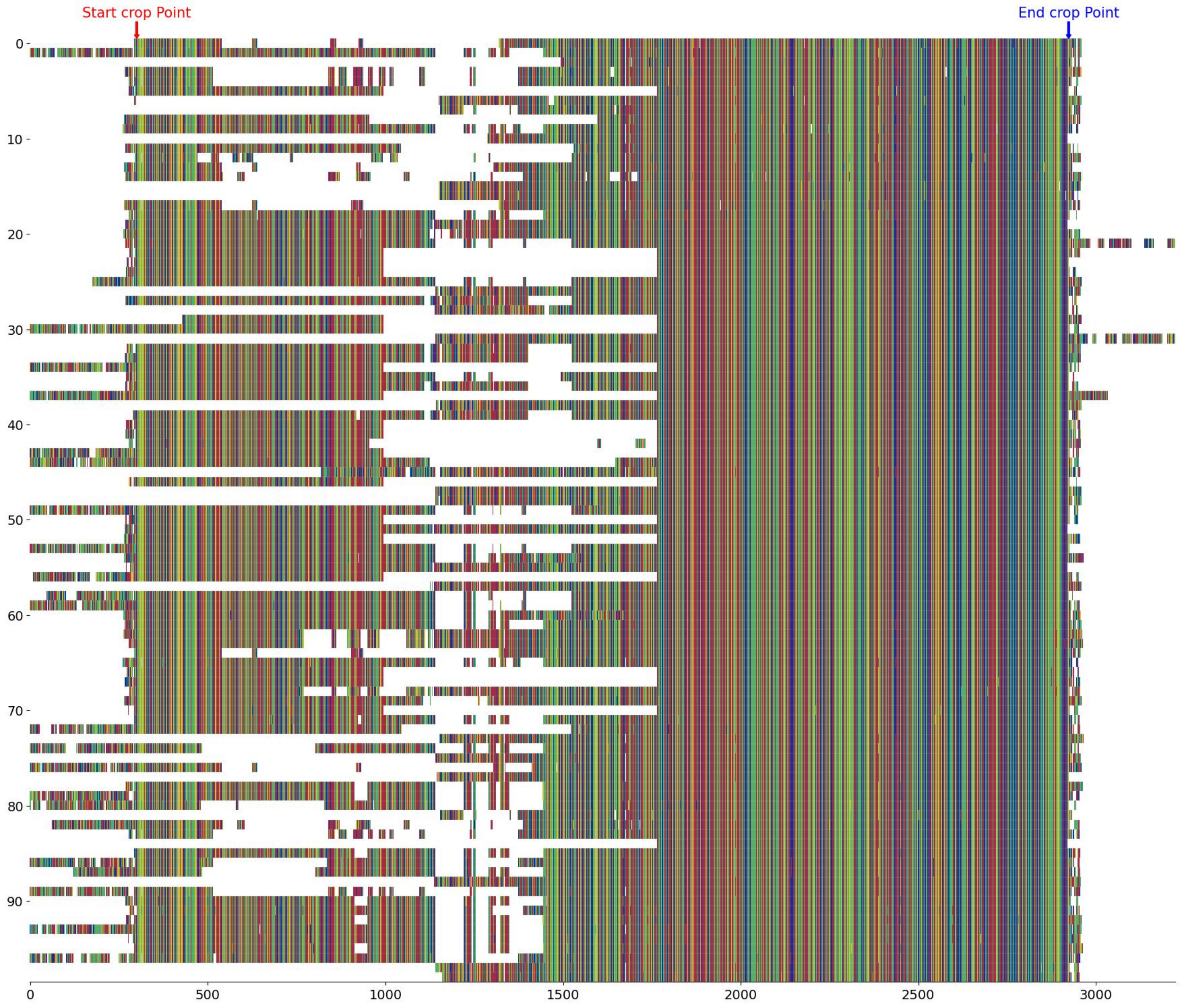
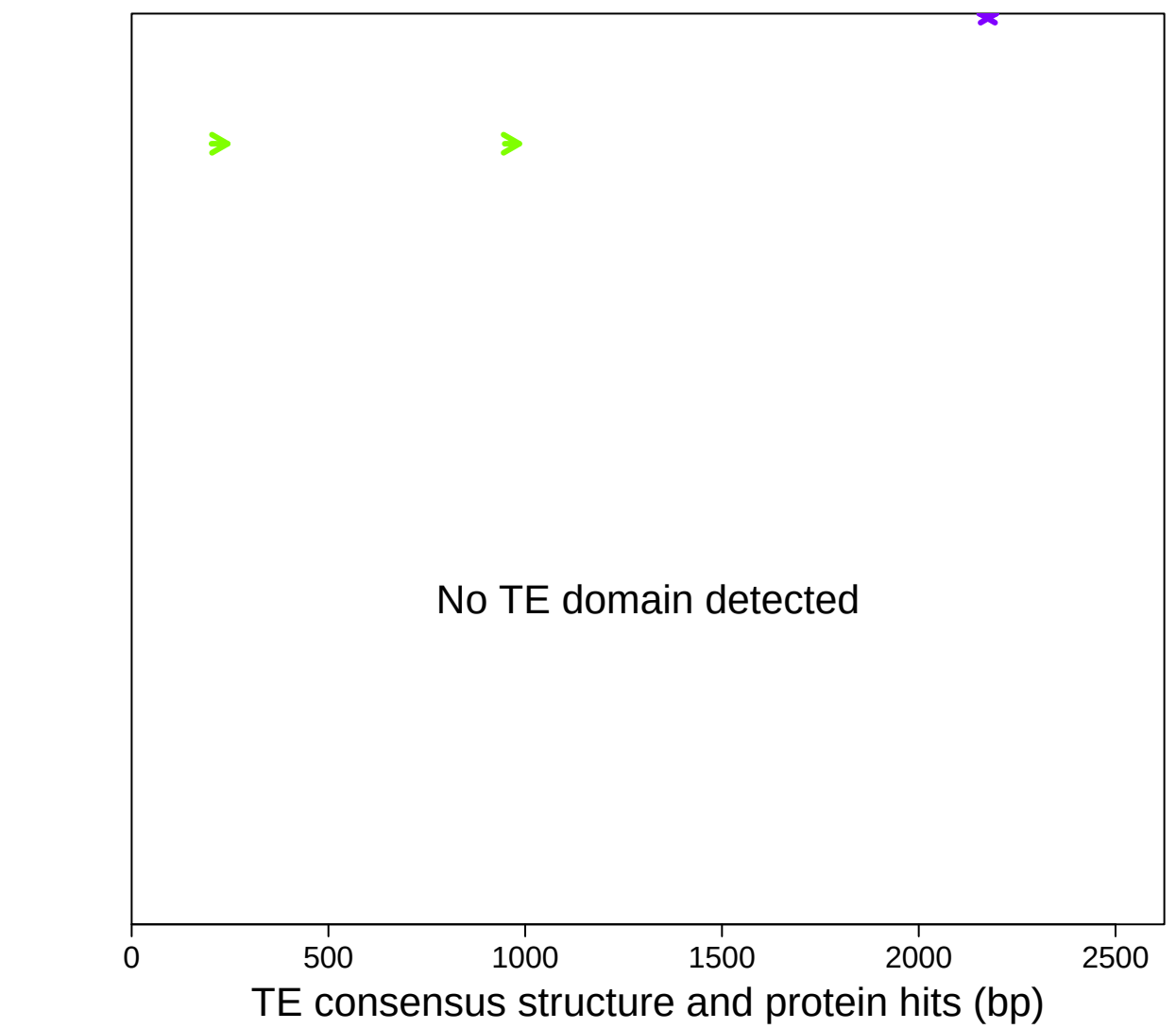
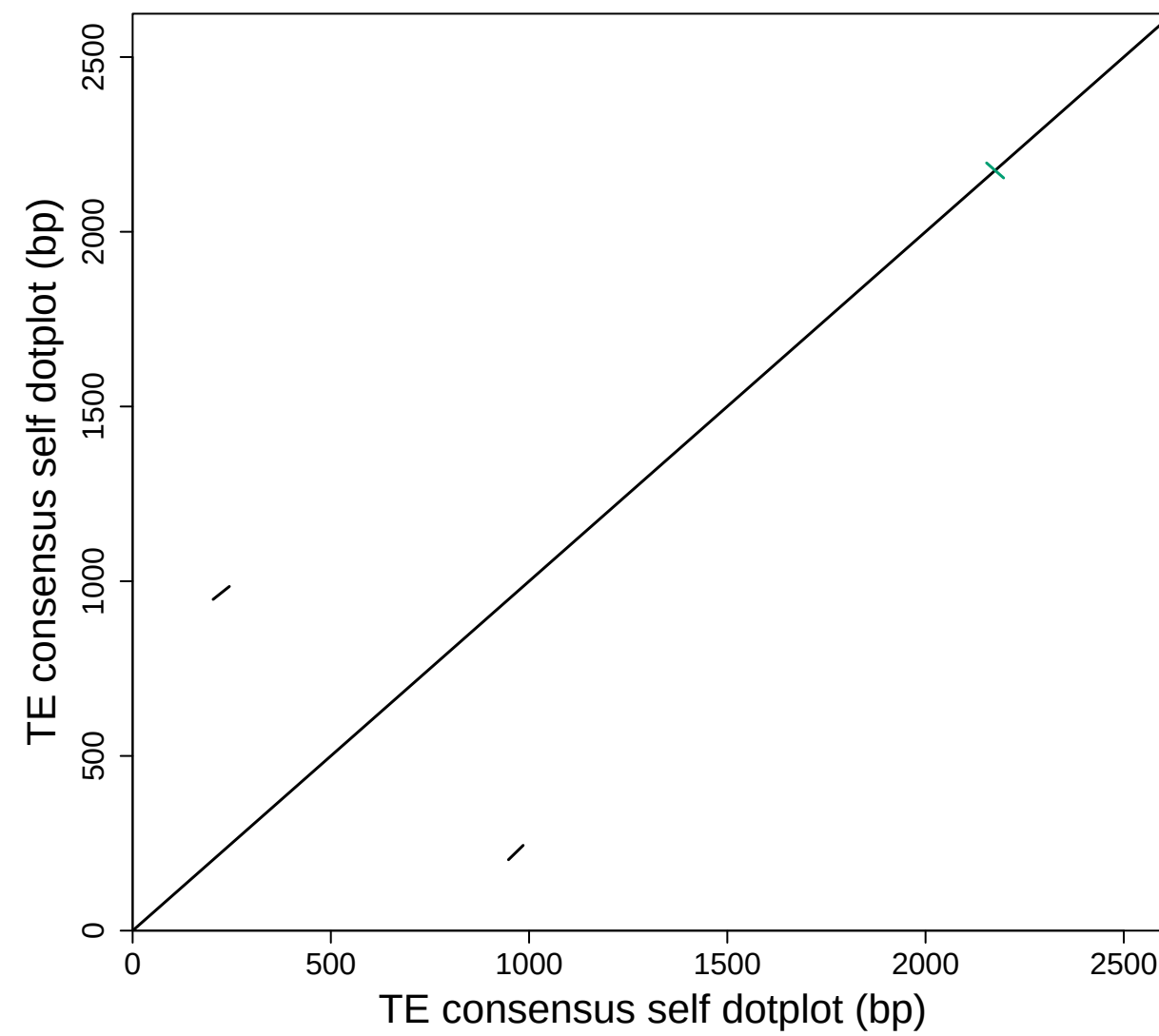
