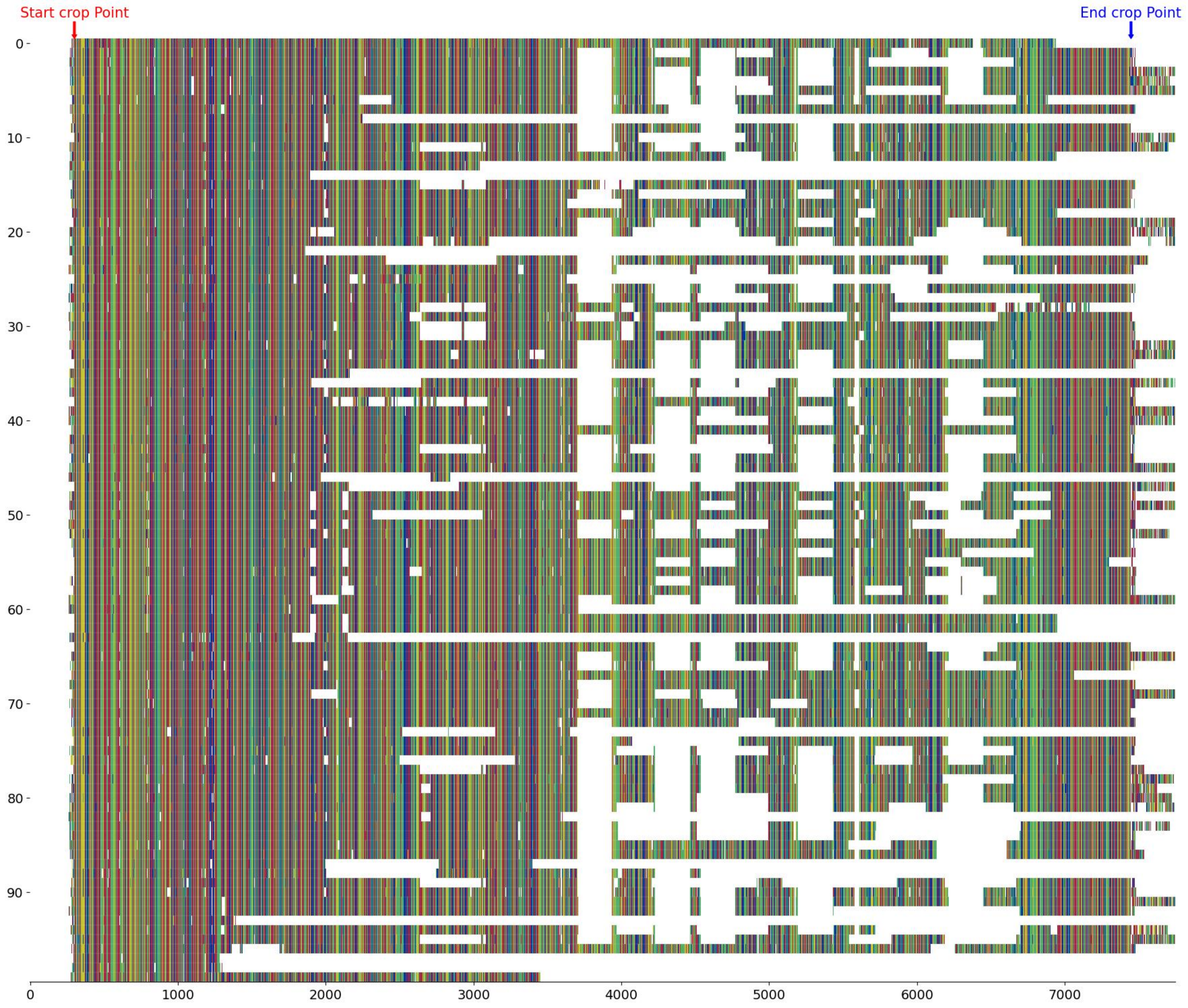




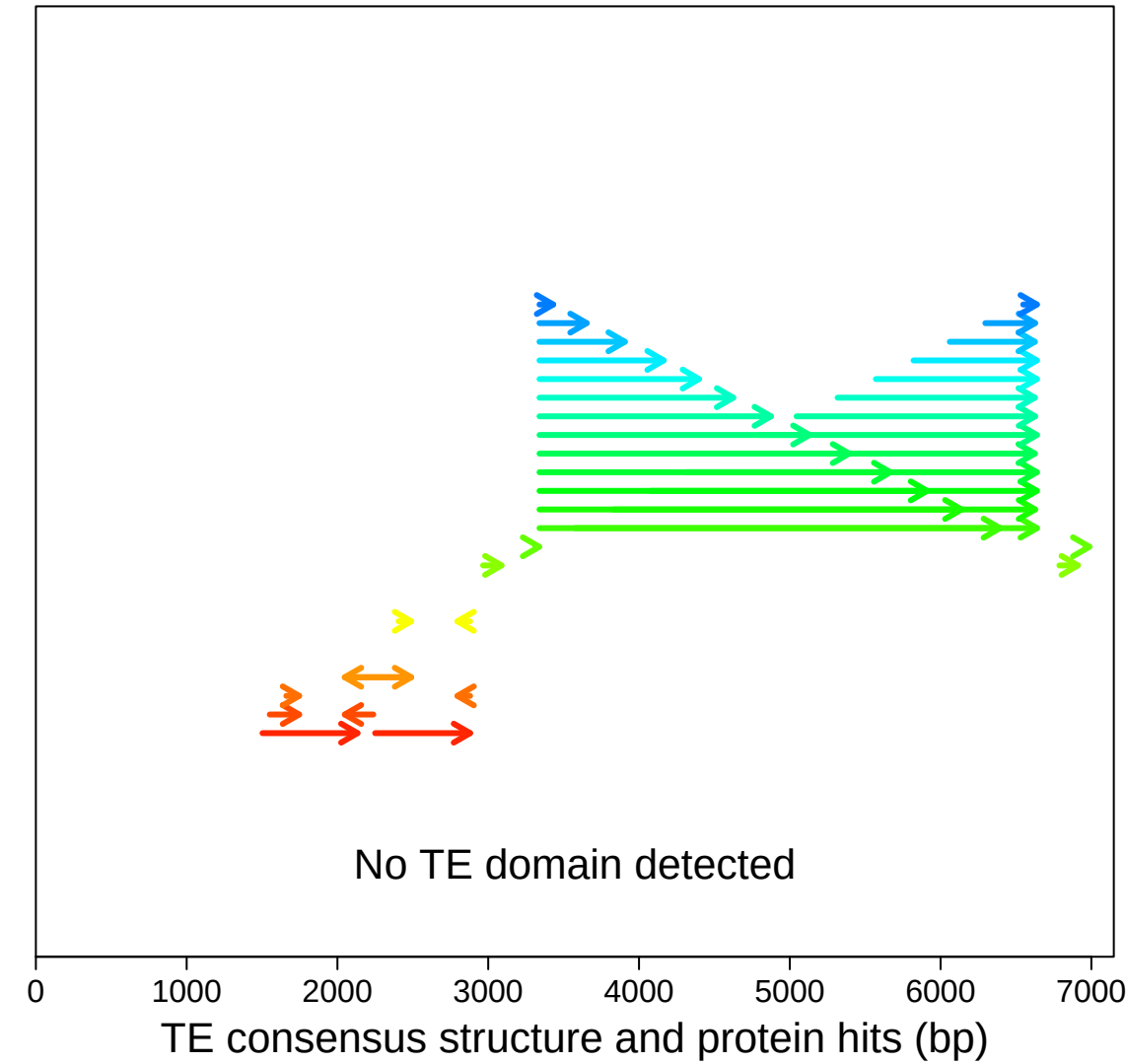
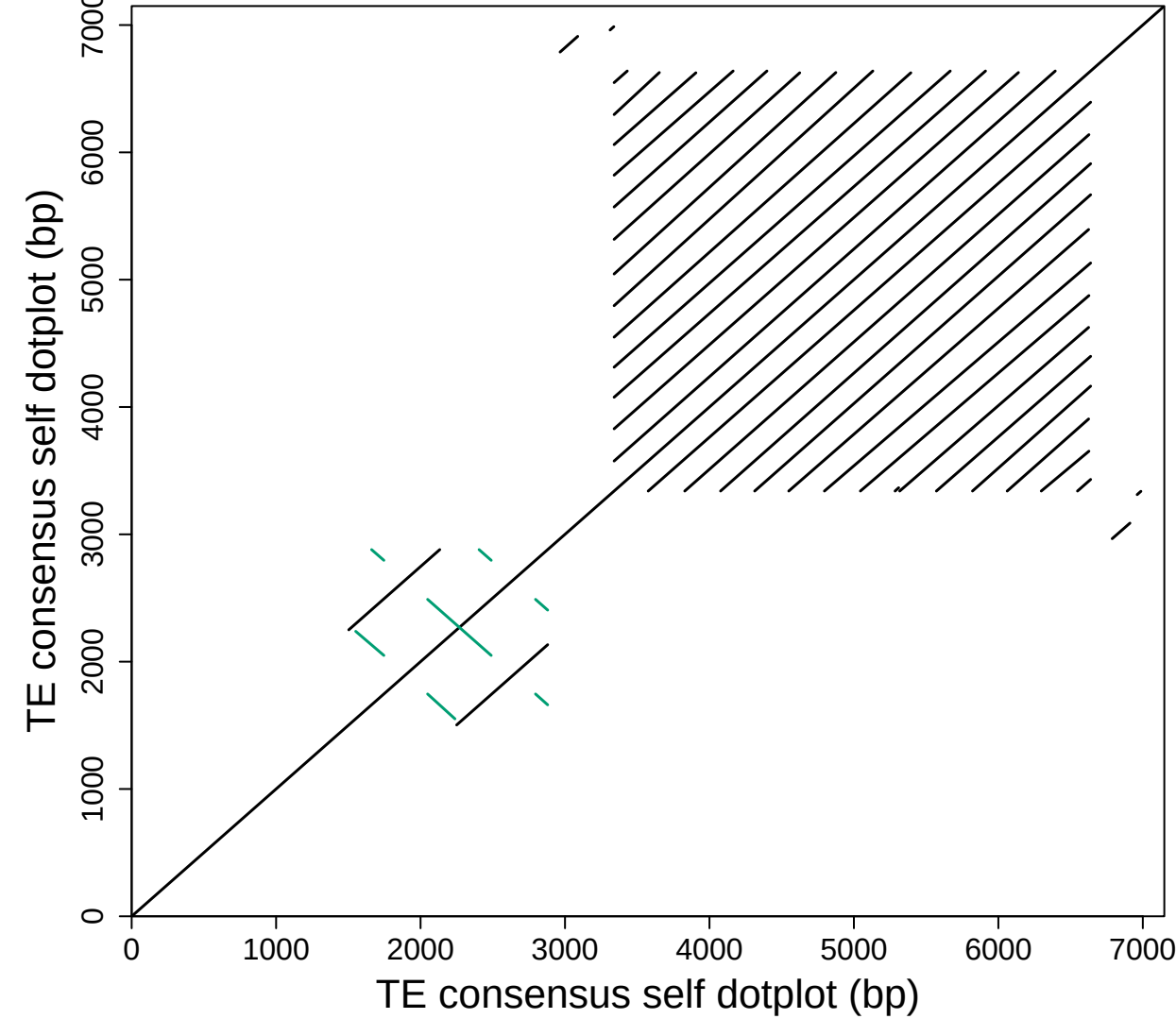
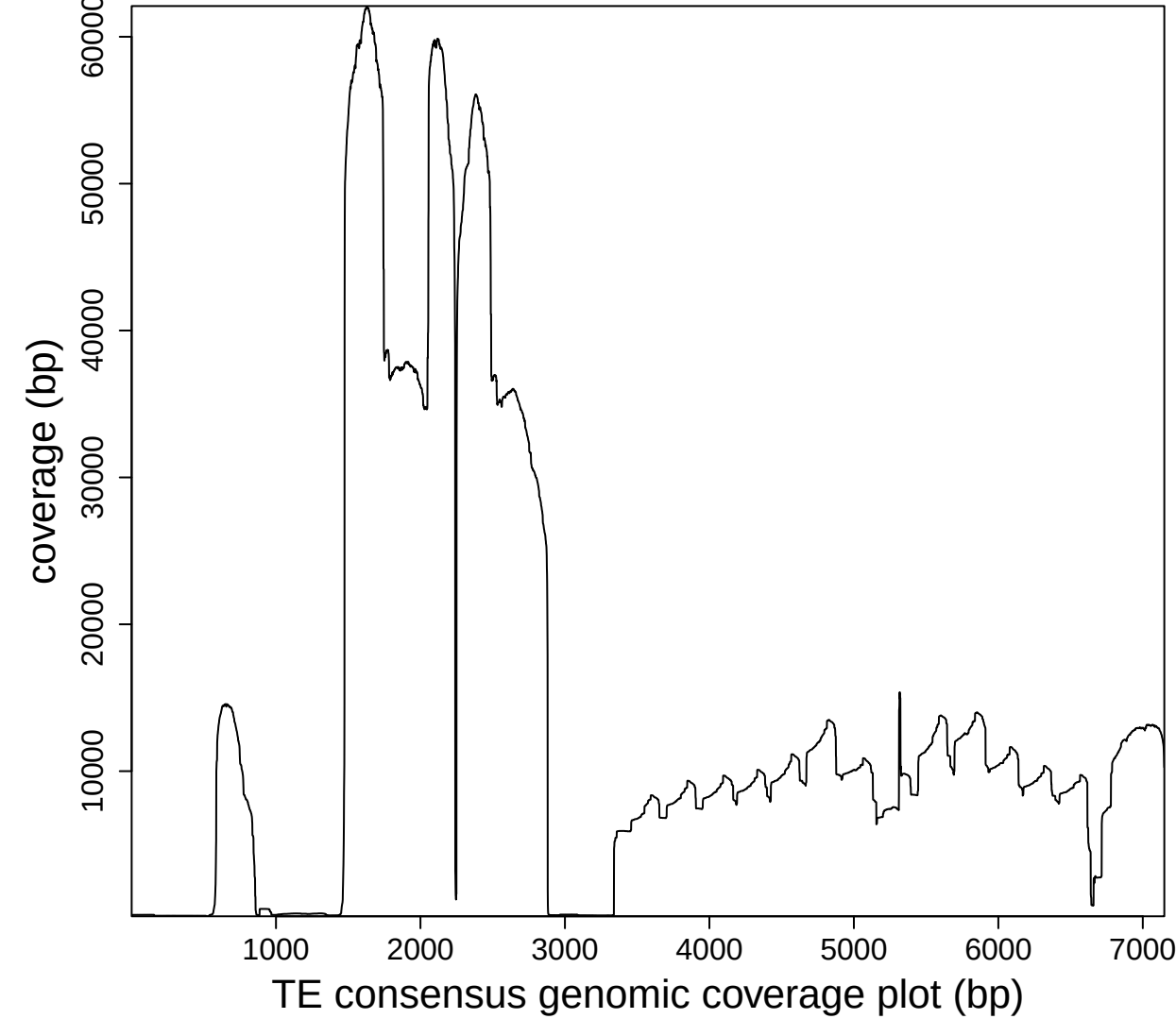
