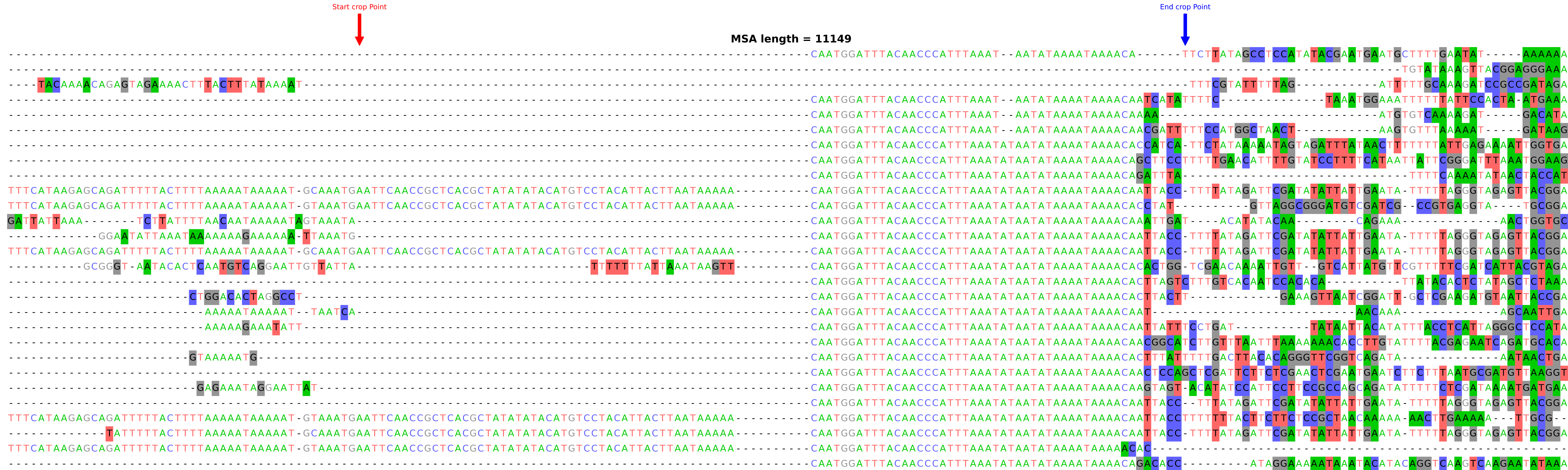
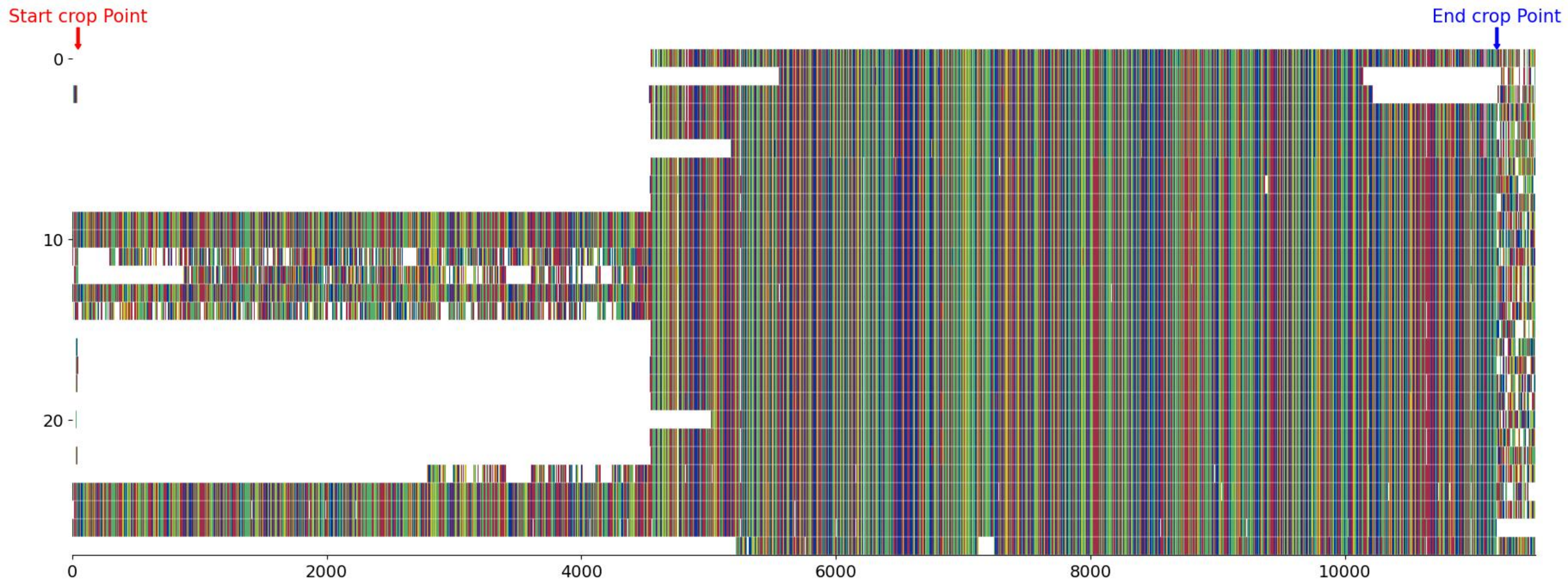
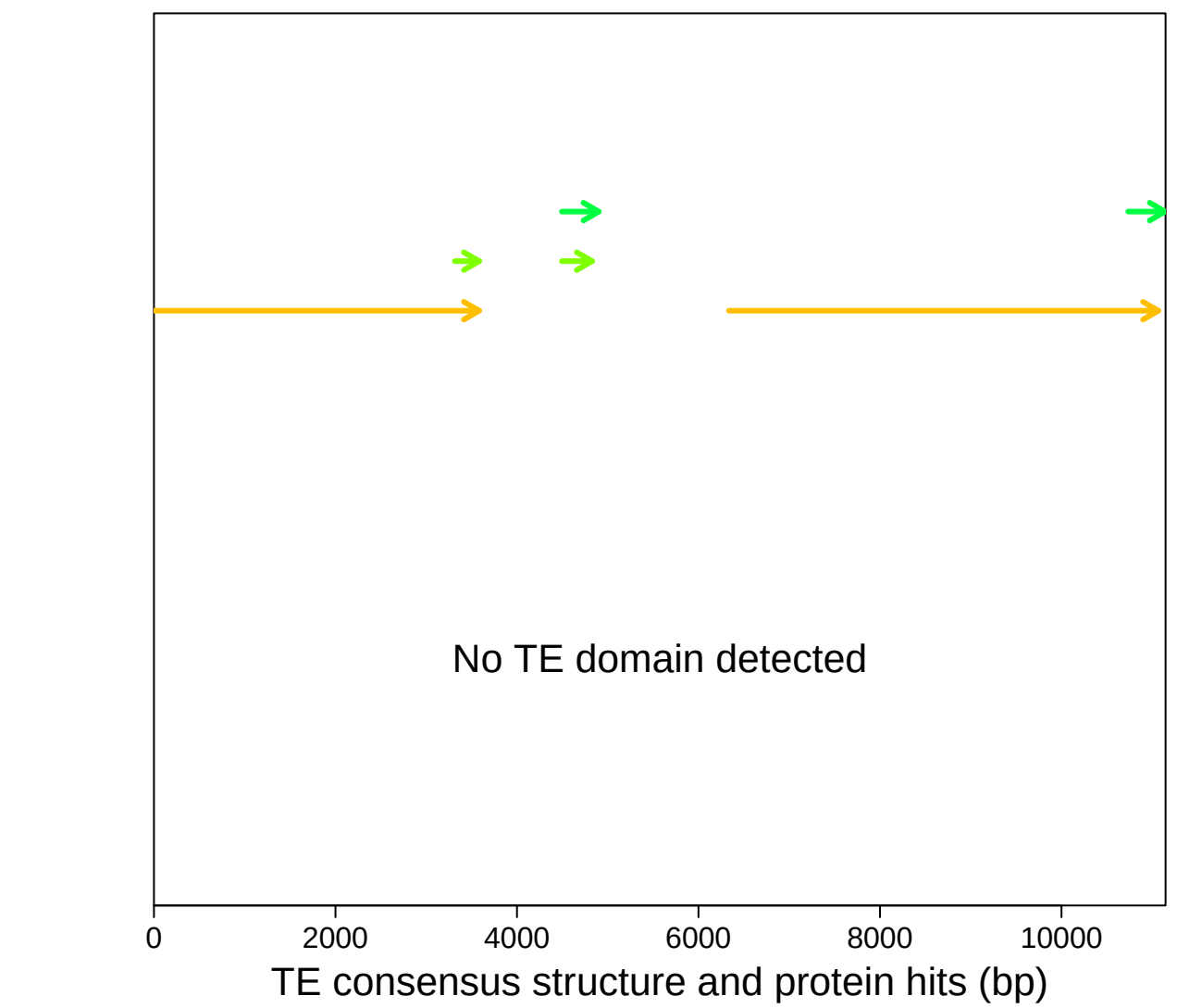
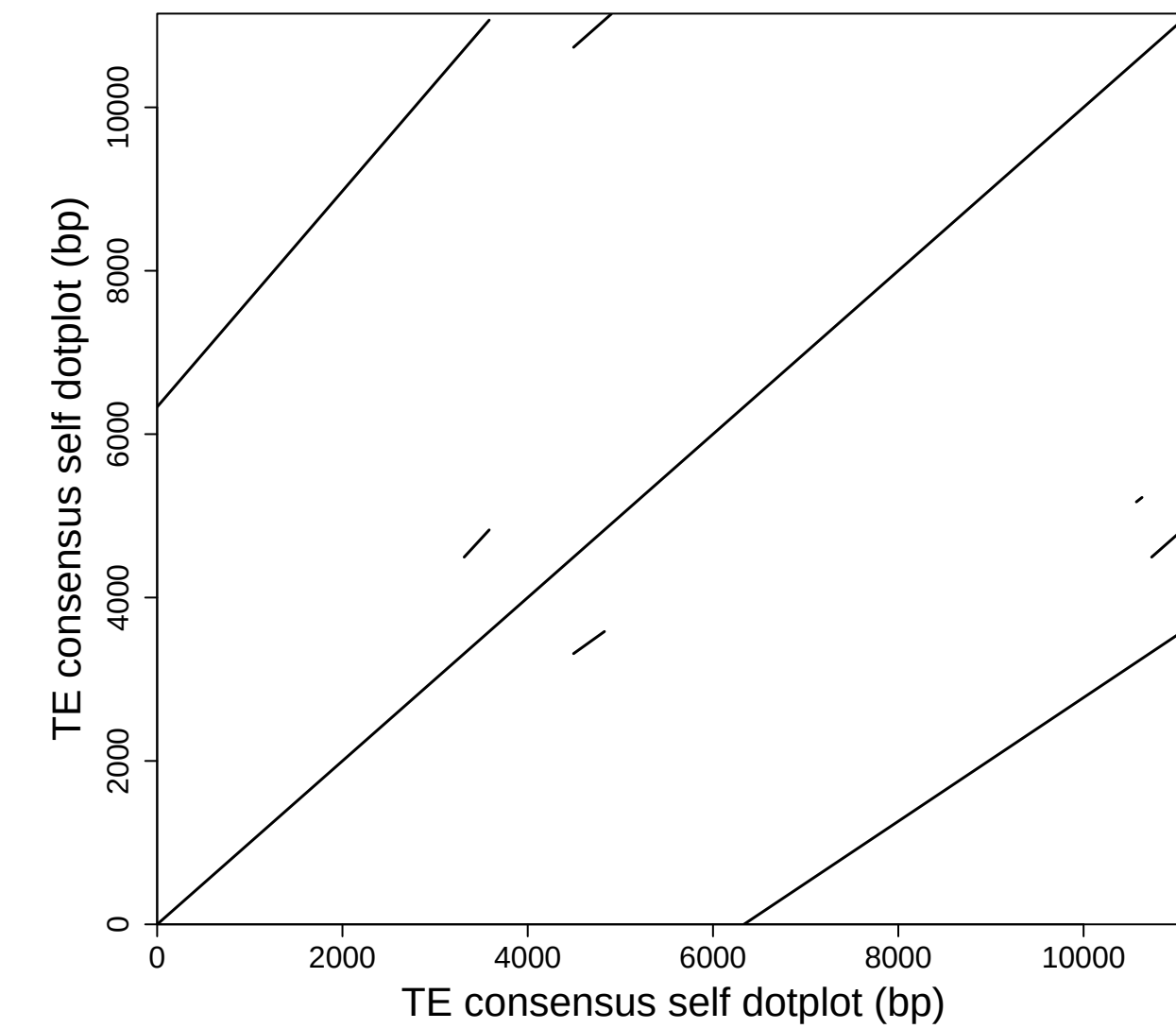