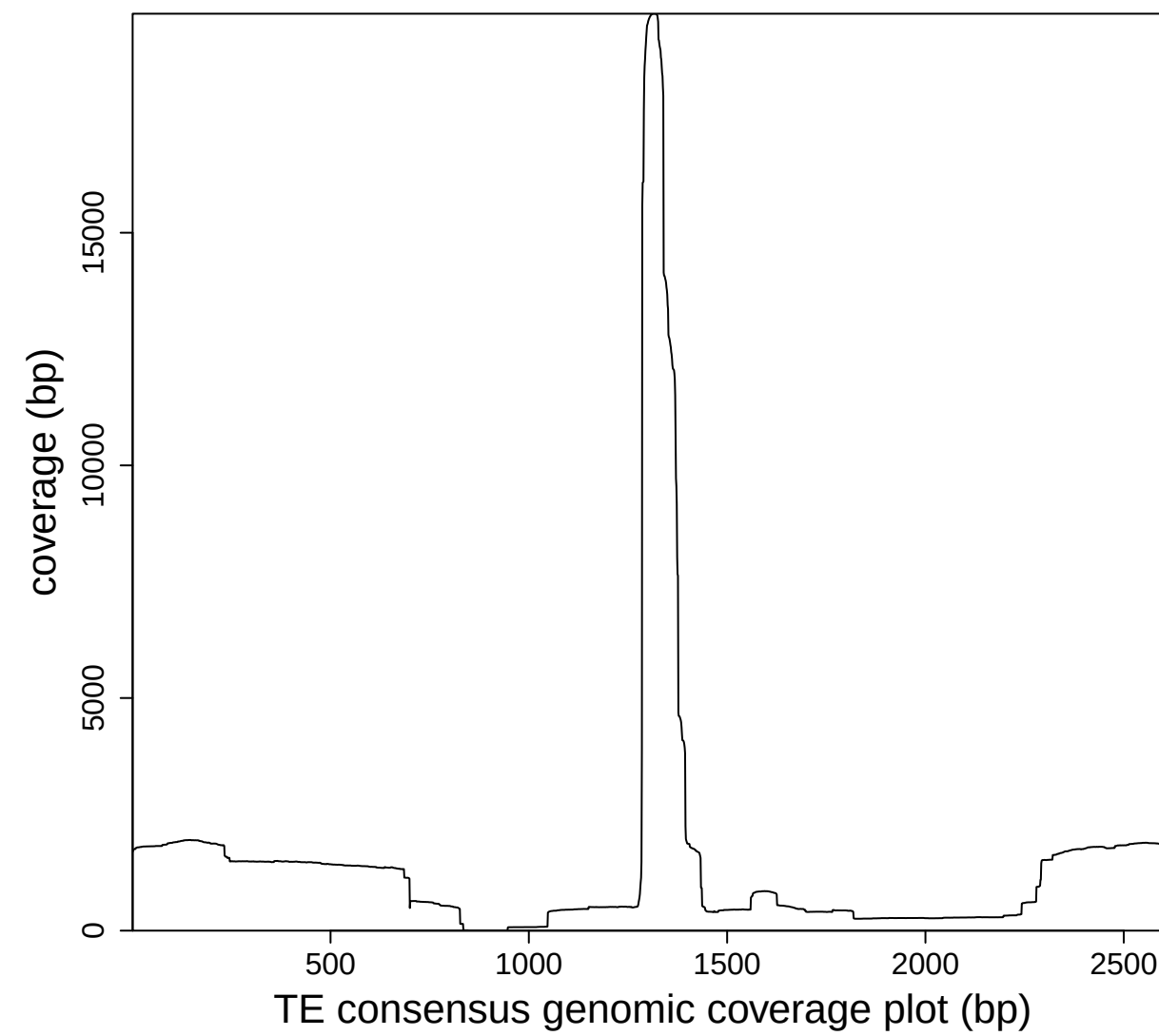
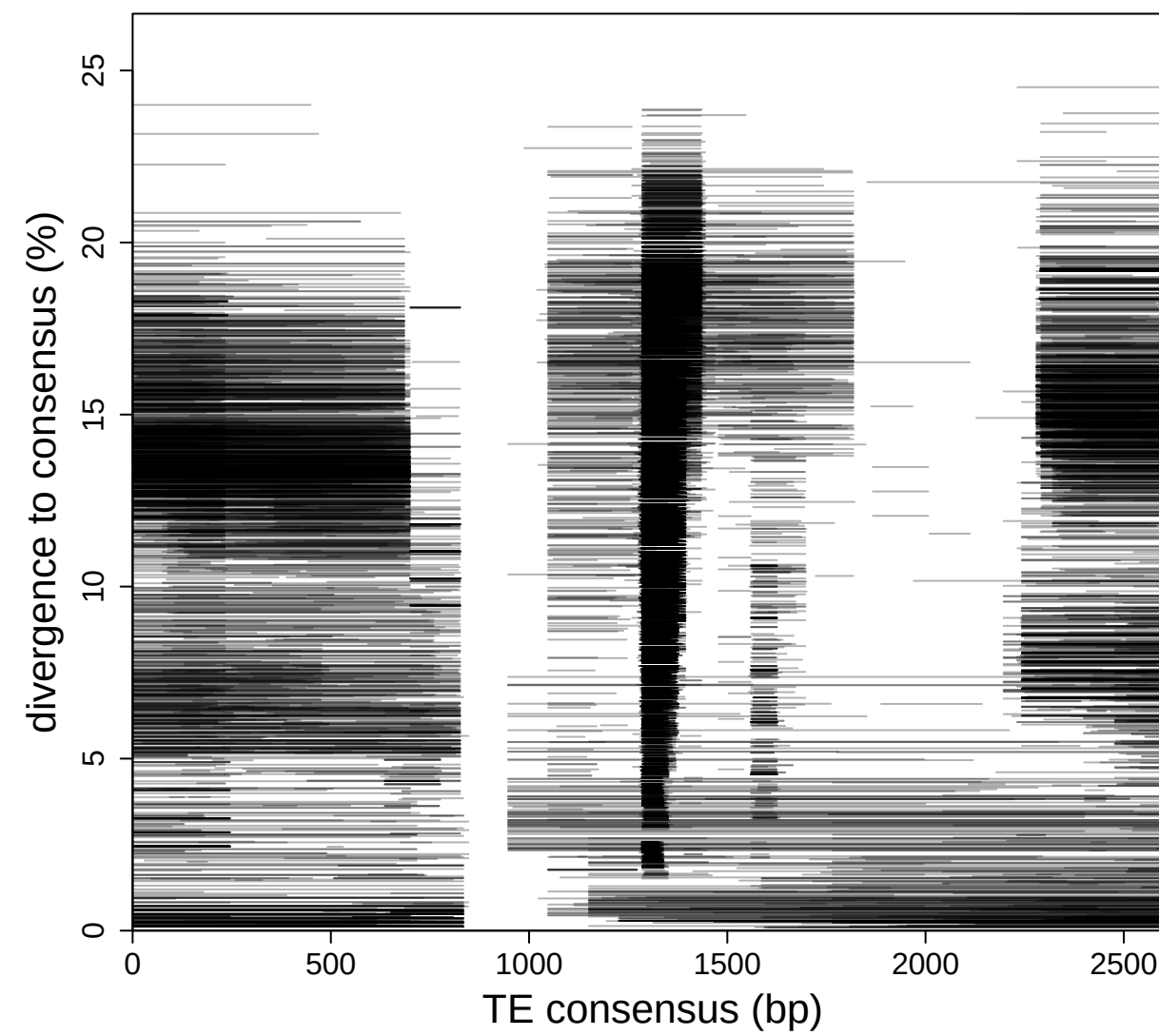






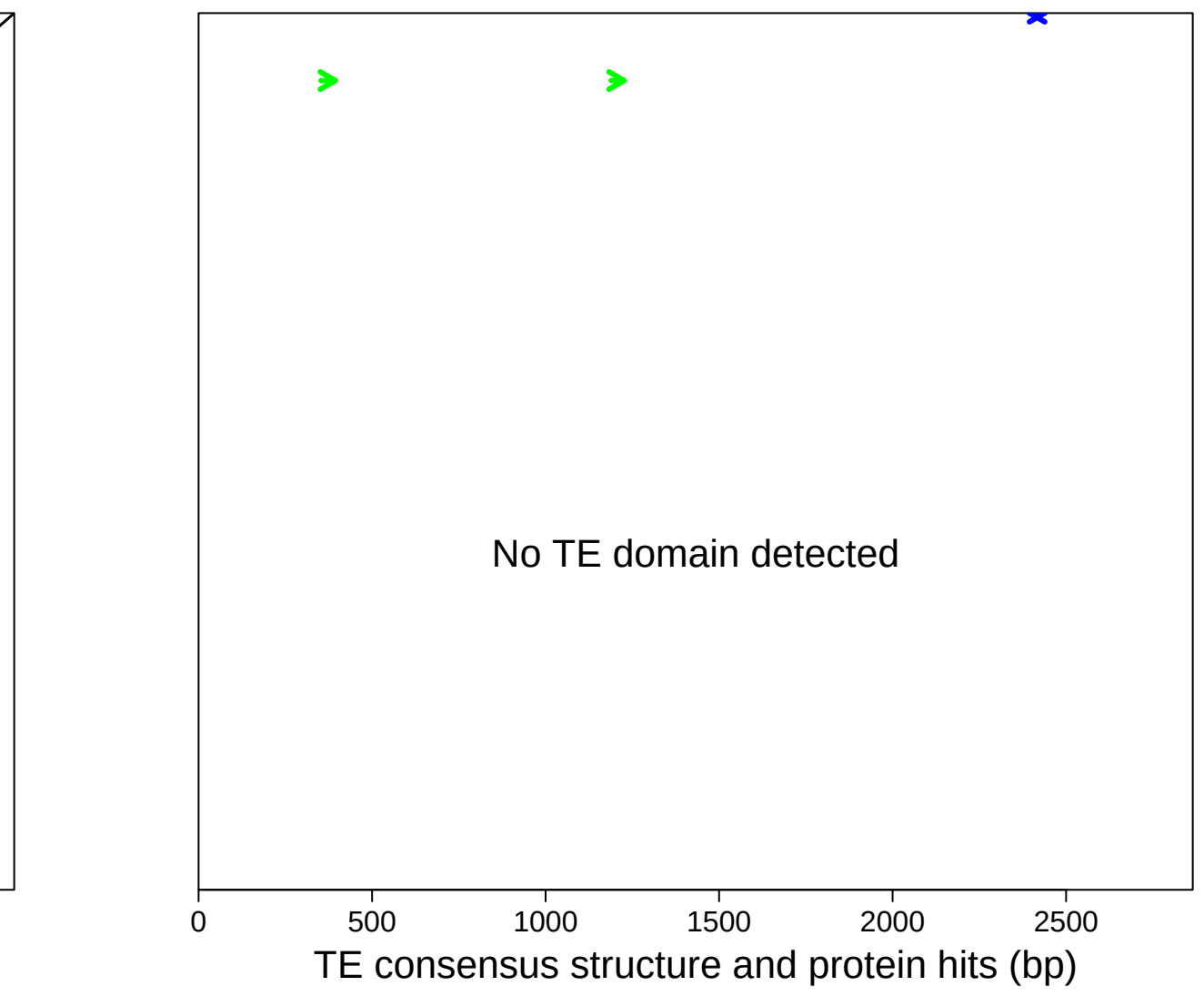
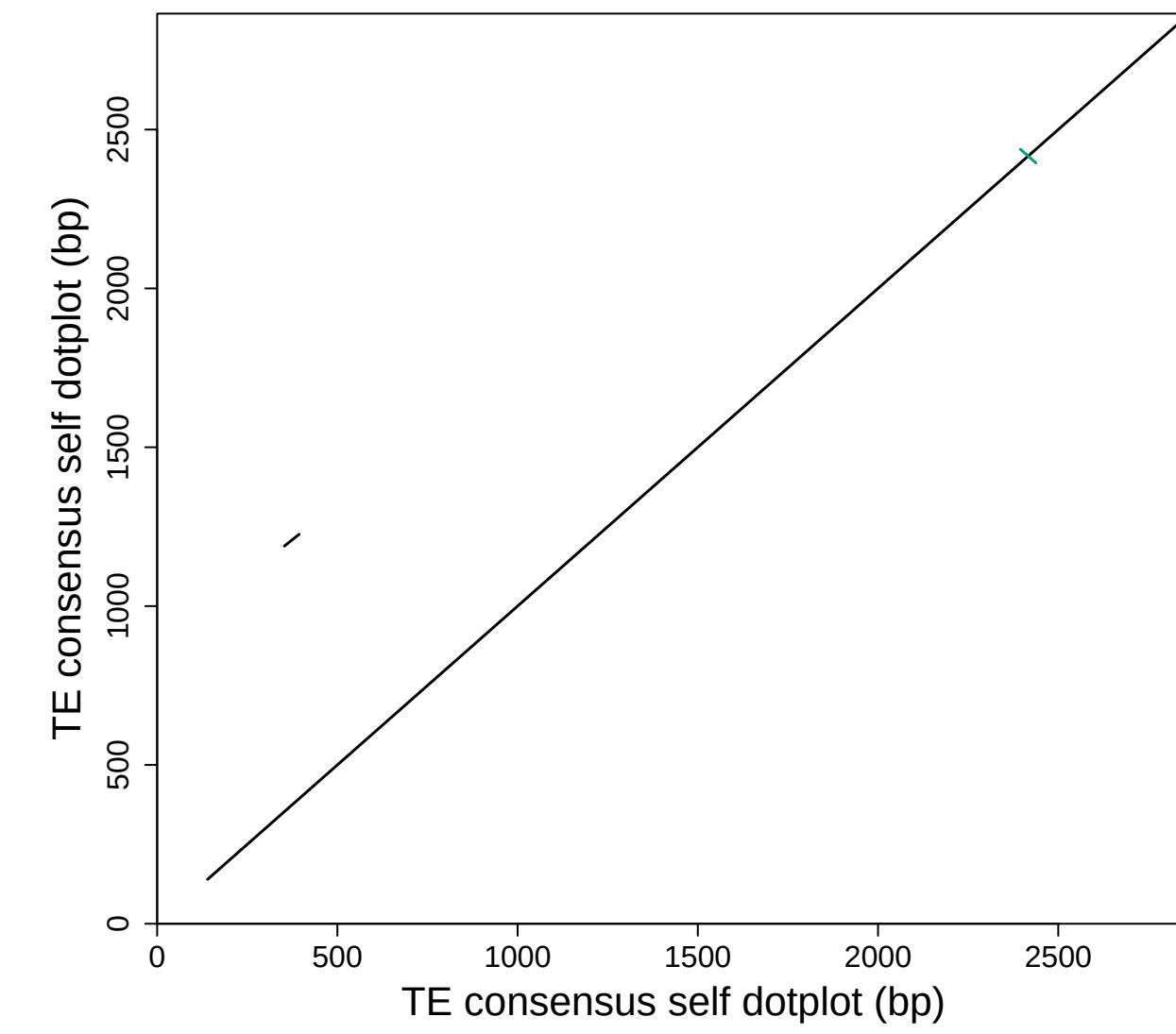