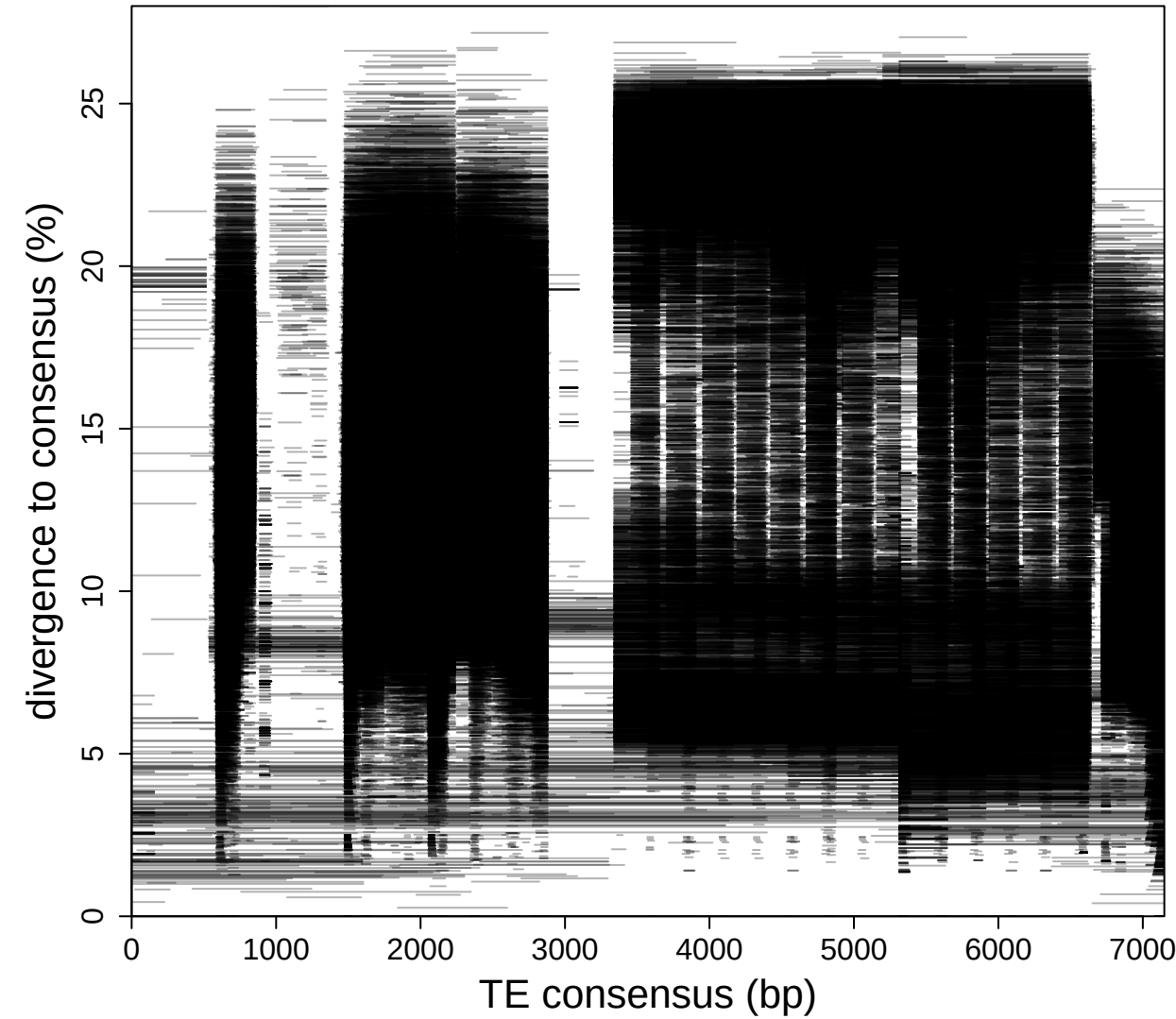
Start crop Point

End crop Point



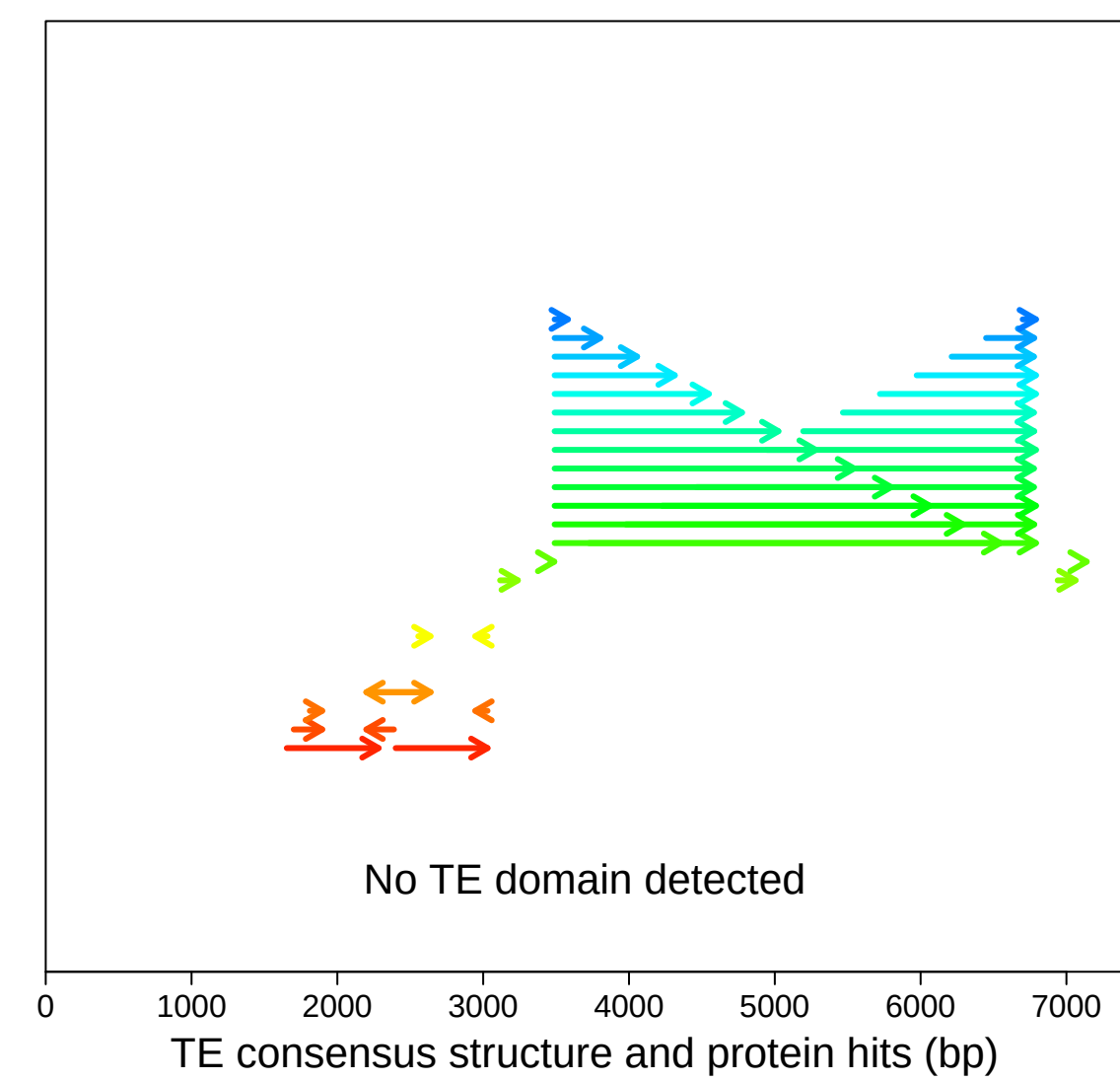
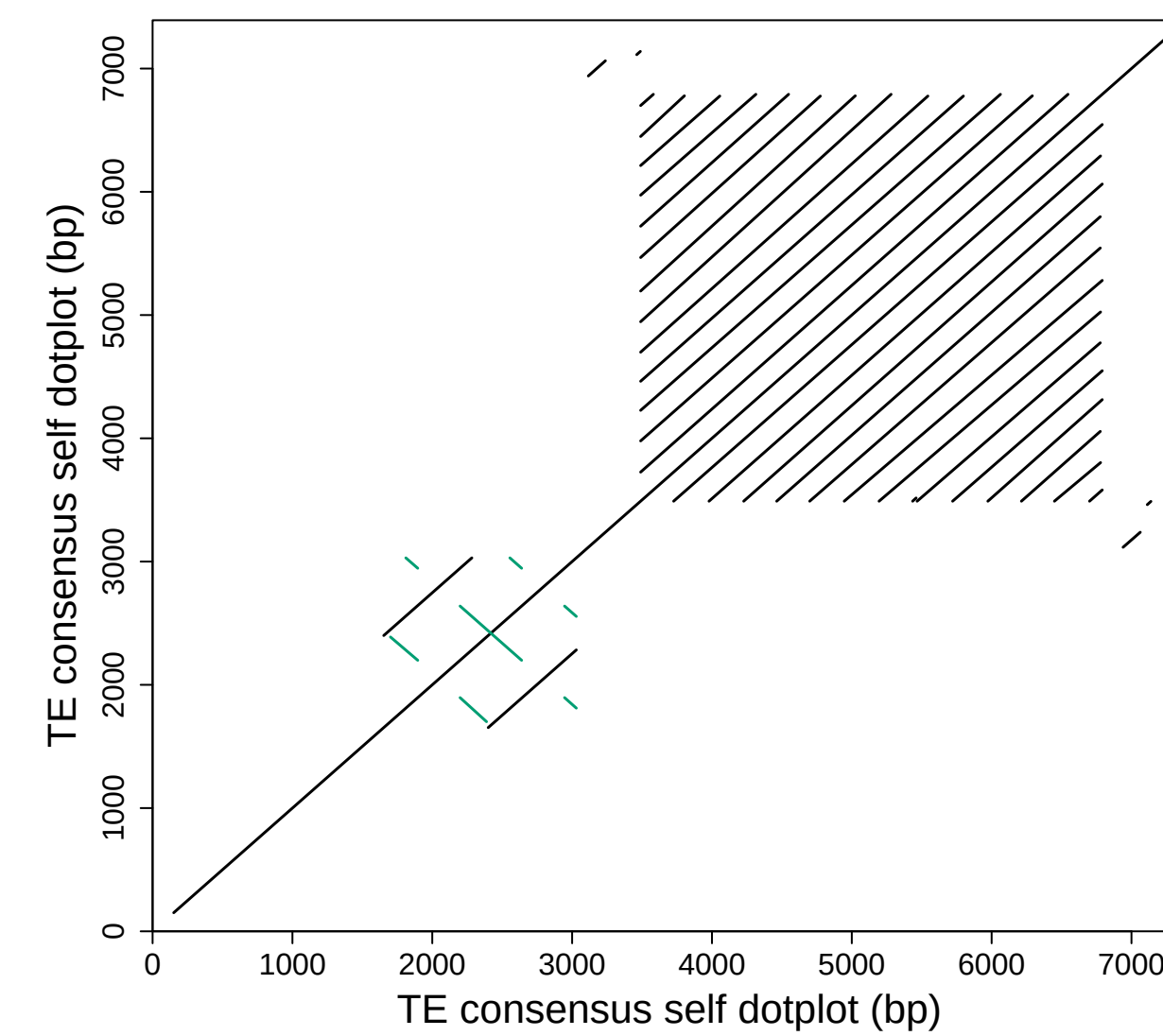
