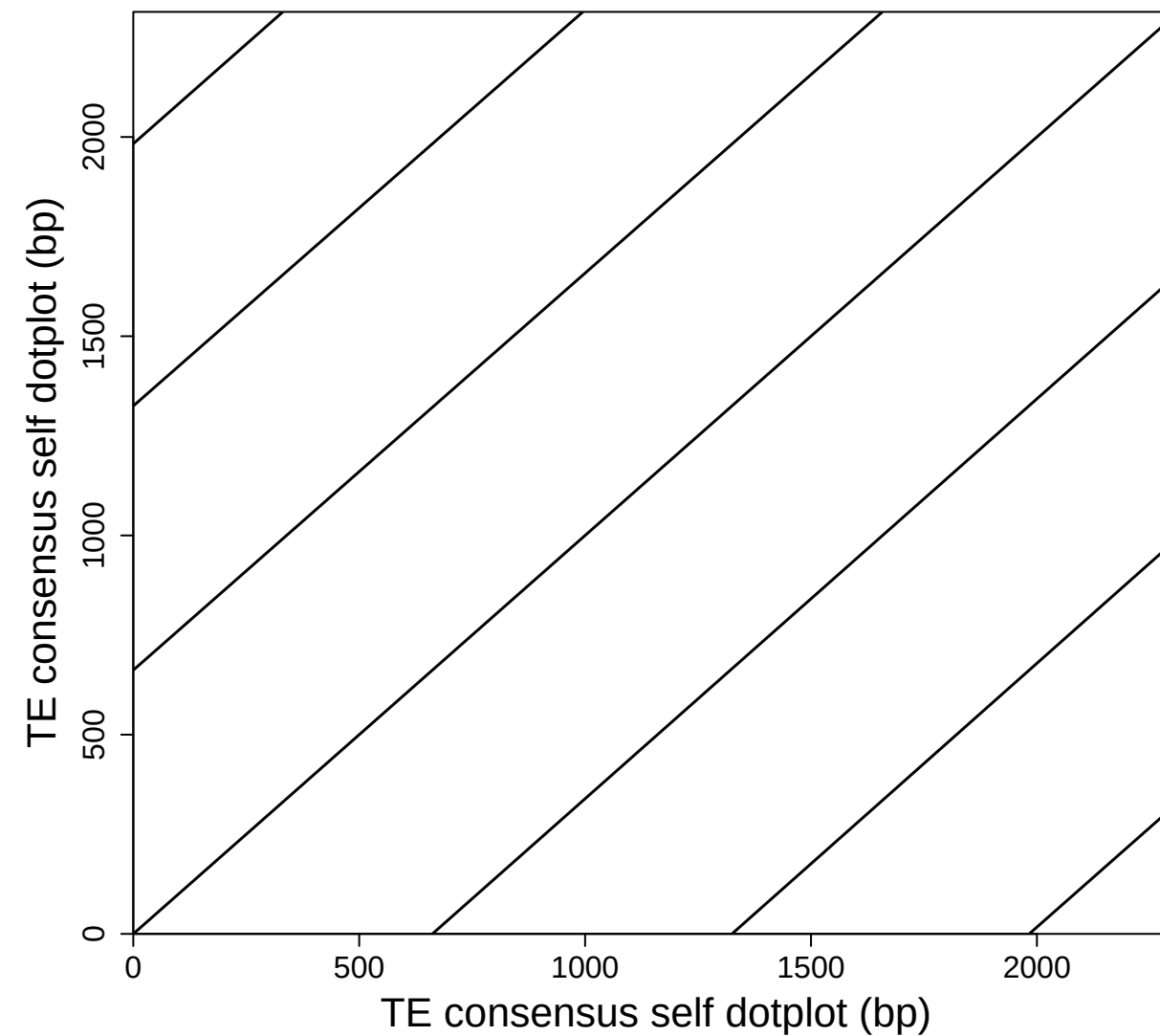
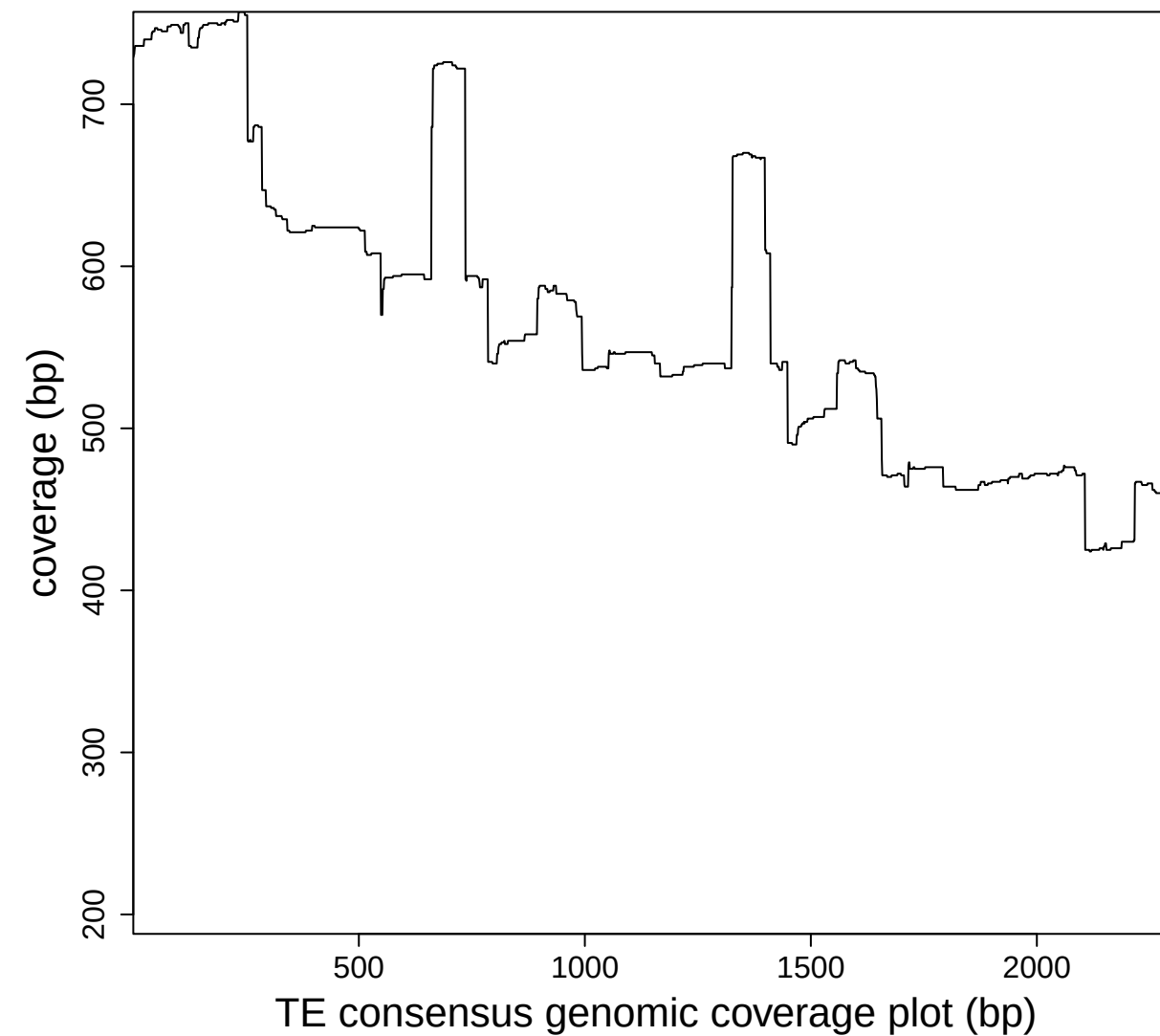