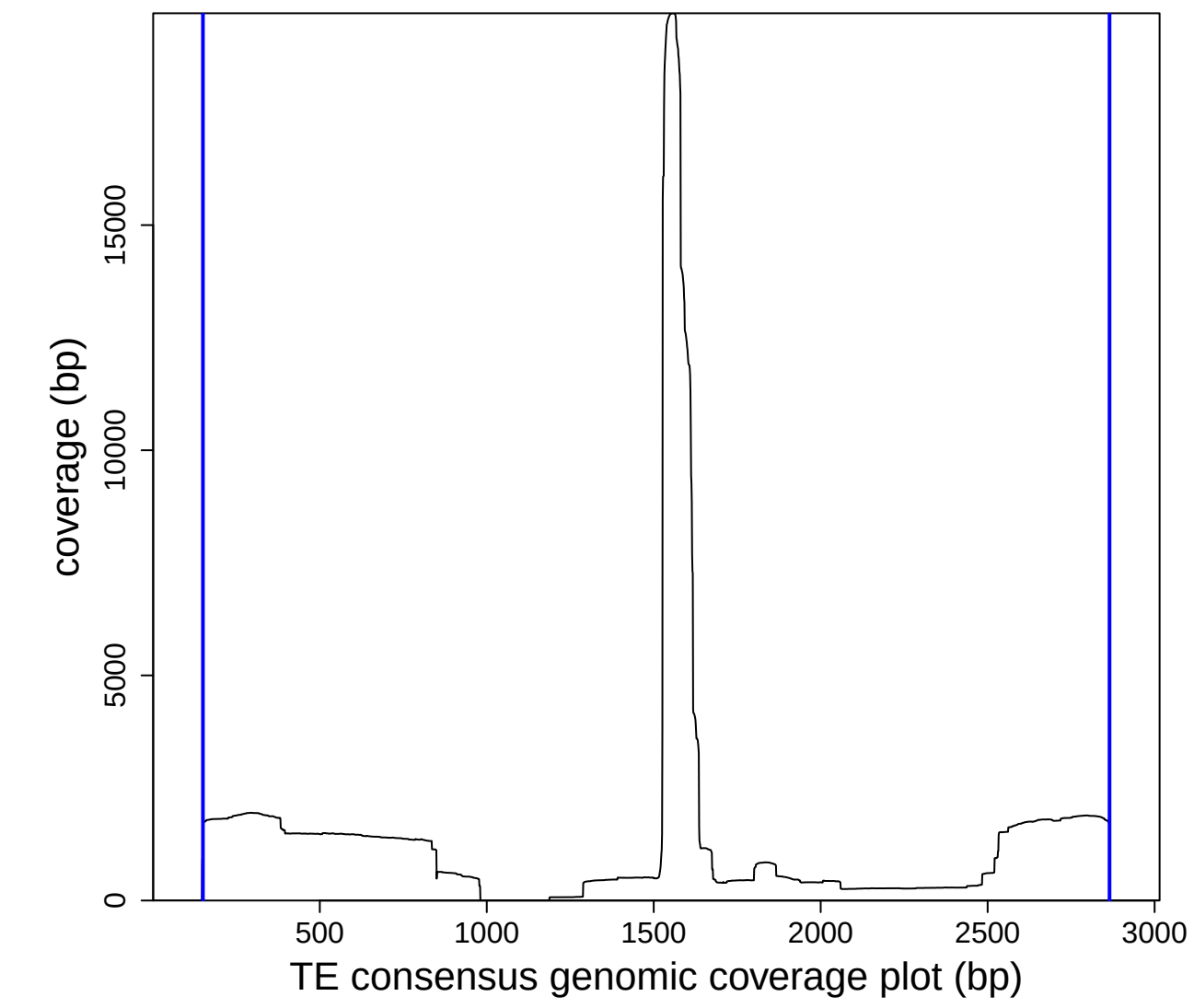
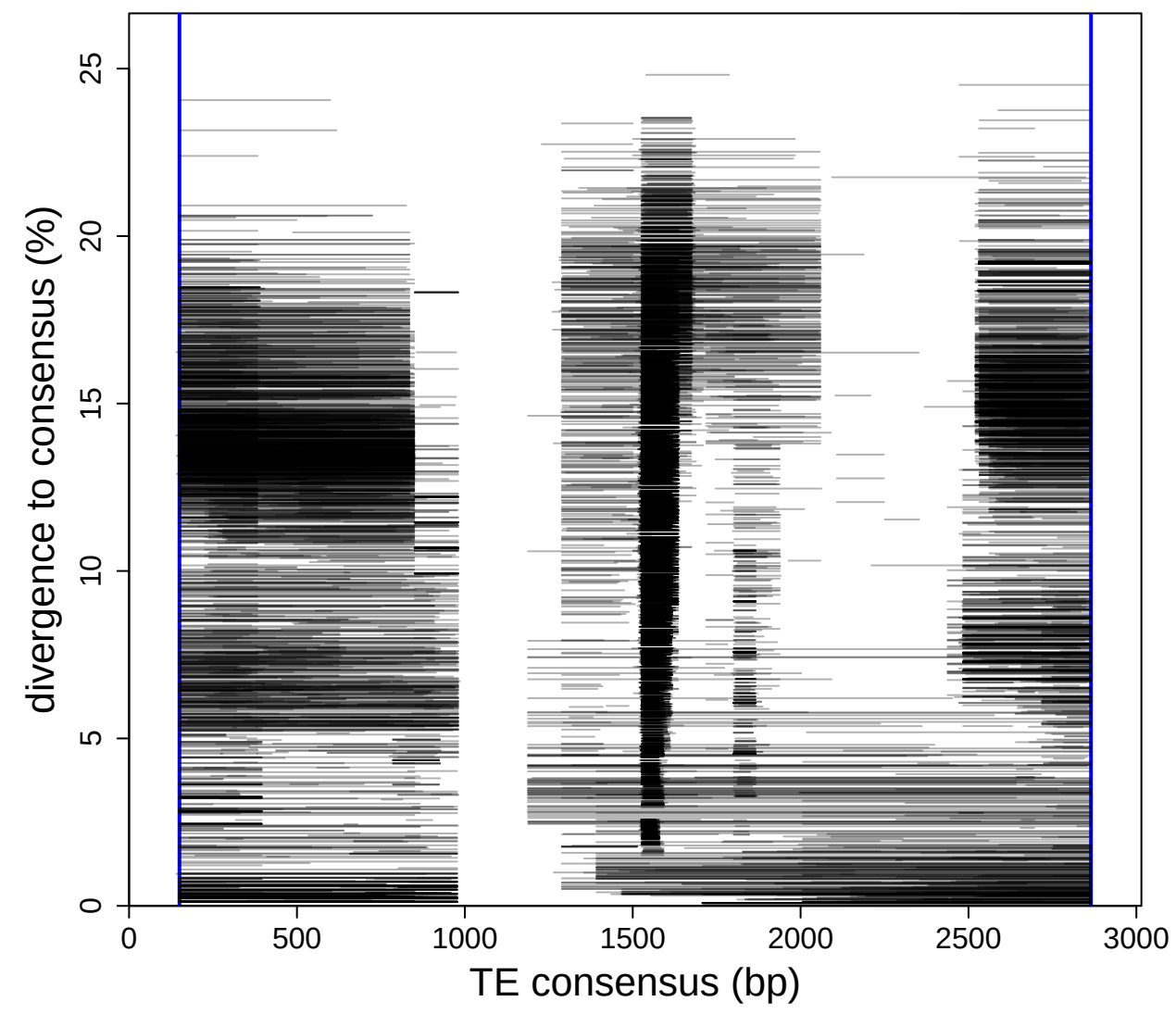
l.b.bed uf.bed g_1.bed fm_1.bed 0 0 bcln.fa aln.fa cl.fa gs.fa