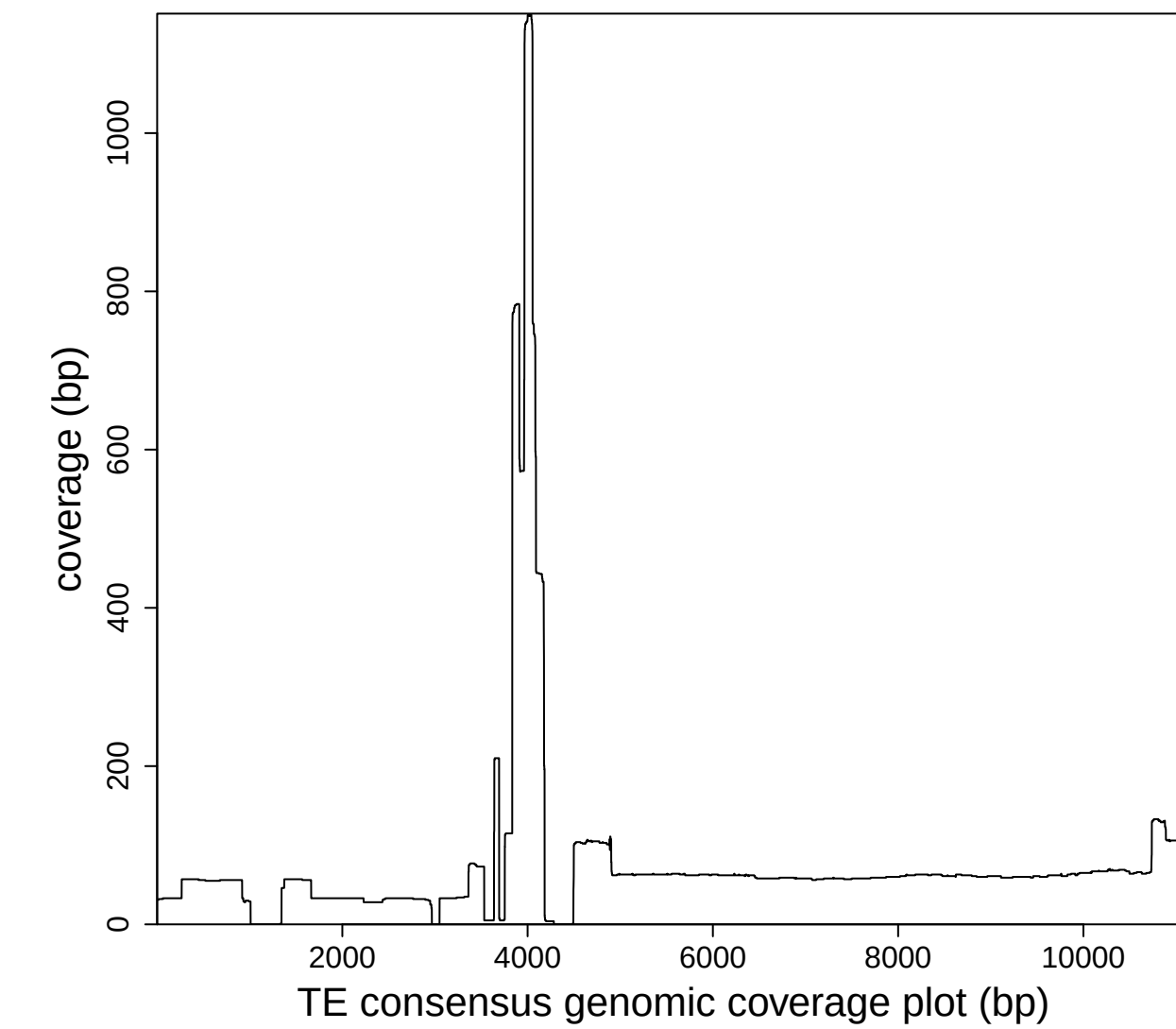
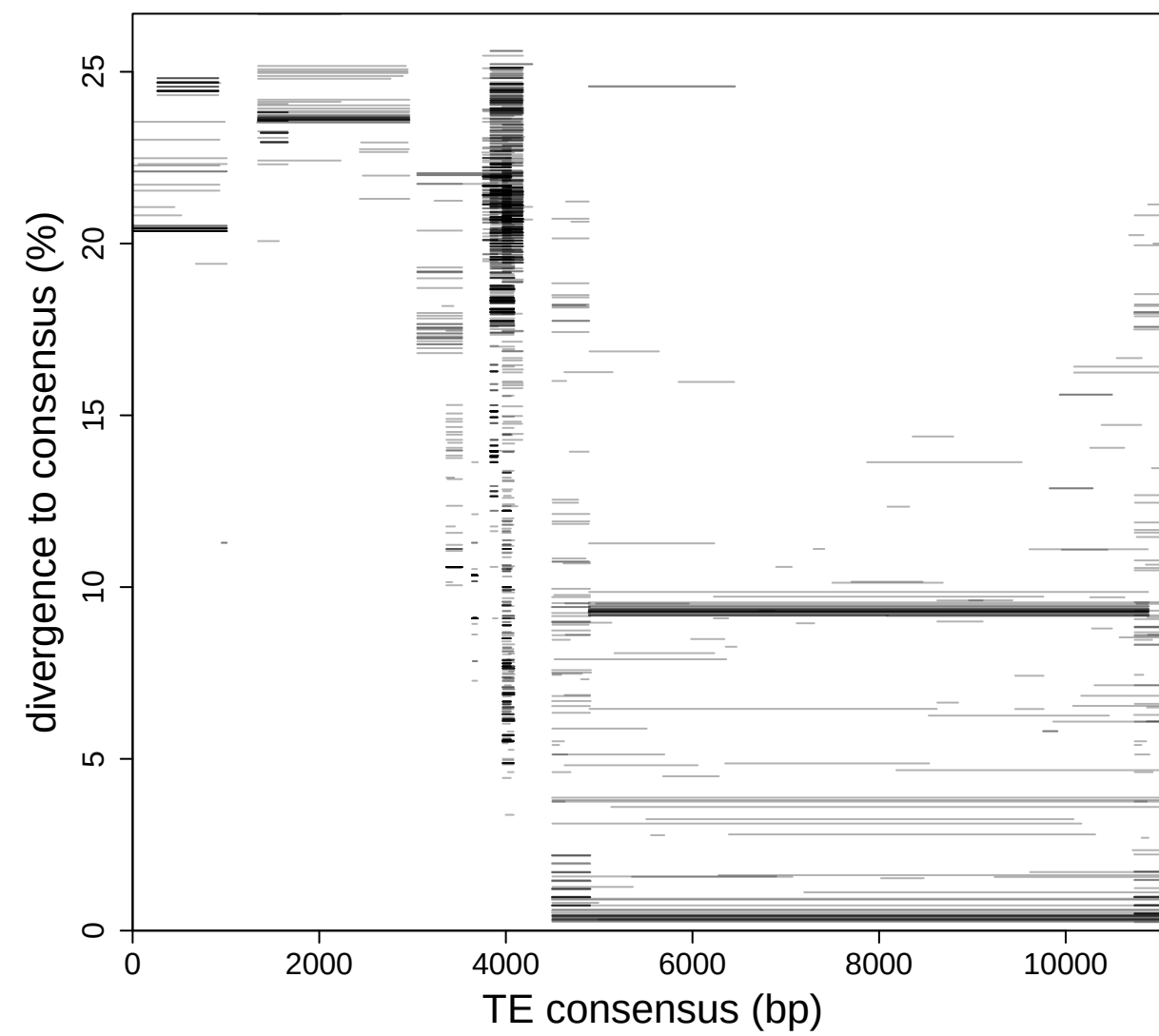






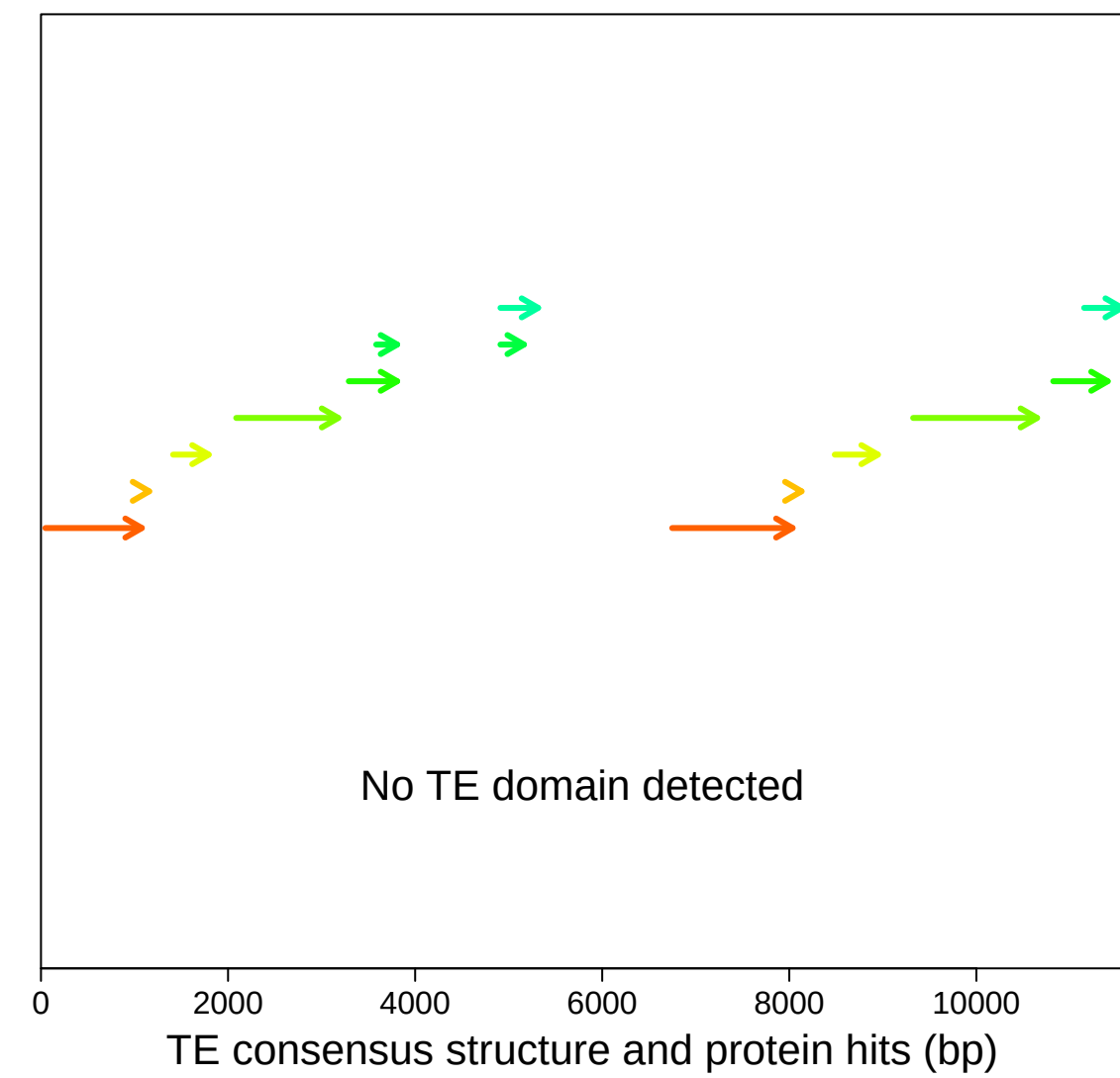