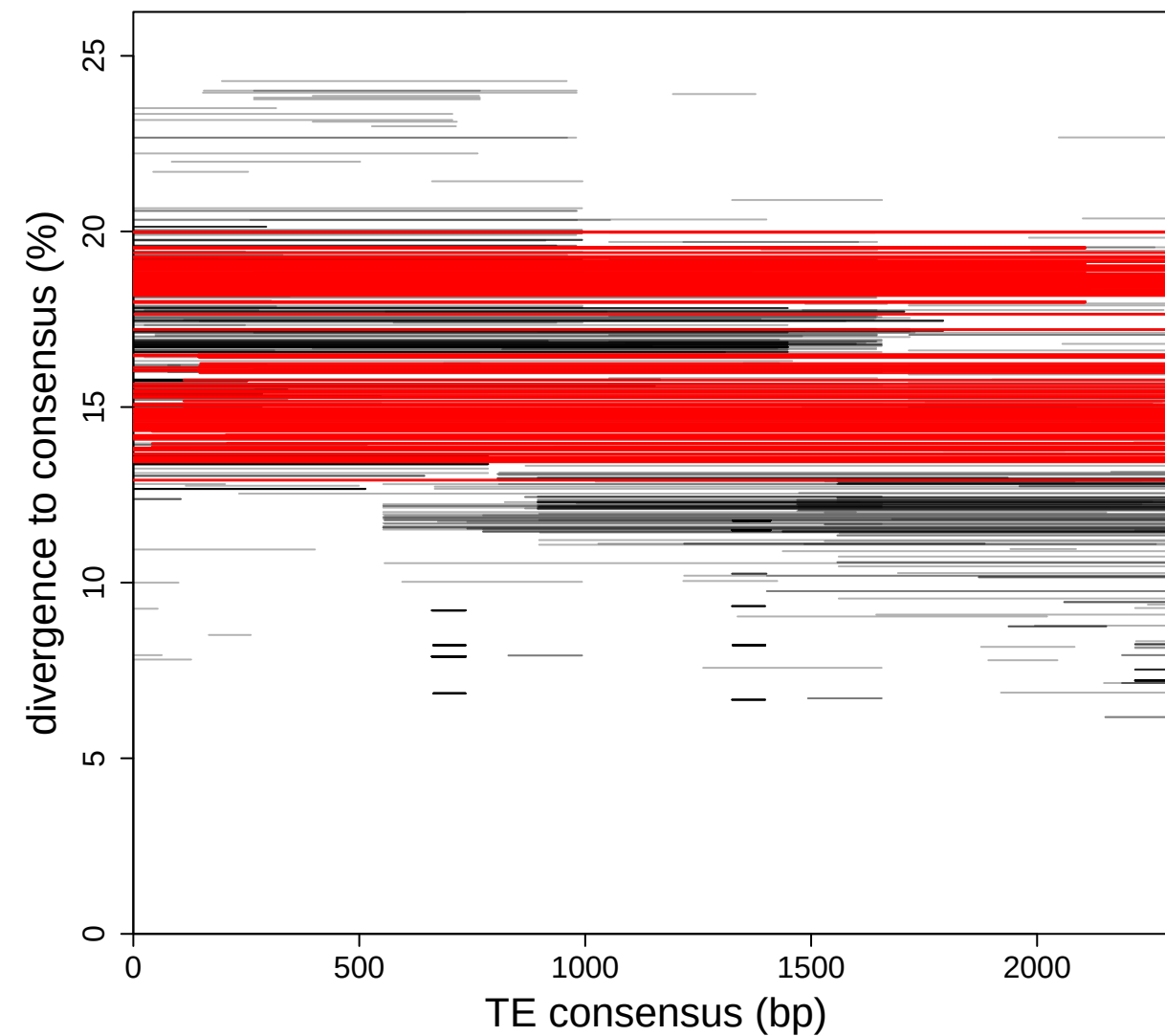
ily_510.fasta.b.bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.f
size: 10893bp; fragments: 8916; full length: 0 (>=9803.7bp)

