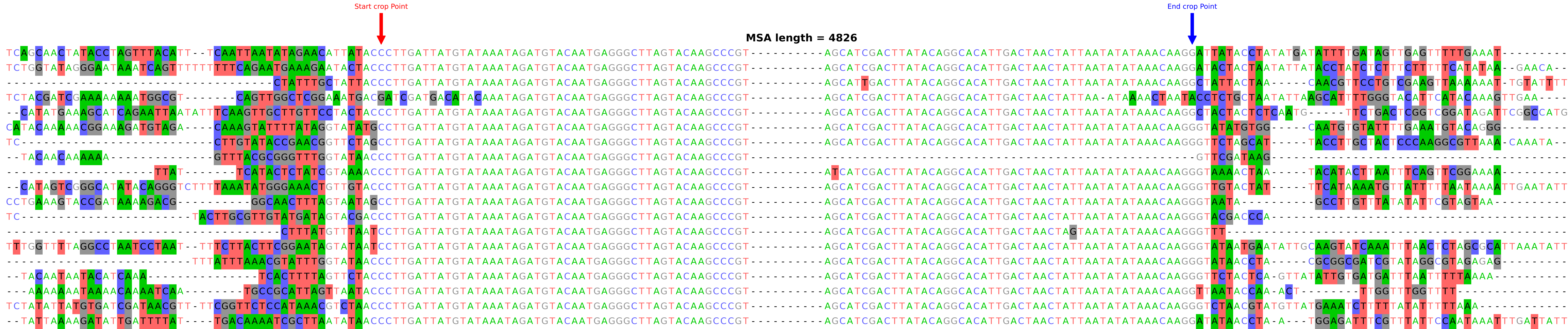