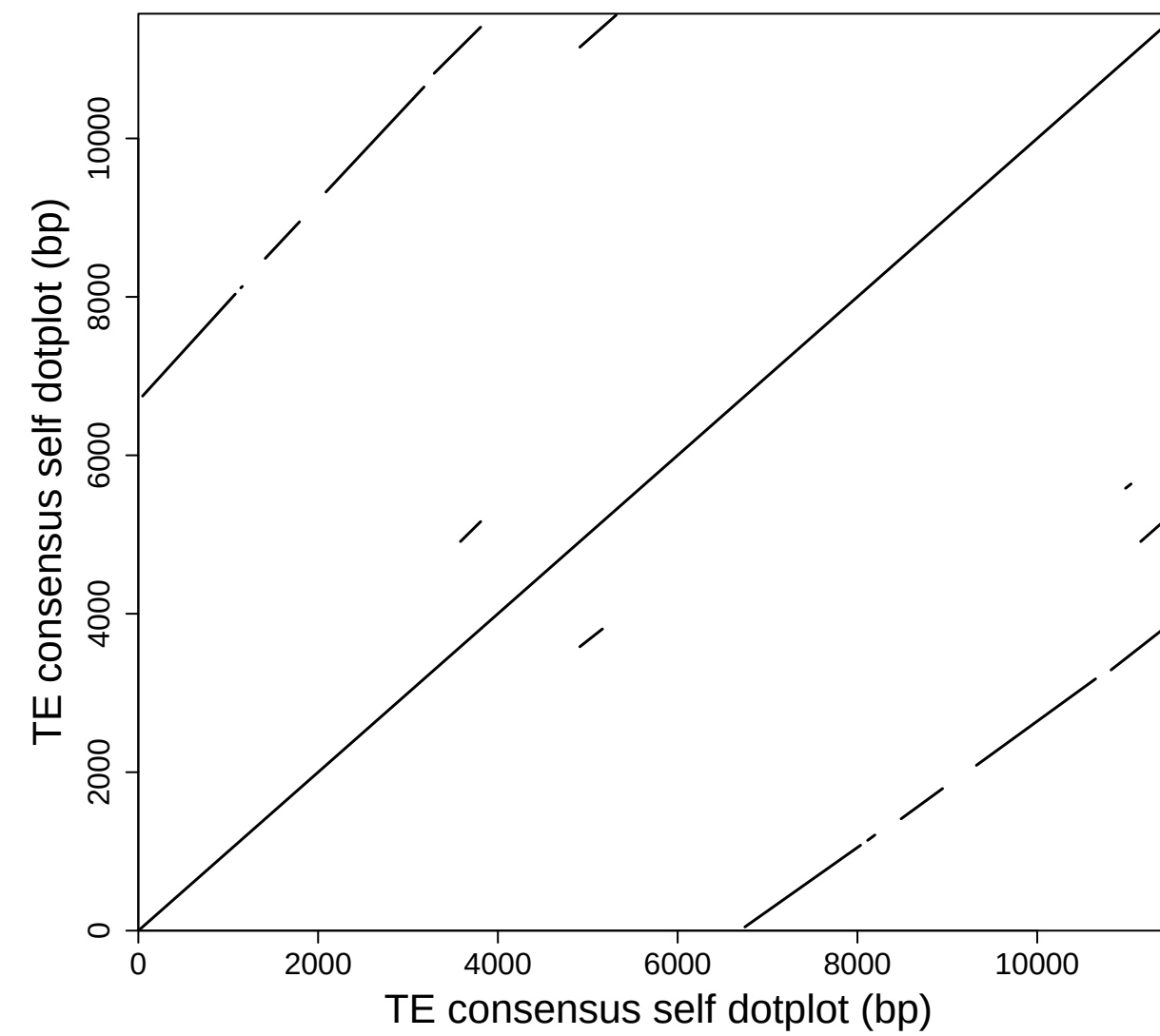
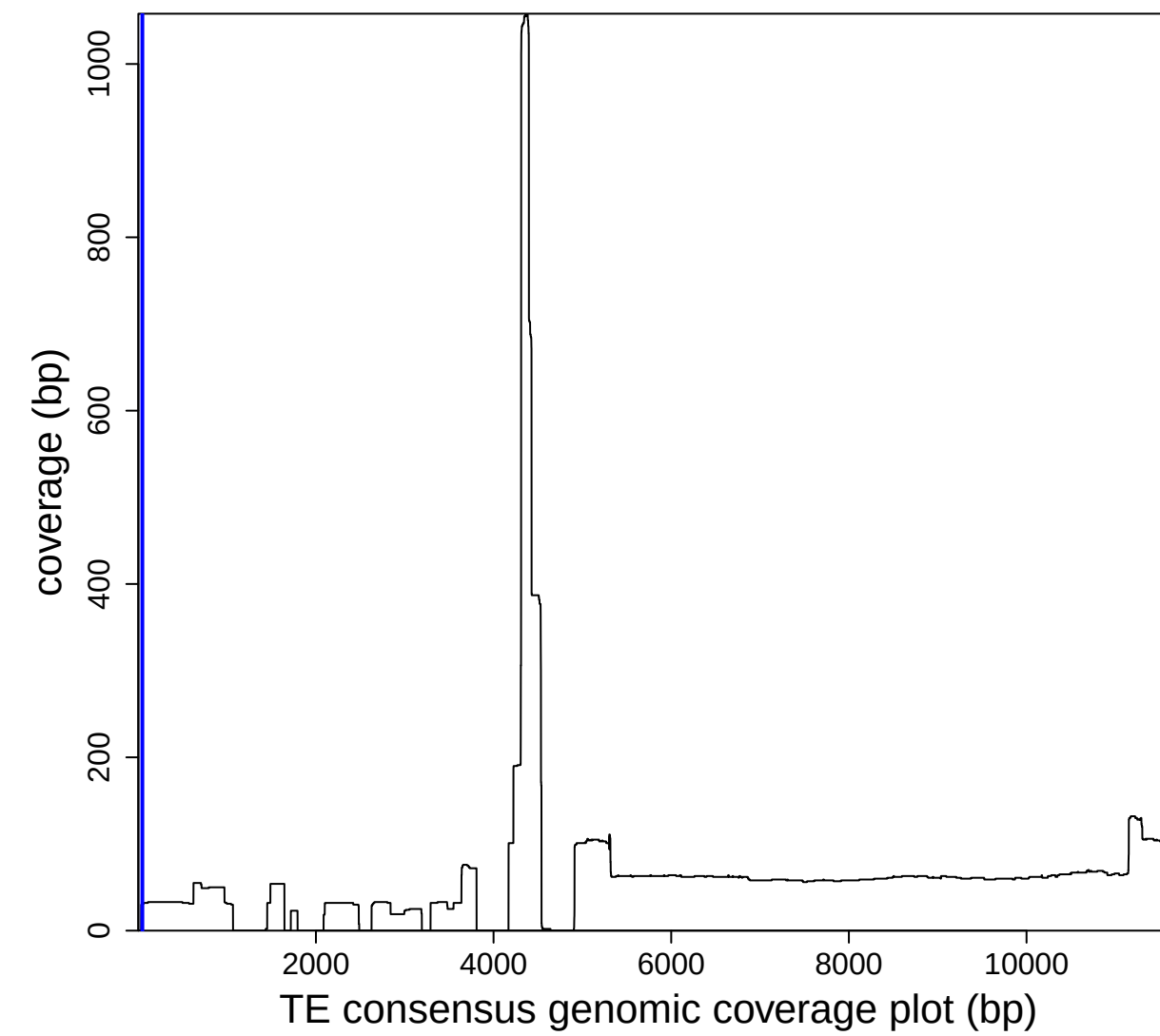
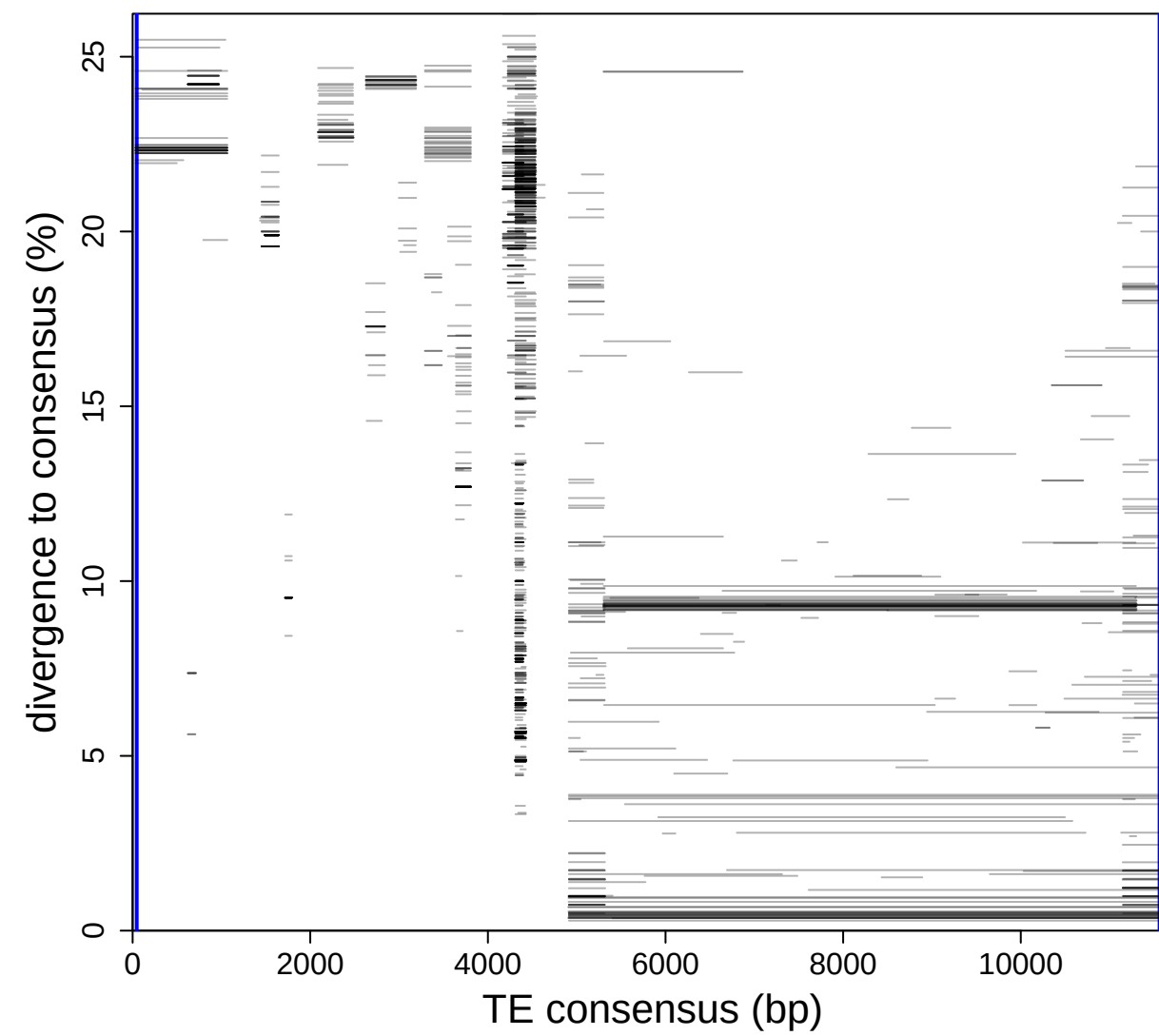
bed uf.bed g_2.bed fm_1.bed 0_0_bcln.fa aln.fa cl.fa gs.fa ce