After TEtrimmer Extended plot Blue lines are boundaries

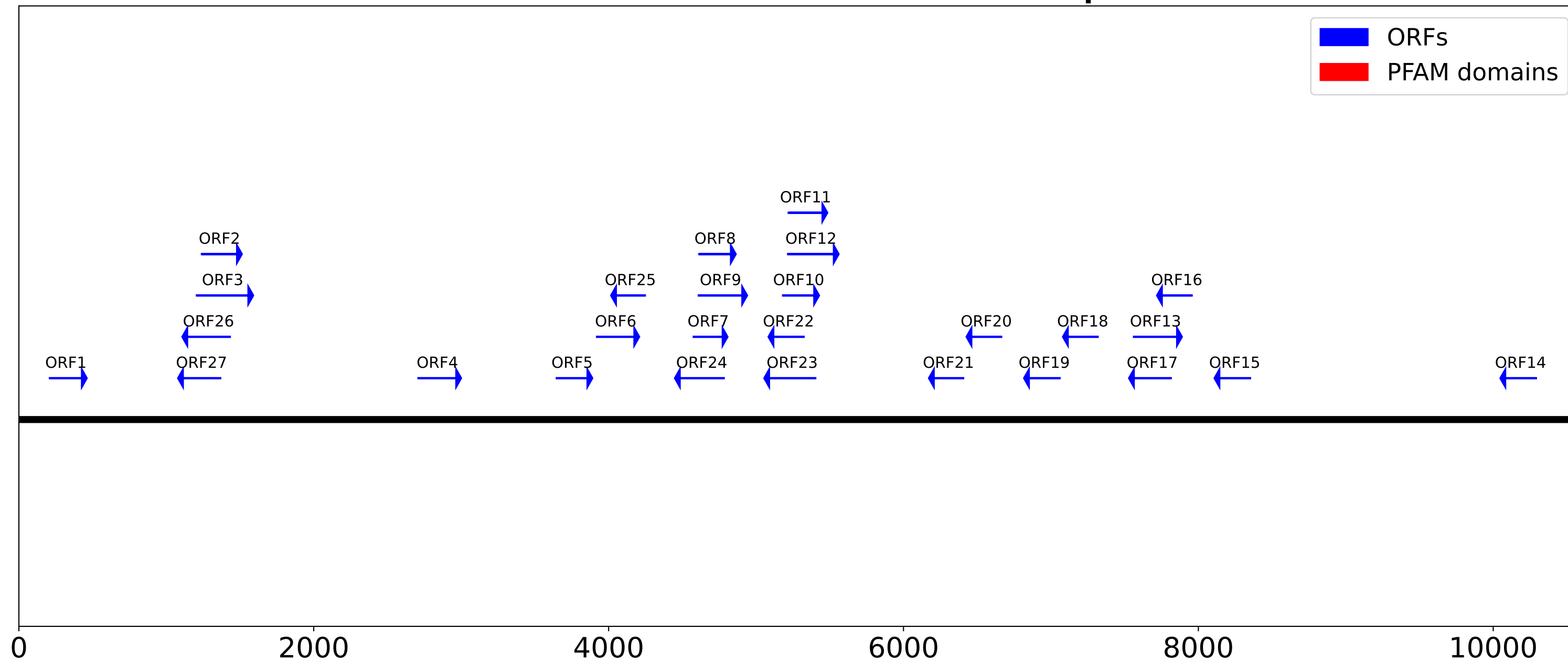


TE: rnd_1_family_510
size: 2314bp; fragments: 1339; full length: 290 (≥ 2082.6 bp)