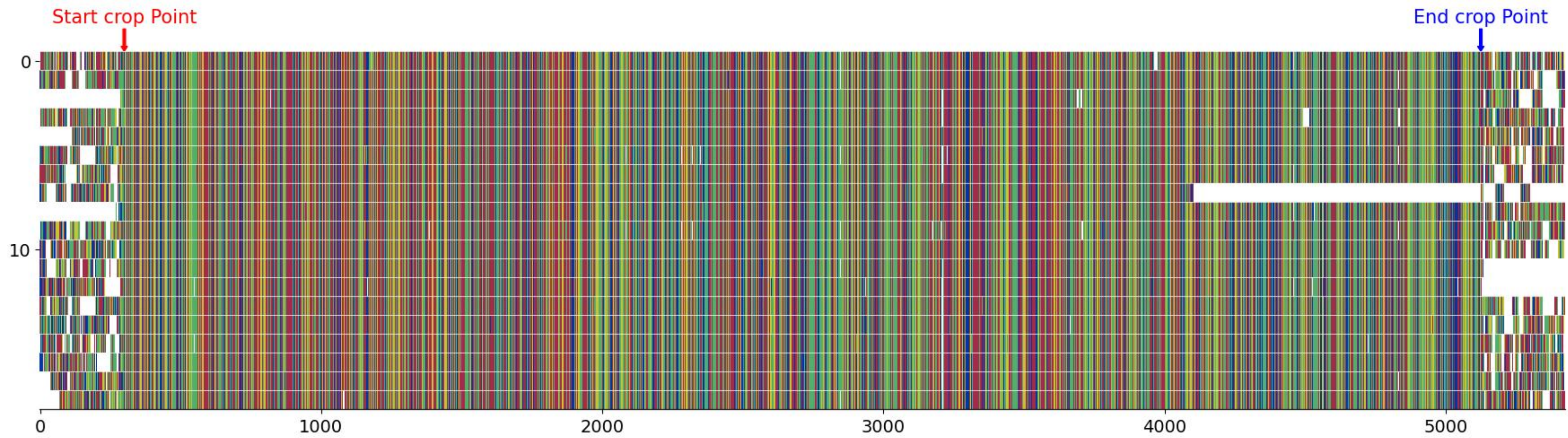




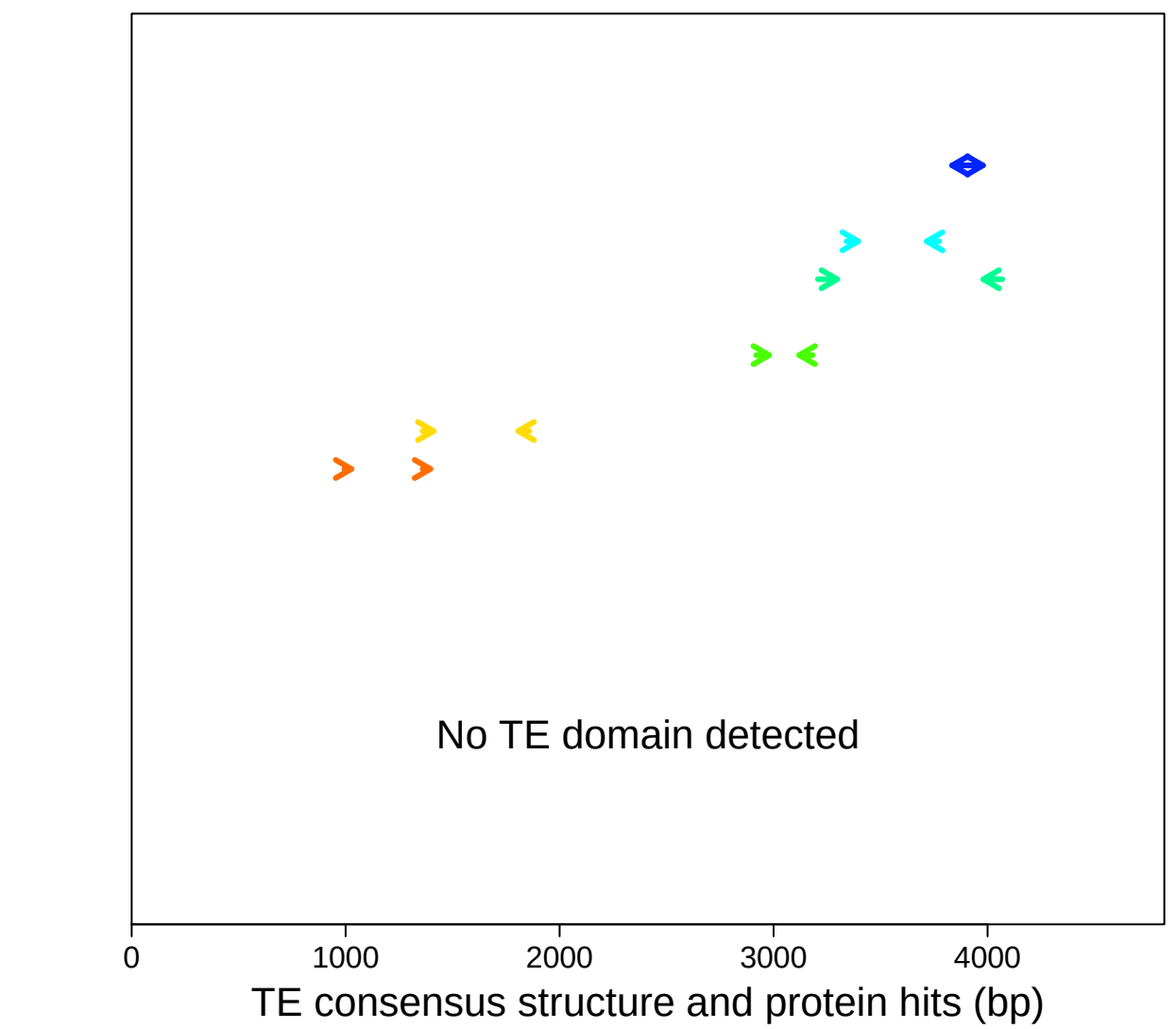