size: 11149bp; fragments: 2048; full length: 0 (>=10034.1bp)

After TEtrimmer 11149 bp



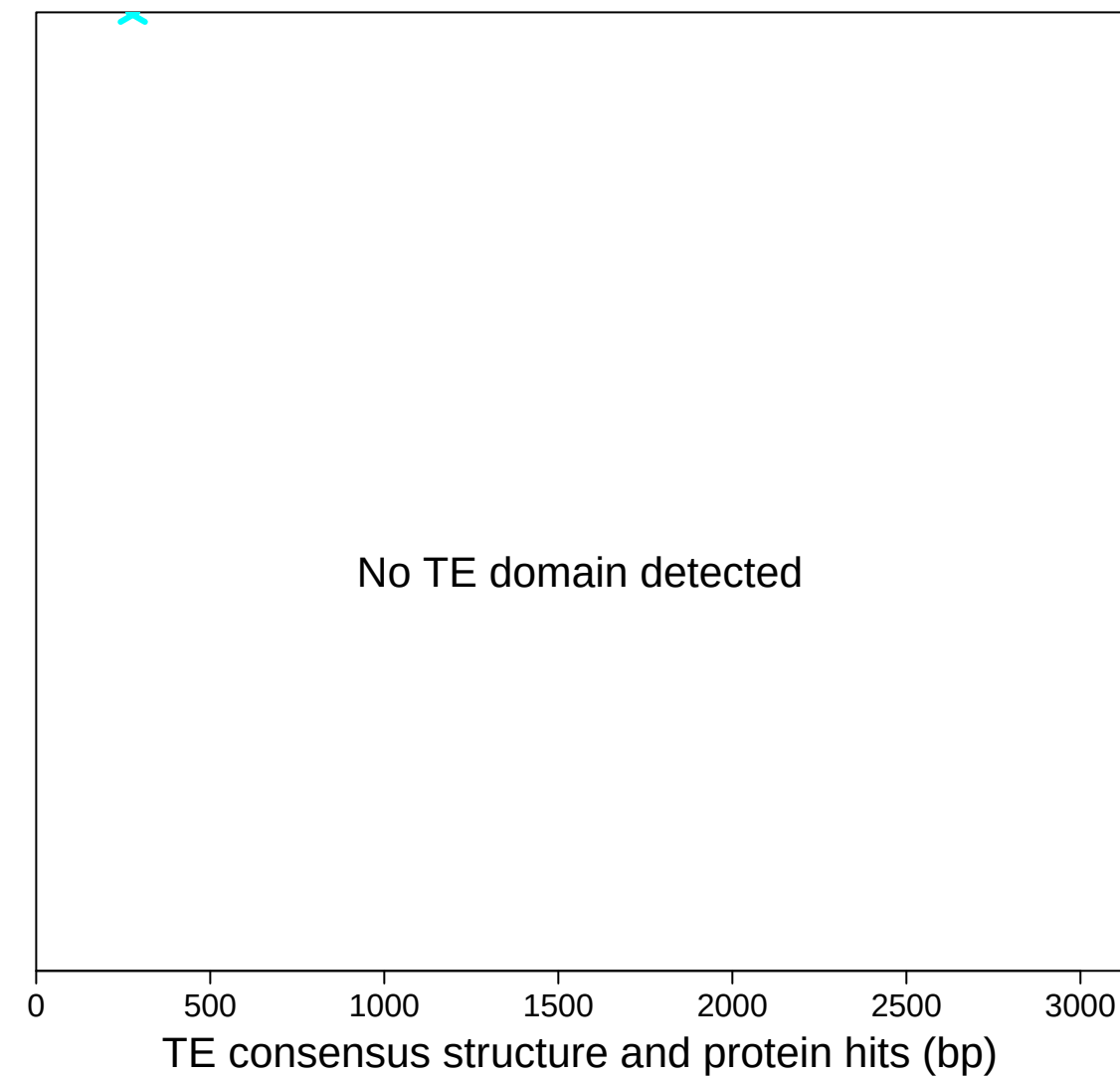