size: 2624bp; fragments: 24654; full length: 0 (>=2361.6bp)

After TEtrimmer 2624 bp



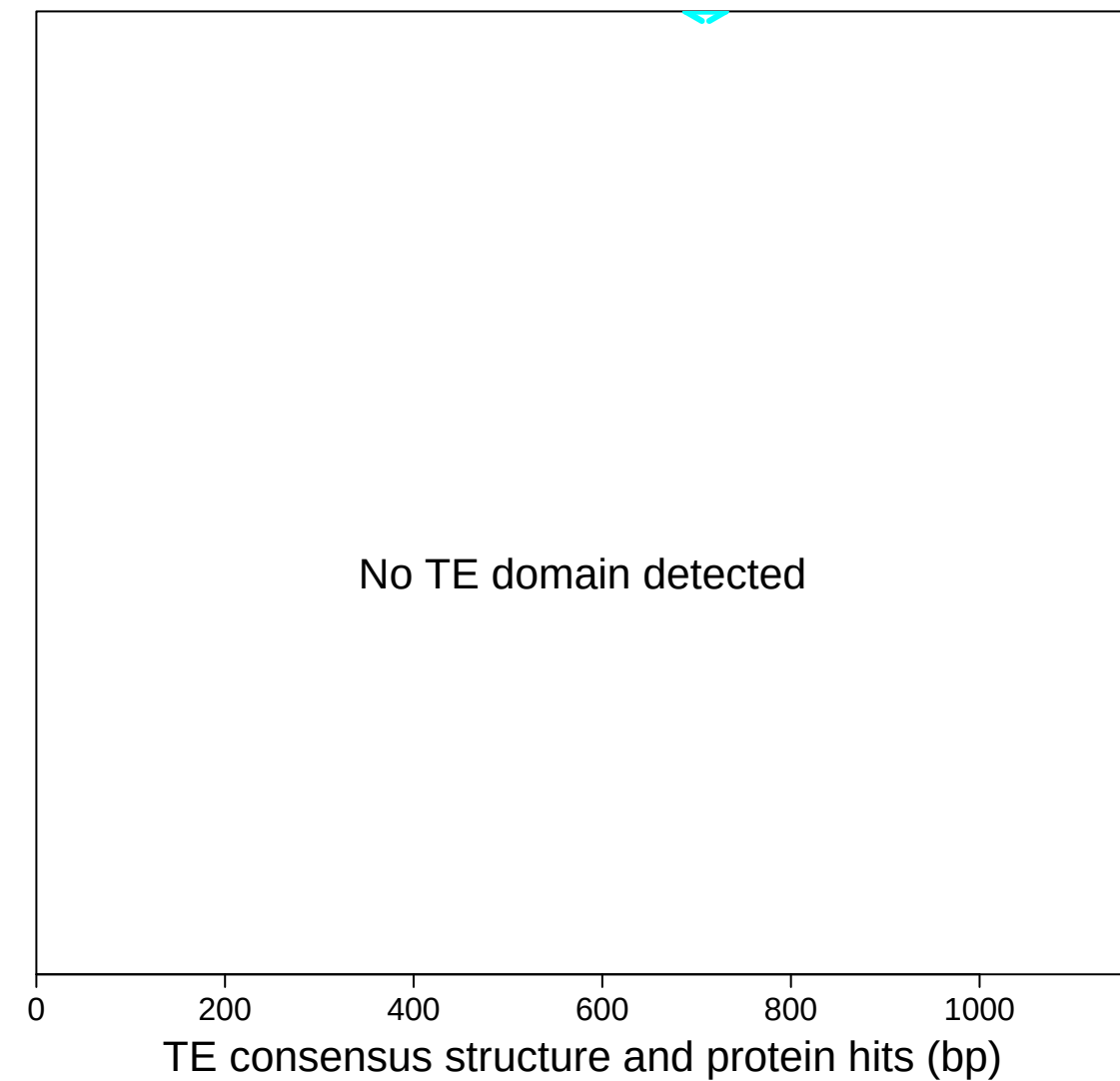