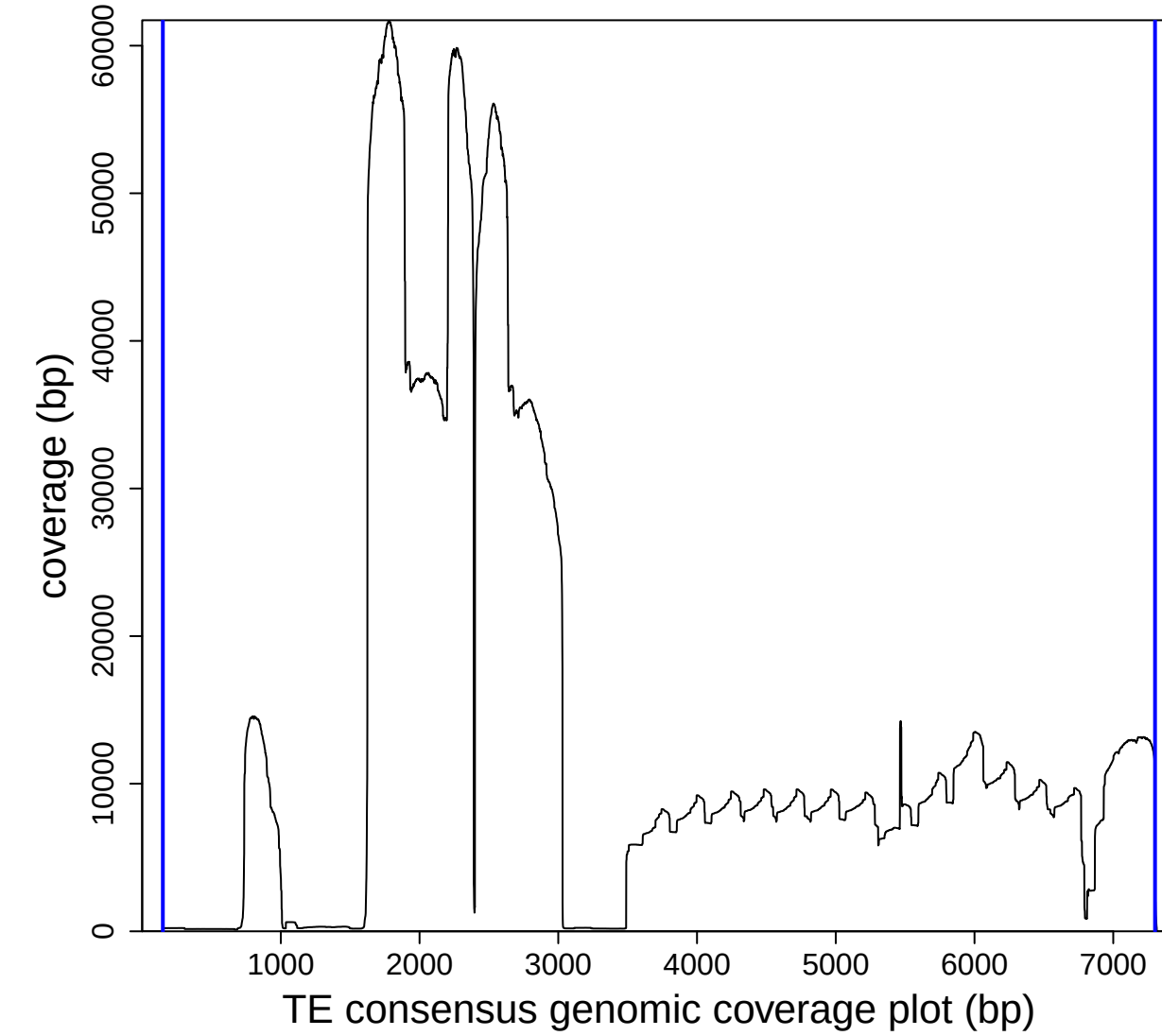
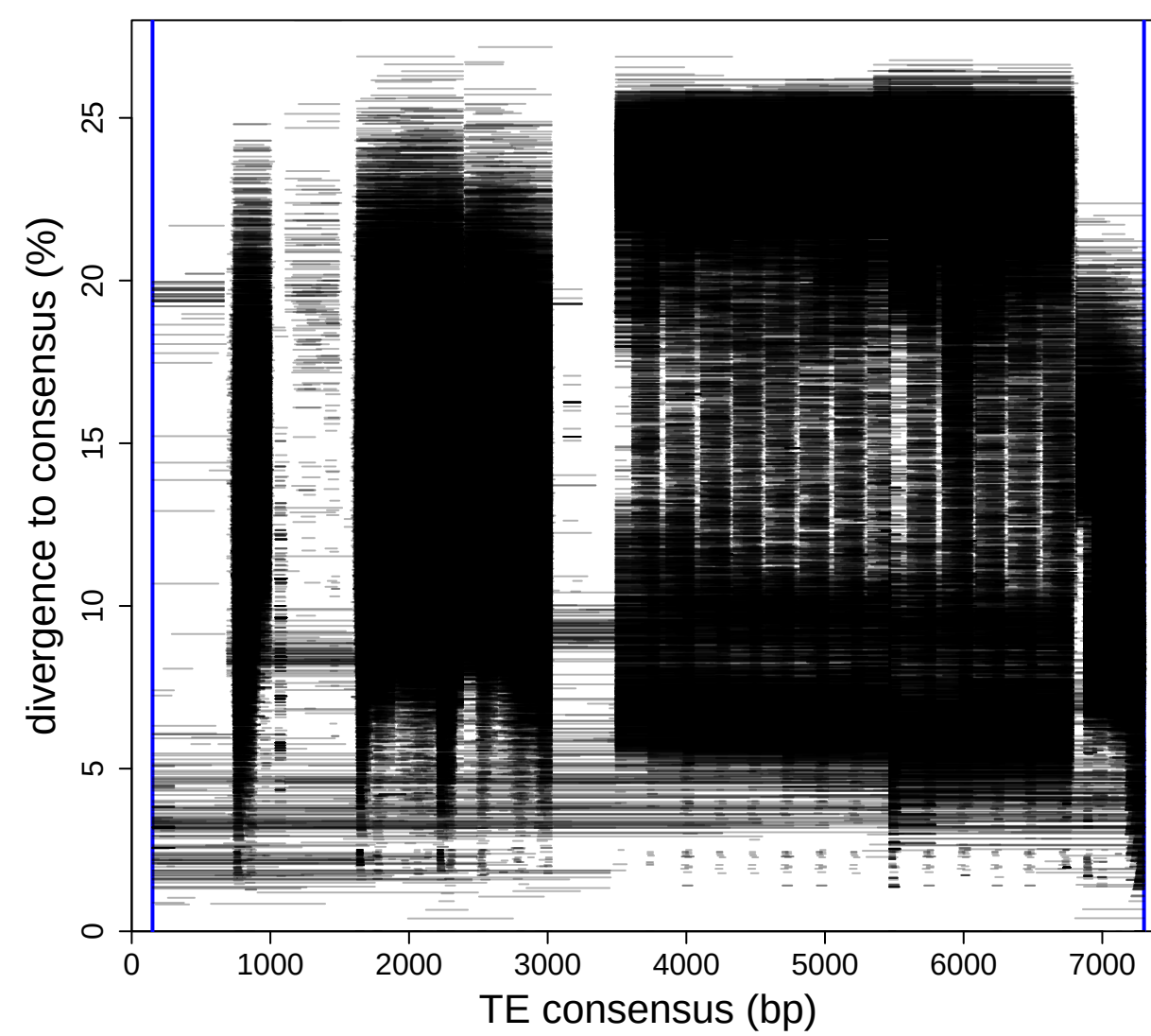
ta.b.bed uf.bed g_1.bed fm_1.bed 0 0 bcln.fa aln.fa cl.fa gs.f