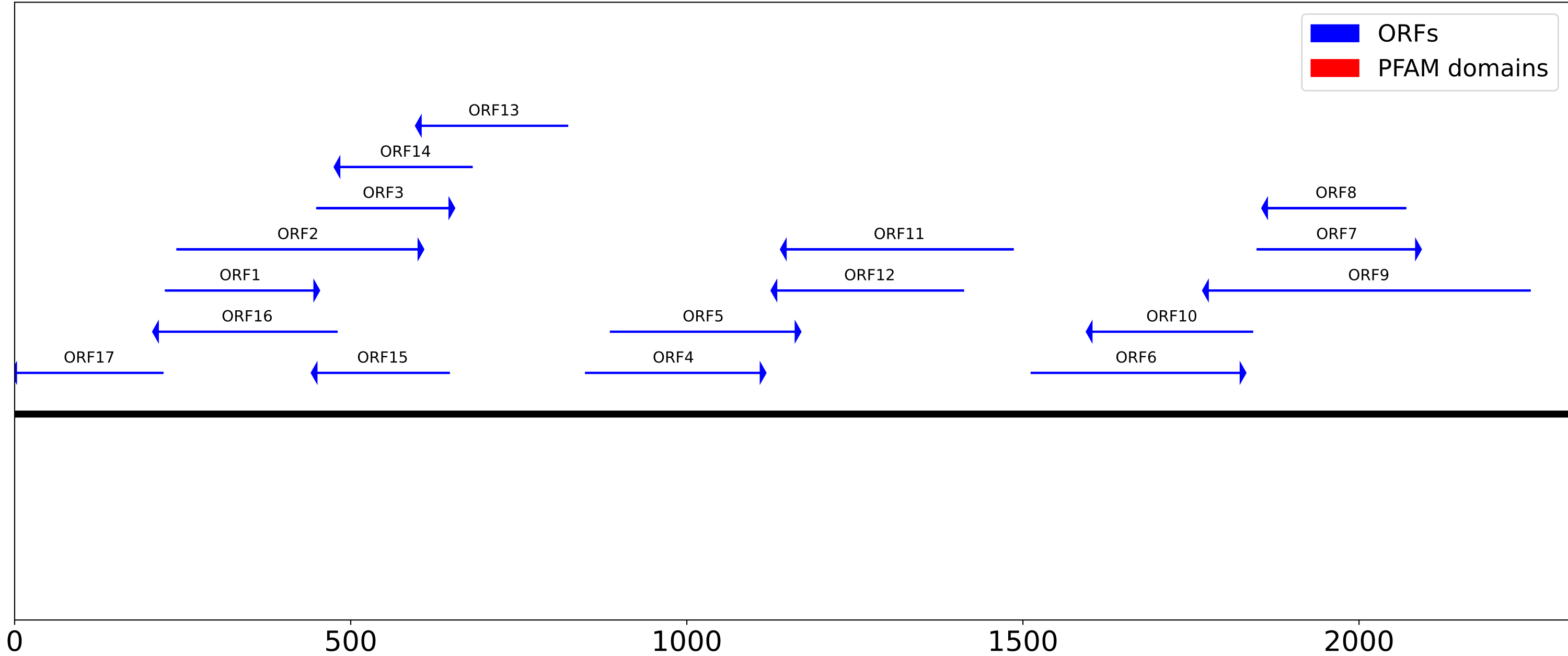
Before TEtrimmer 2314 bp



After TEtrimmer ORF and PFAM domain plot



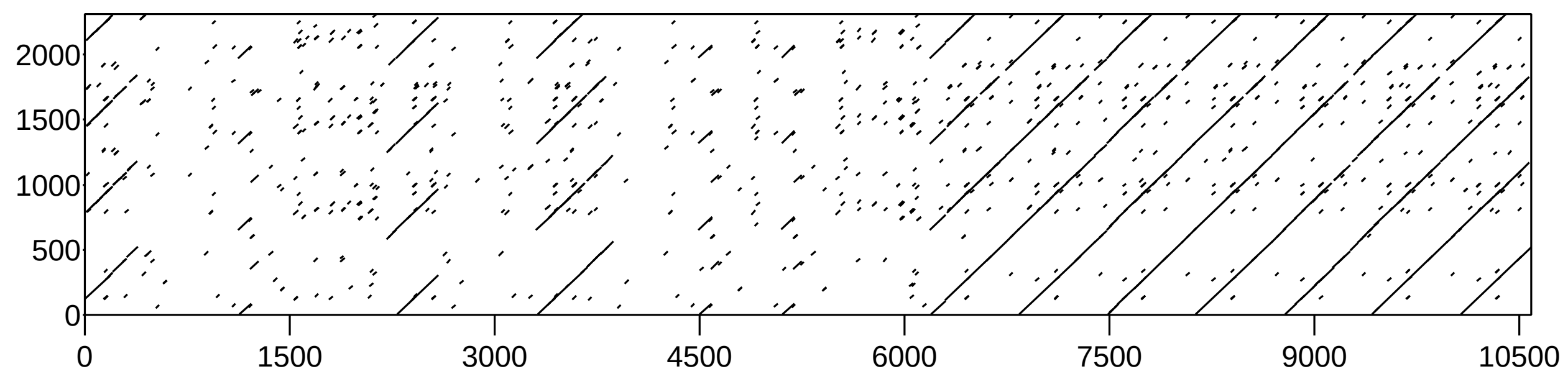
Before TEtrimmer ORF and PFAM domain plot



Dotplot

(windowsize = 25, threshold = 50.00 09/10/25)

onsensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)