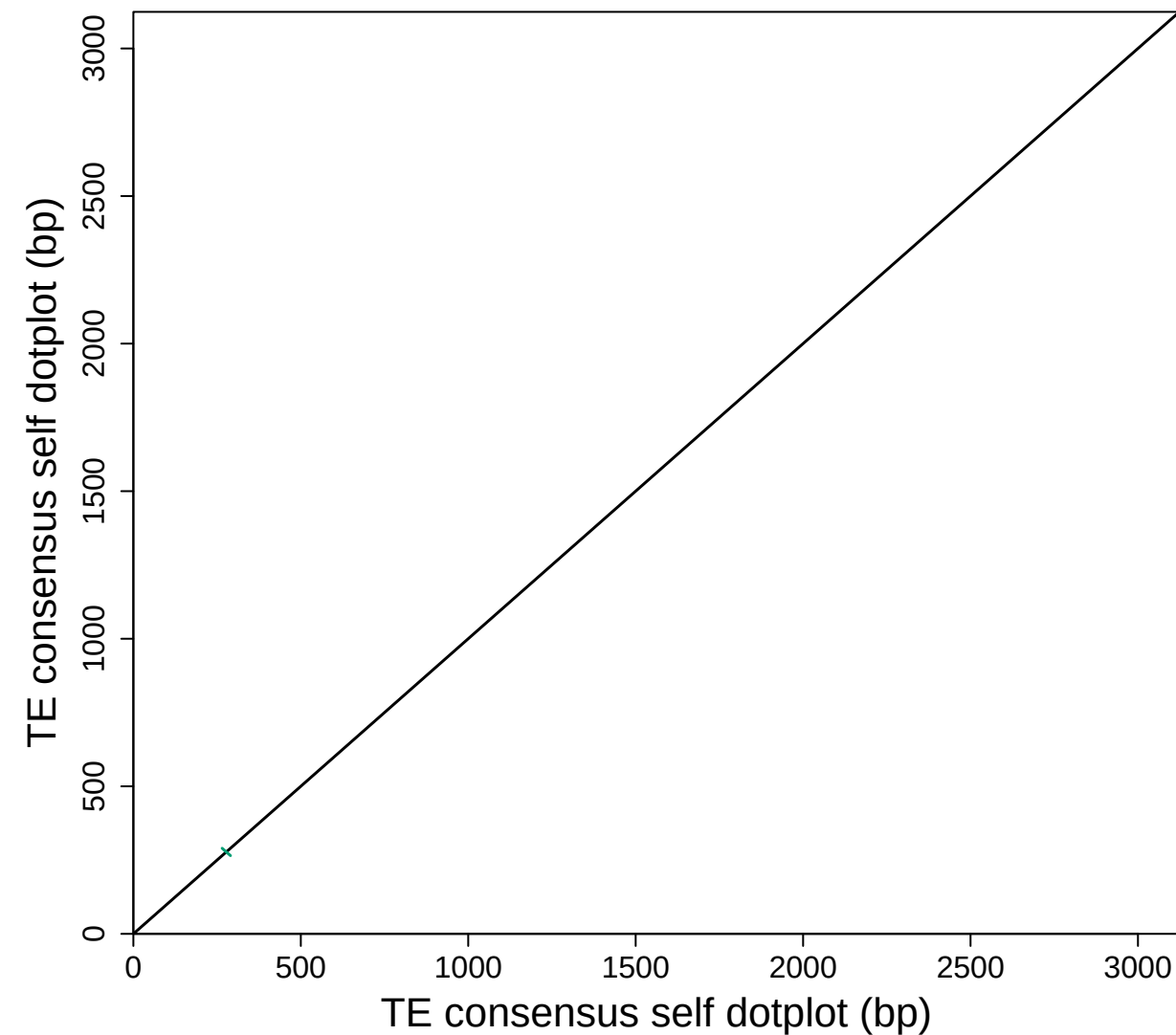
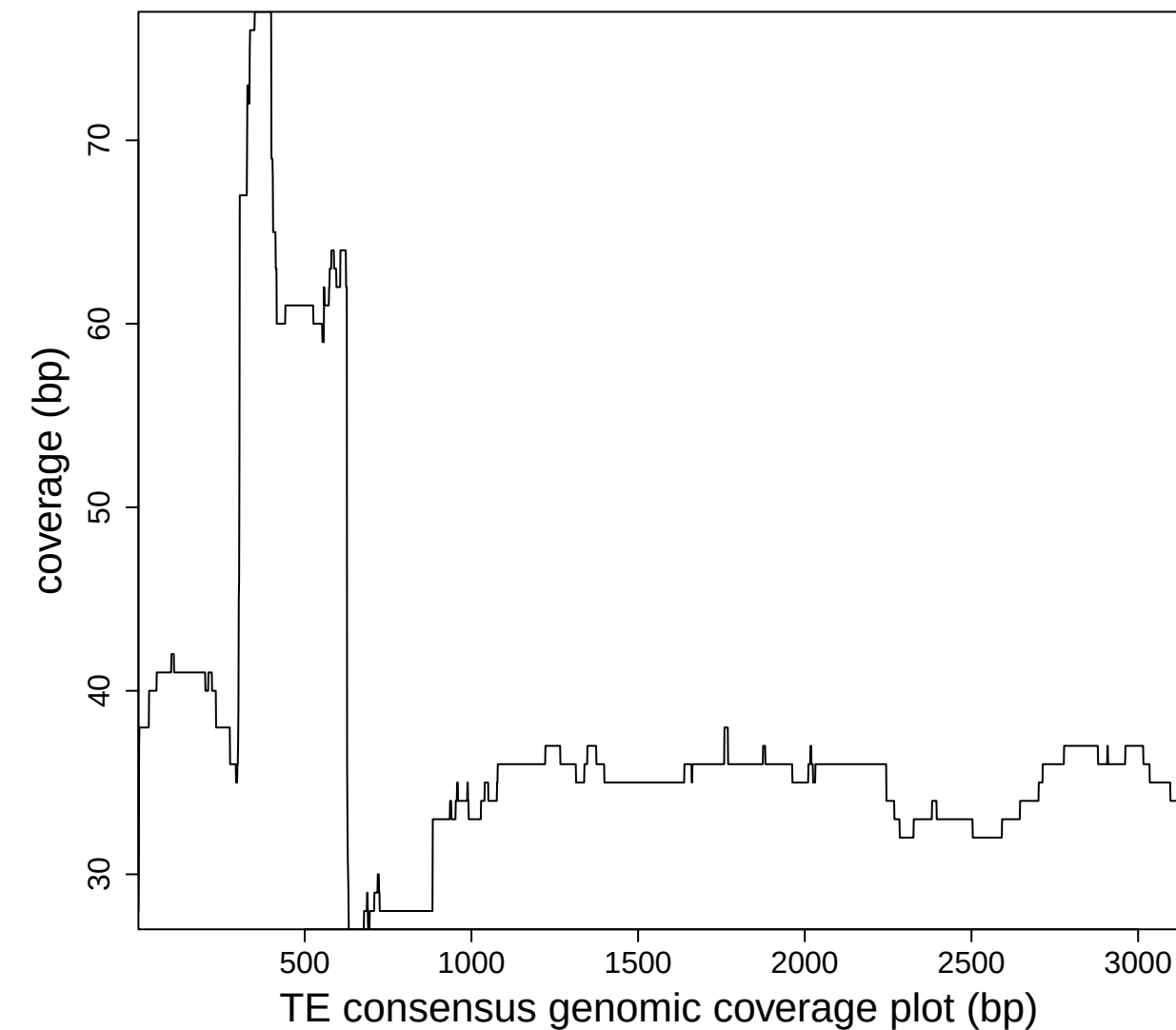
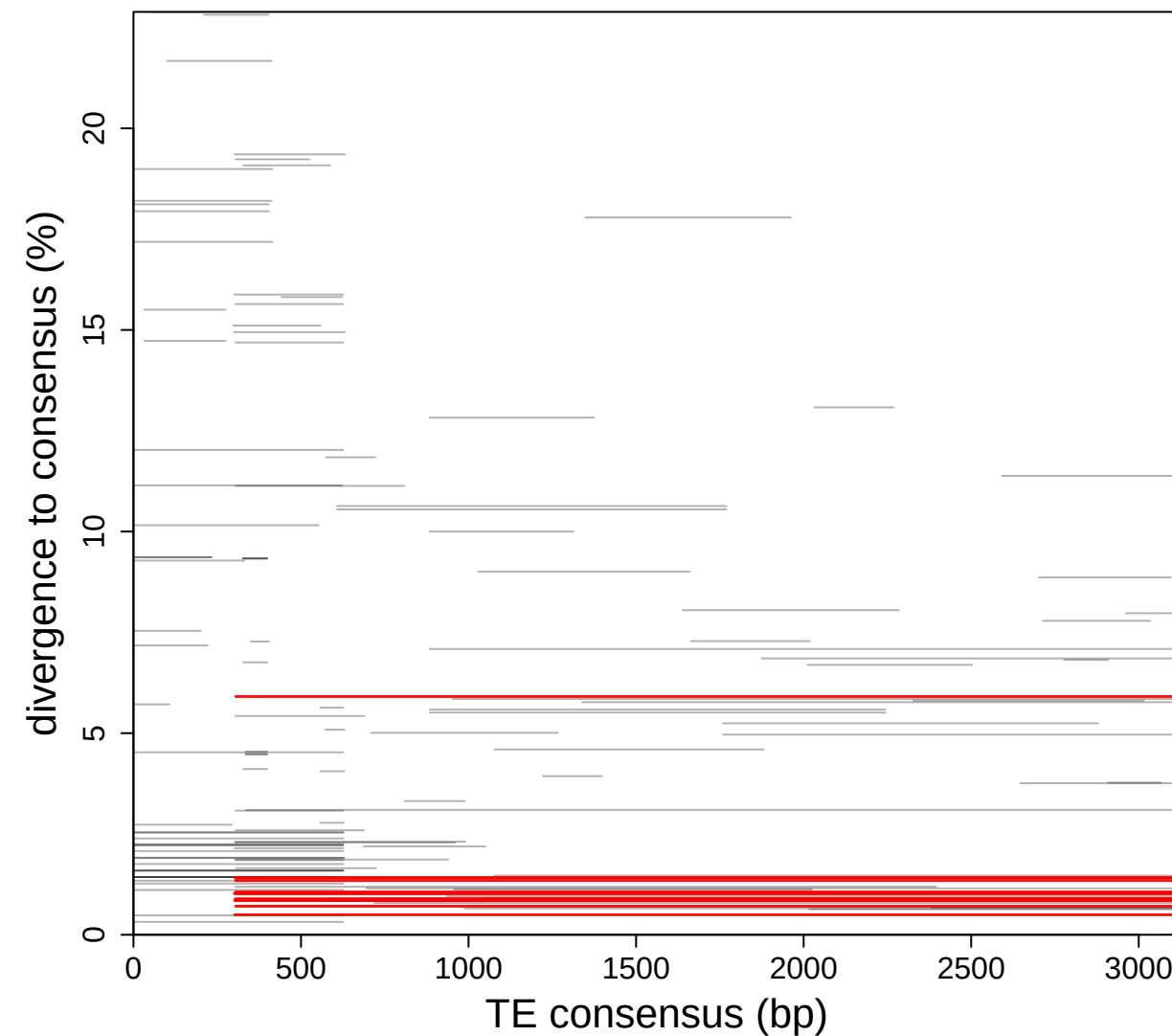
ily_678.fasta.b.bed_uf.bed_g_2.bed_fm_1.bed_0_0_bcln.fa_aln.f