size: 7149bp; fragments: 267931; full length: 0 (>=6434.1bp)

After TEtrimmer 7149 bp



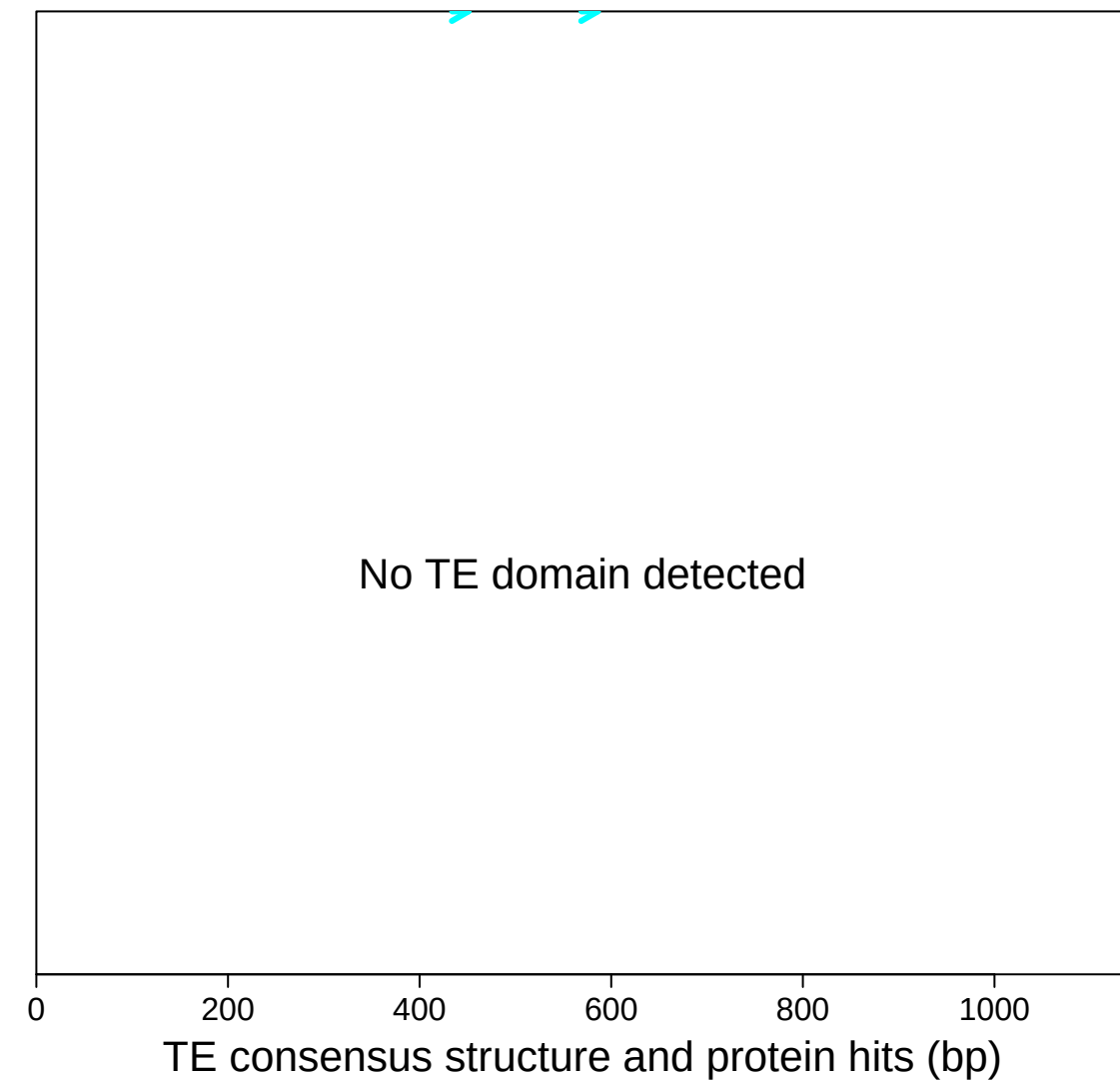
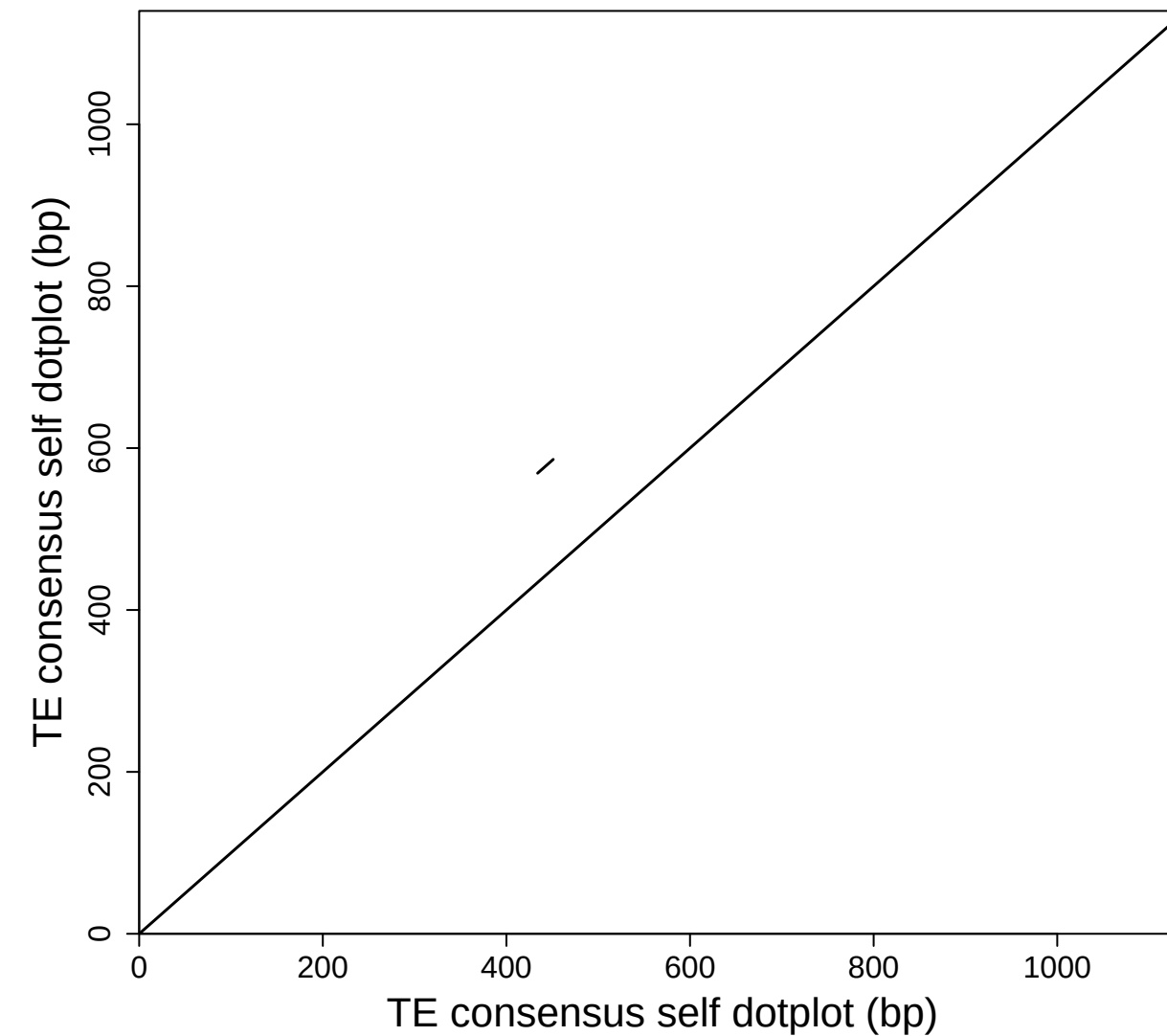