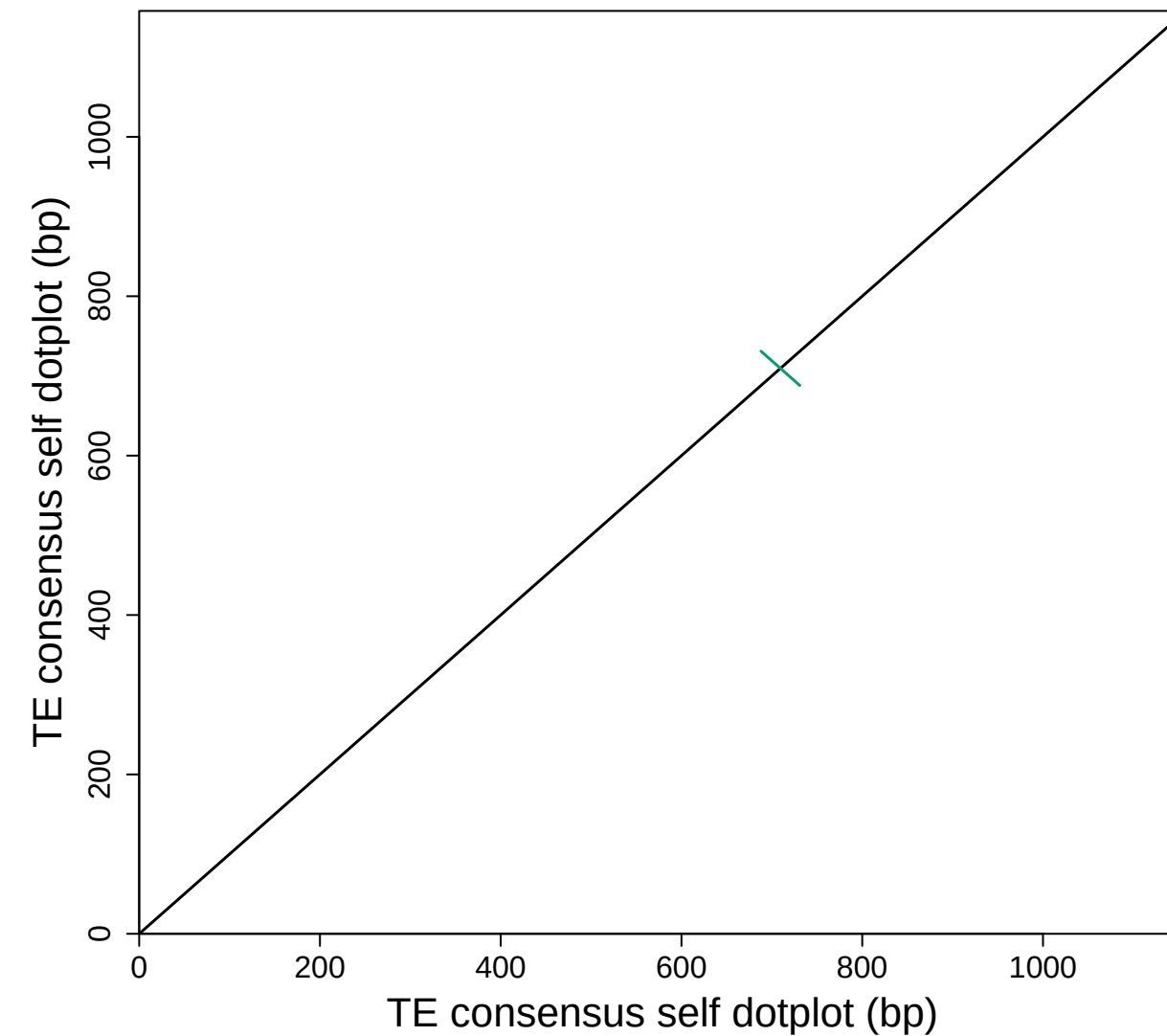
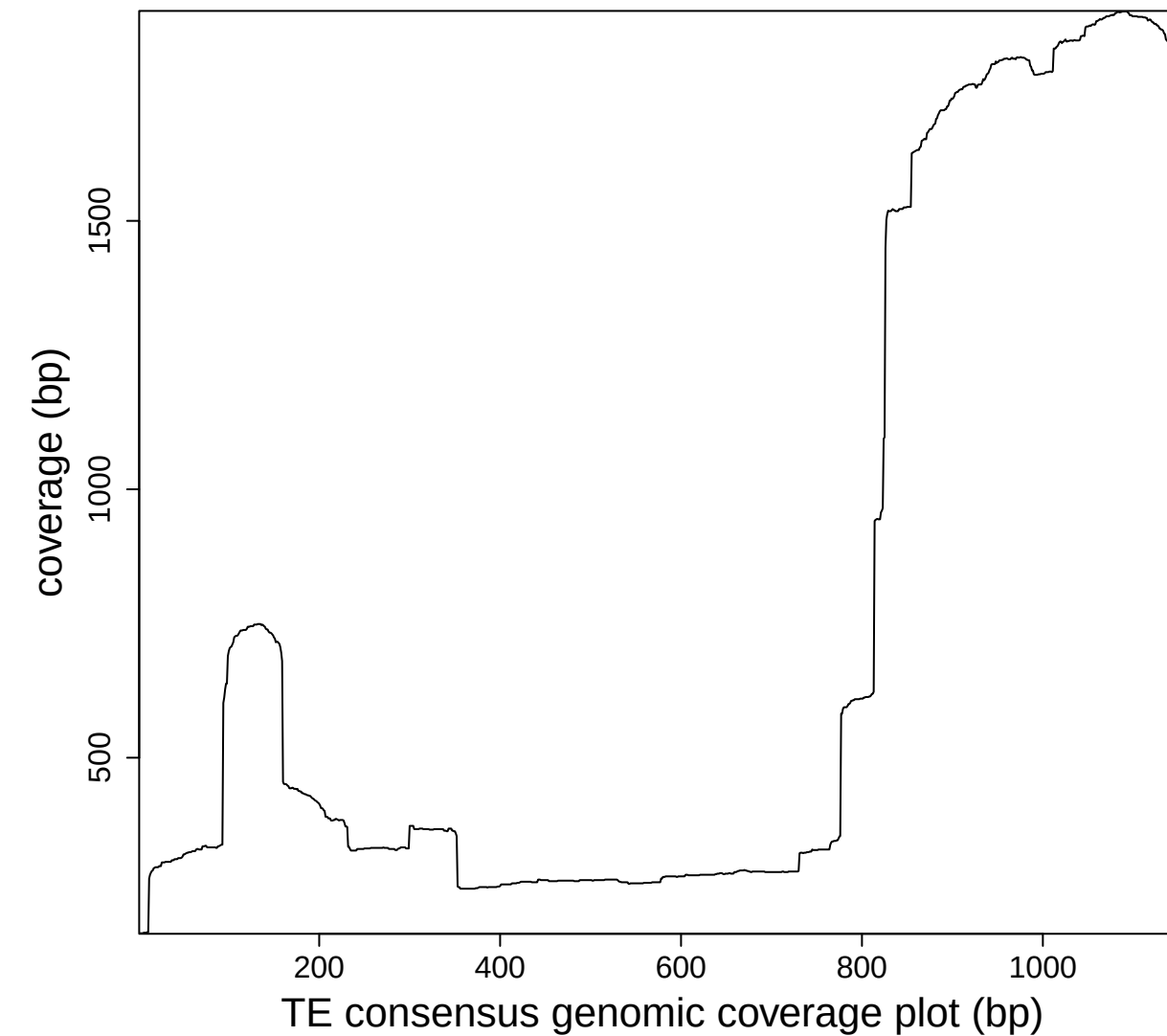
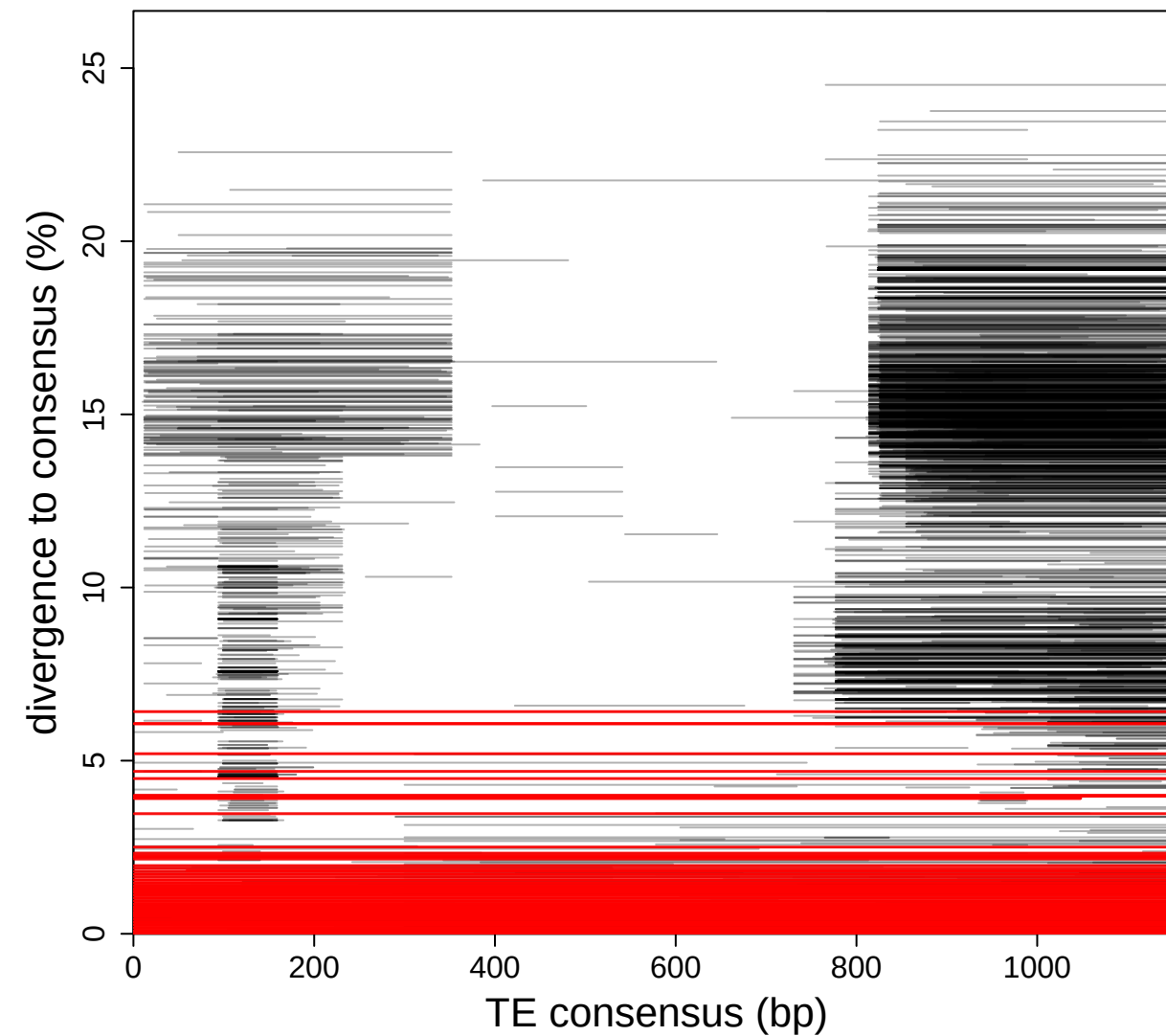