size: 11713bp; fragments: 1629; full length: 0 (>=10541.7bp)

After TEtrimmer Extended plot Blue lines are boundaries

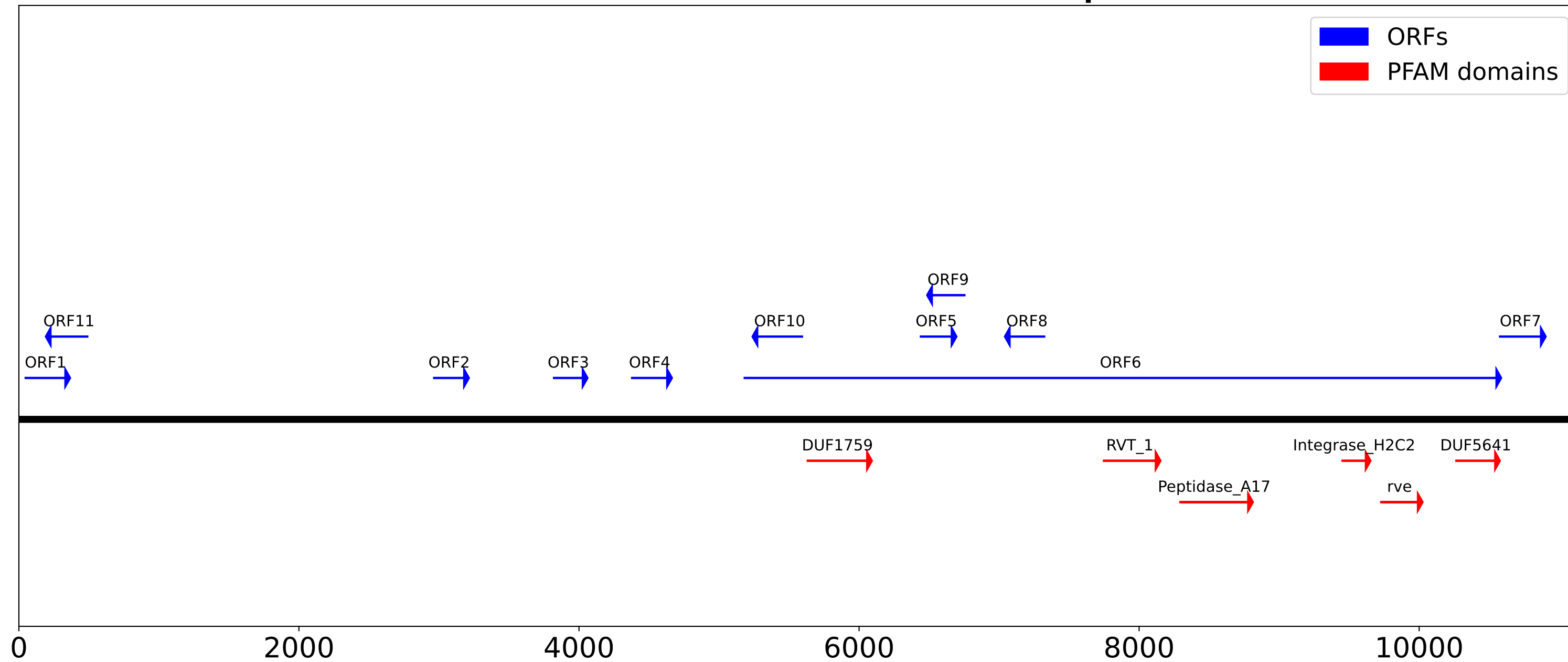


TE: rnd_5_family_678
size: 3124bp; fragments: 135; full length: 14 (≥ 2811.6 bp)

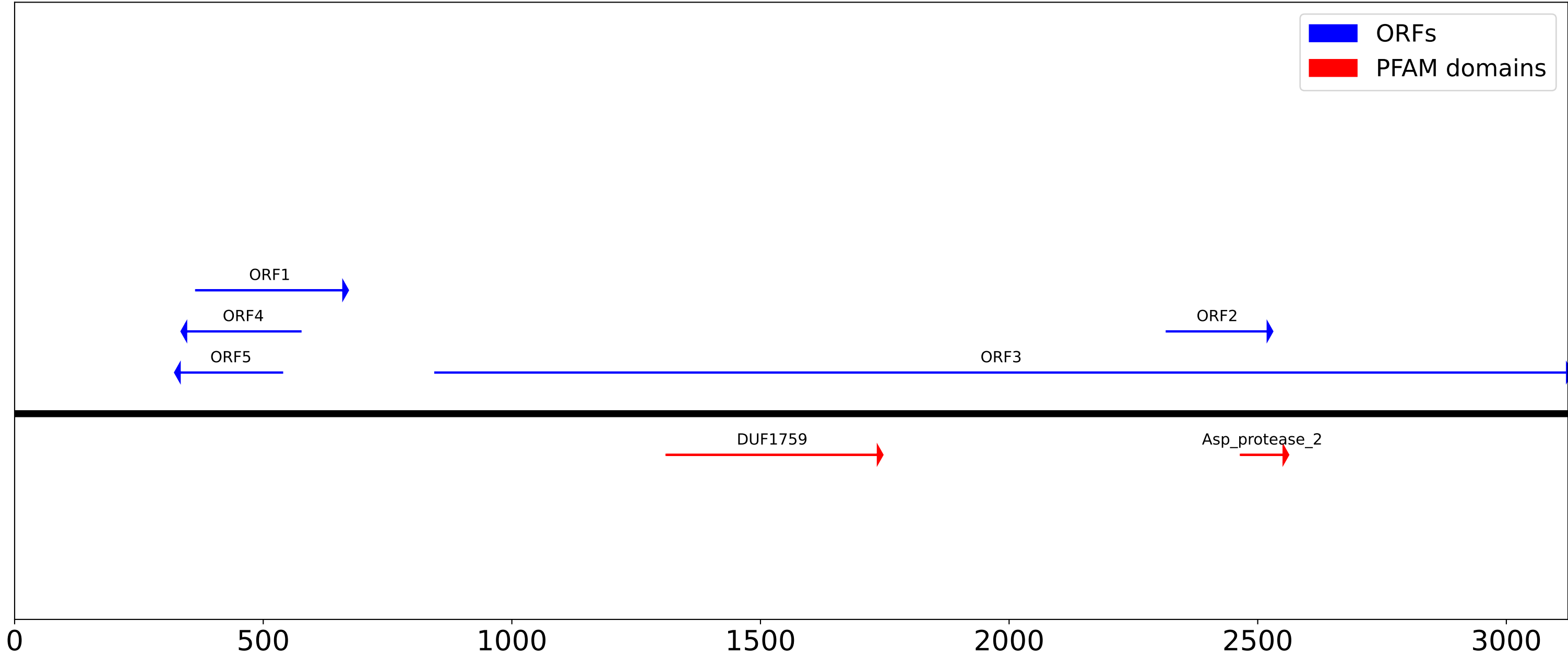