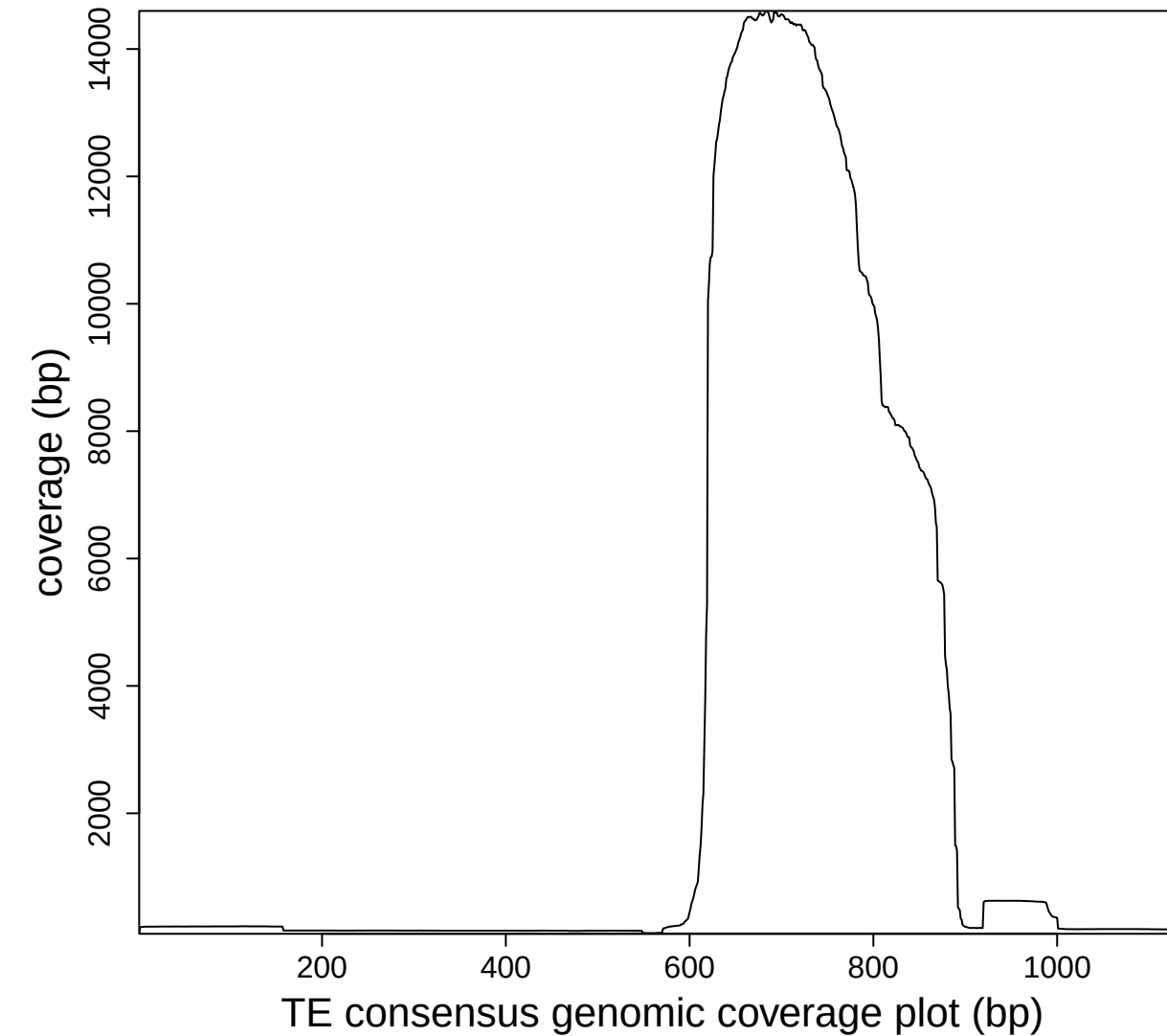
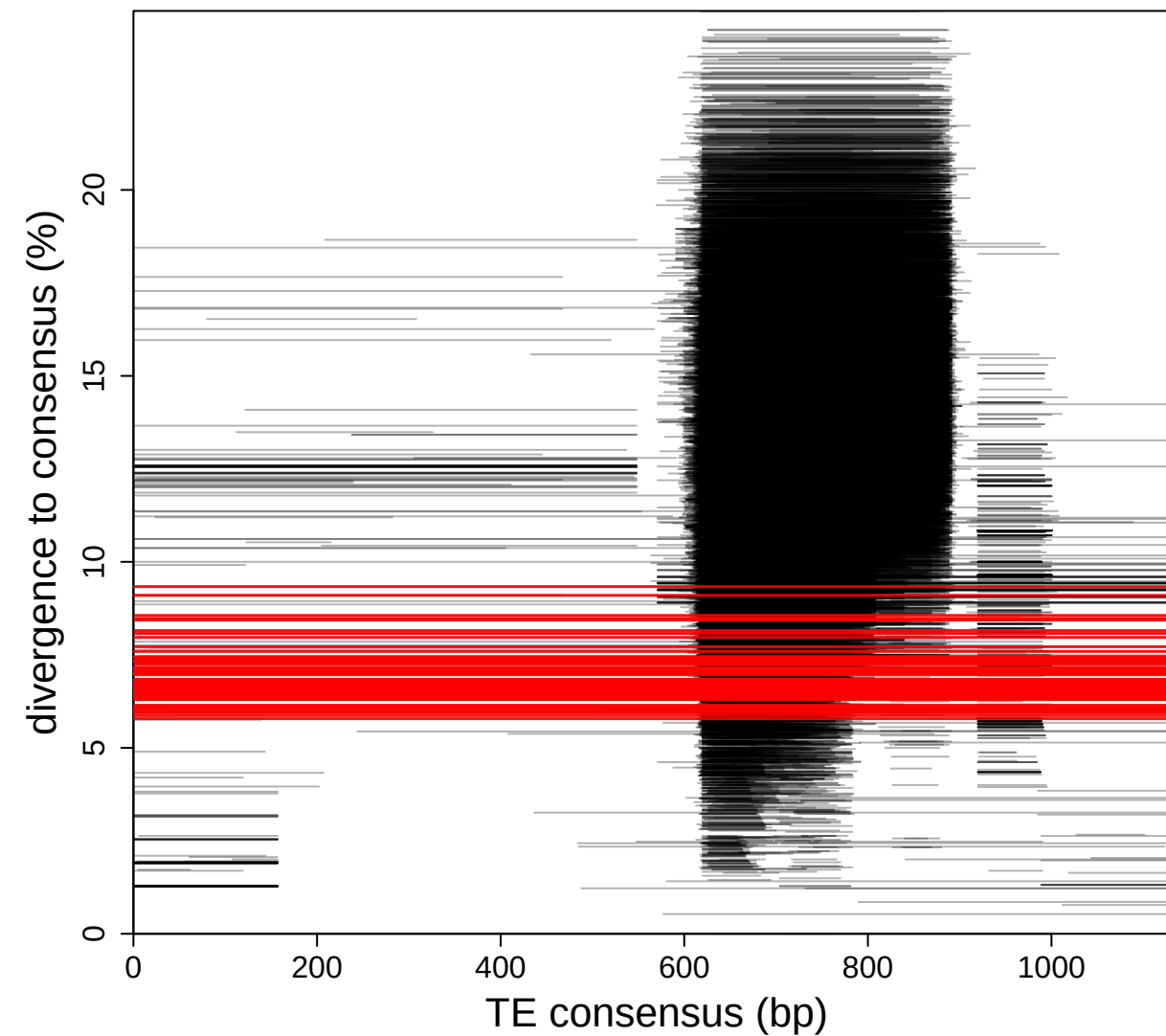
89.fasta.b.bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.1