6.fasta.b.bed uf.bed g 1.bed fm 1.bed 0 0 bcln.fa aln.fa cl.fa
size: 3015bp; fragments: 24642; full length: 0 (>=2713.5bp)

After TEtrimmer Extended plot Blue lines are boundaries

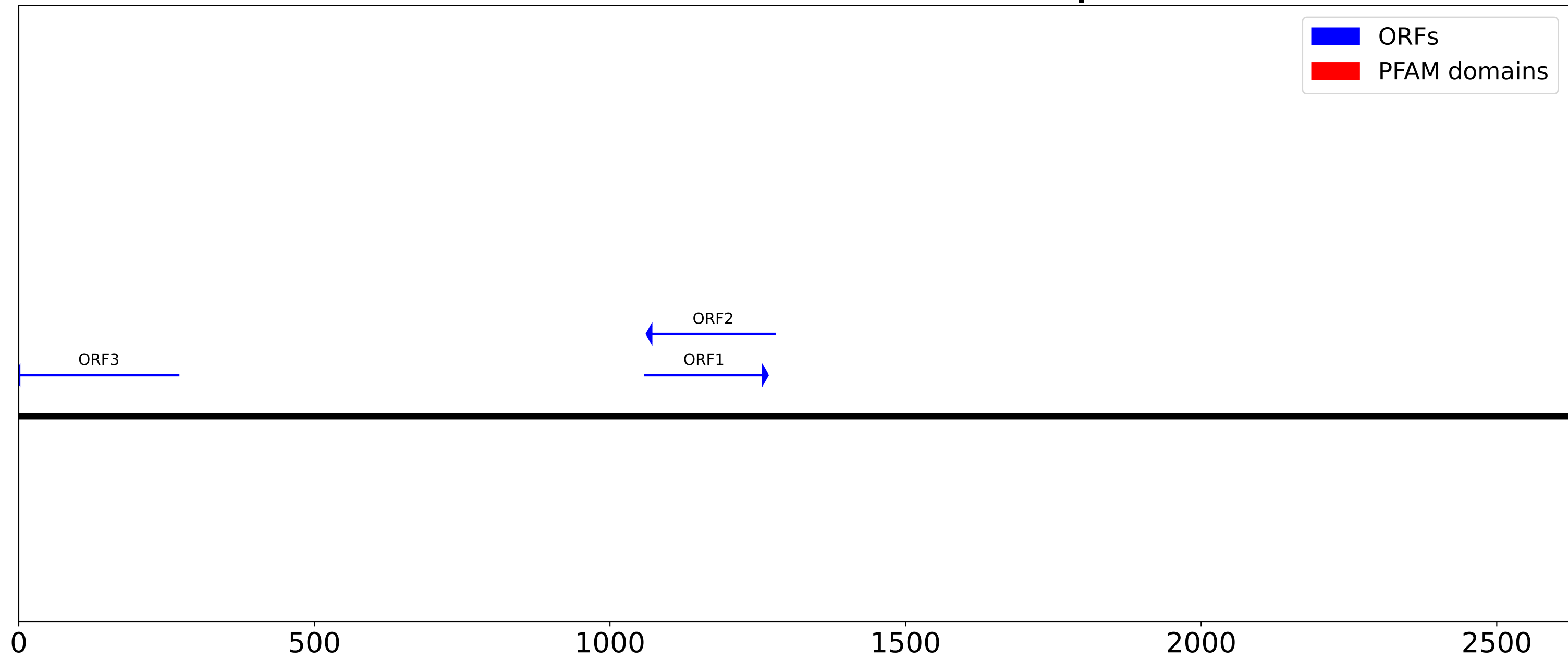


TE: rnd_1_family_566
size: 1158bp; fragments: 2620; full length: 149 (≥ 1042.2 bp)