Before TEtrimmer 3124 bp



After TEtrimmer ORF and PFAM domain plot



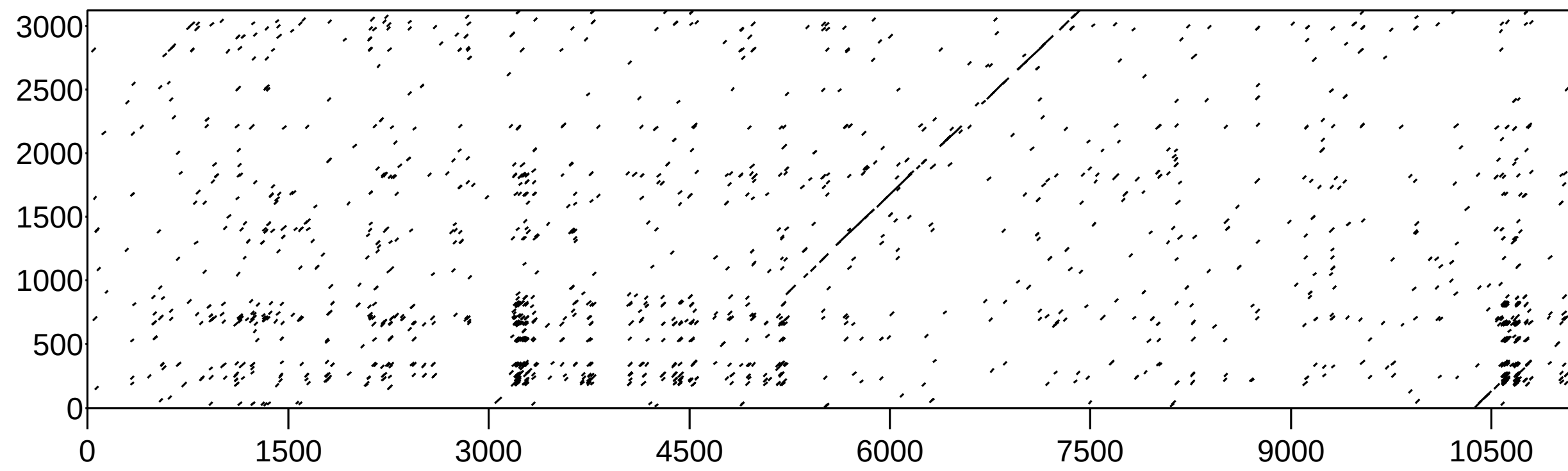
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)