size: 7451bp; fragments: 260676; full length: 0 (>=6705.9bp)

After TEtrimmer Extended plot Blue lines are boundaries

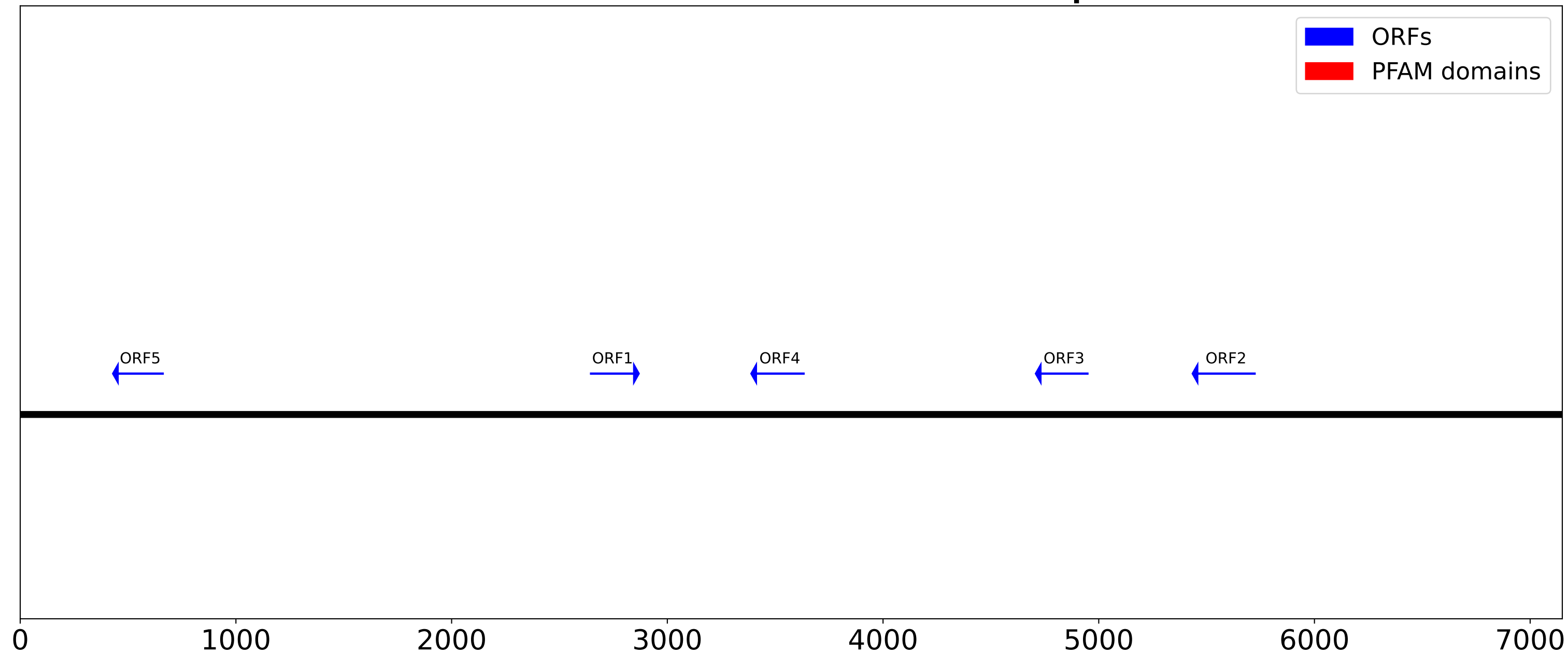


TE: rnd_5_family_13289