Before TEtrimmer 1158 bp



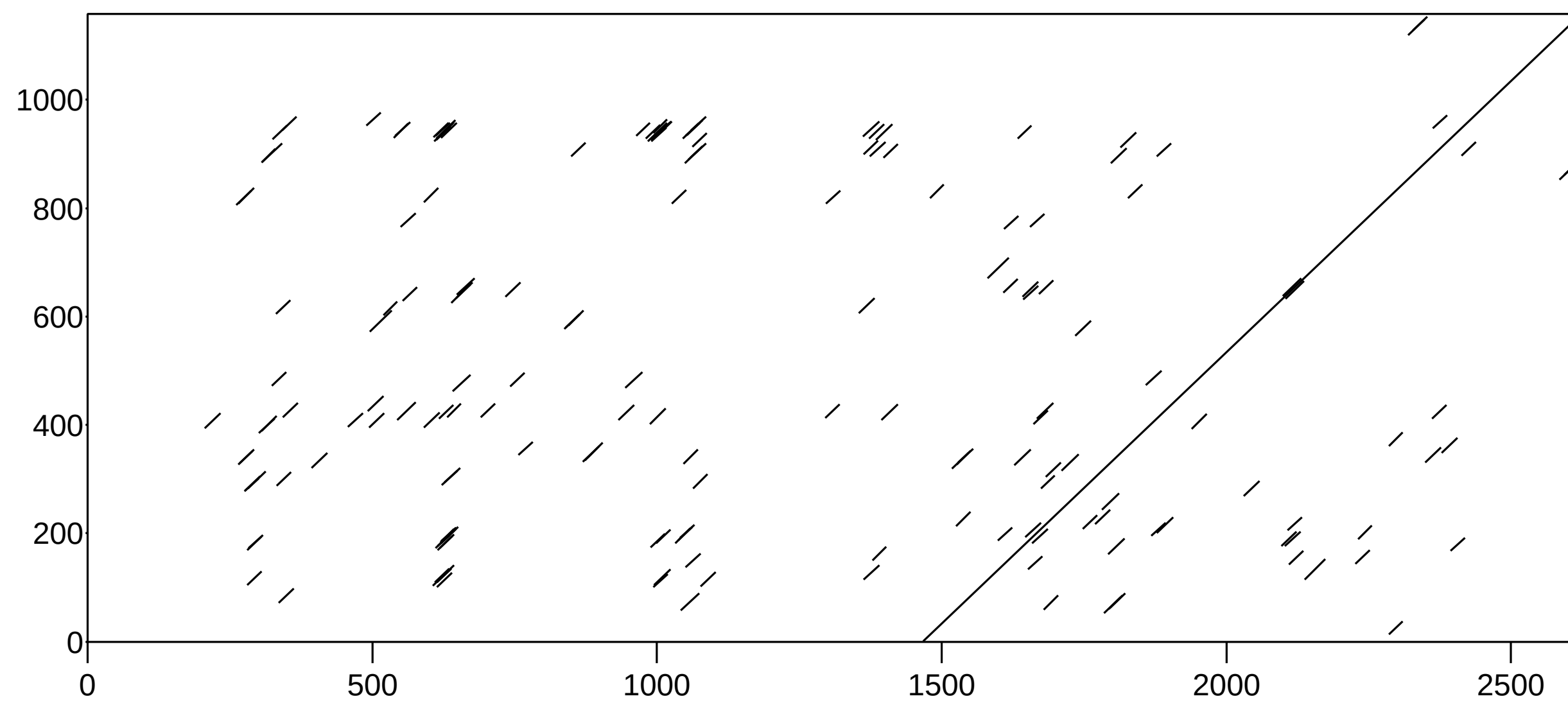
After TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)