size: 1140bp; fragments: 16915; full length: 89 (≥ 1026 bp)

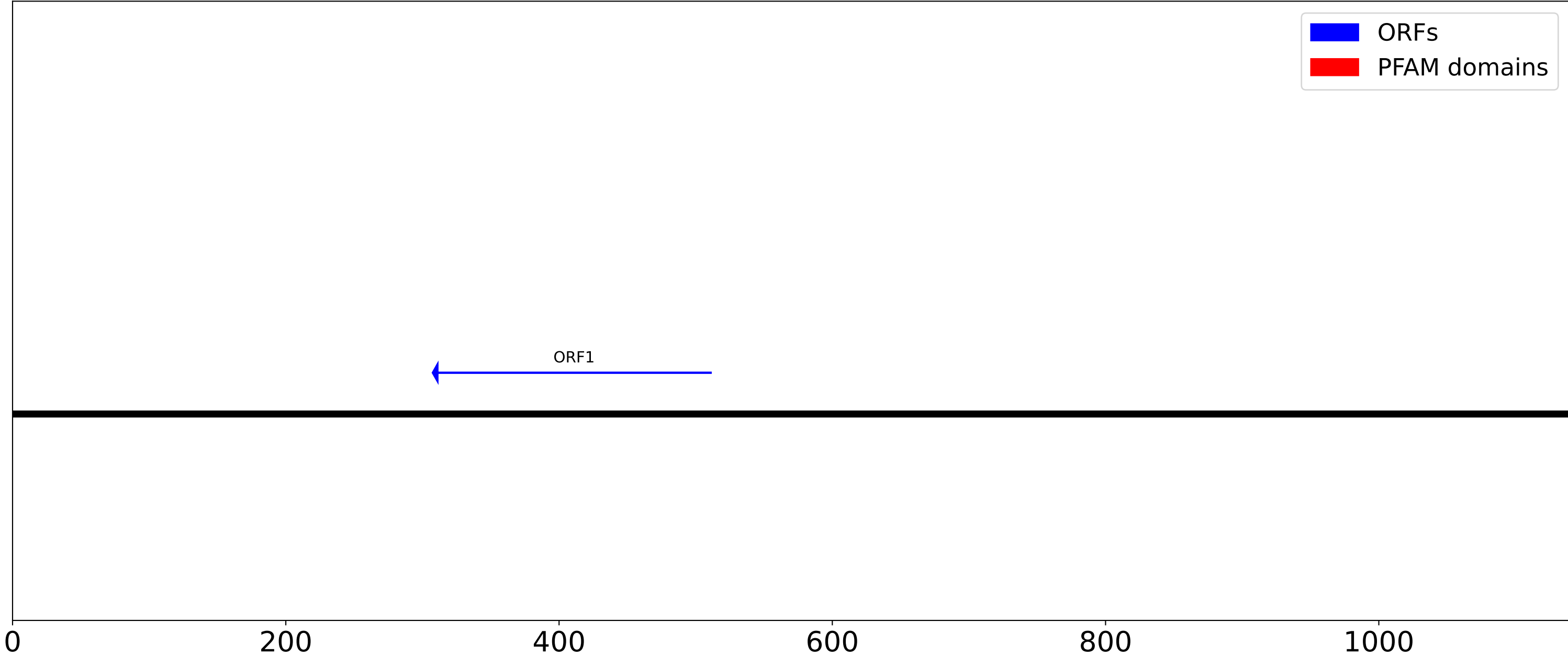
Before TEtrimmer 1140 bp



After TEtrimmer ORF and PFAM domain plot



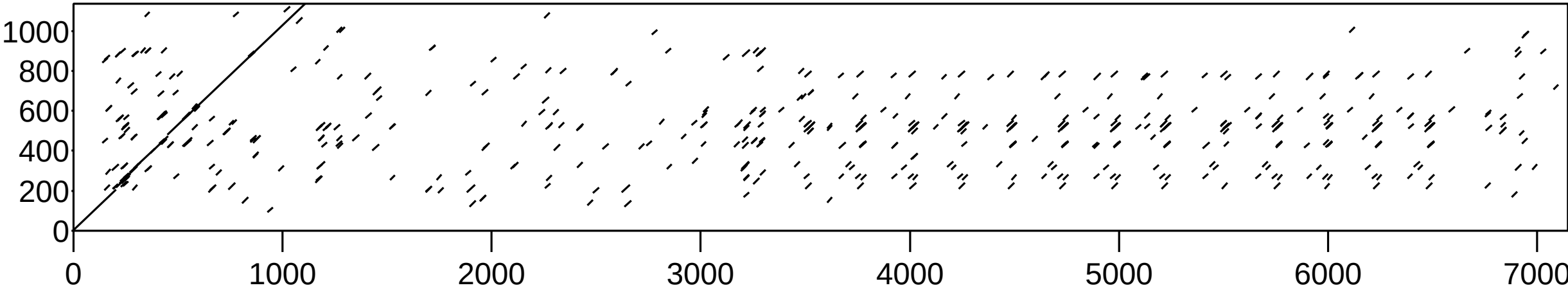
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

nsensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)