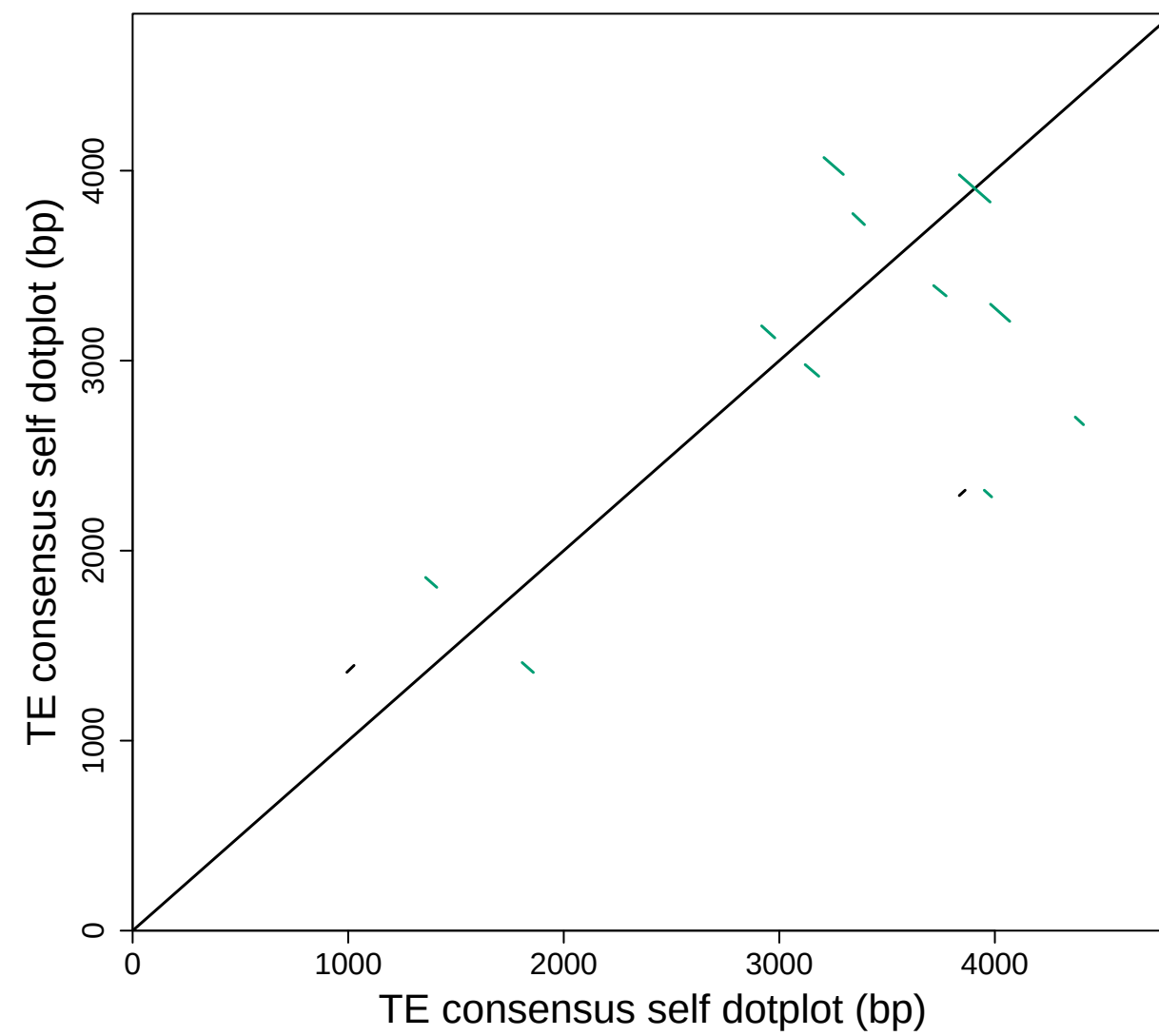
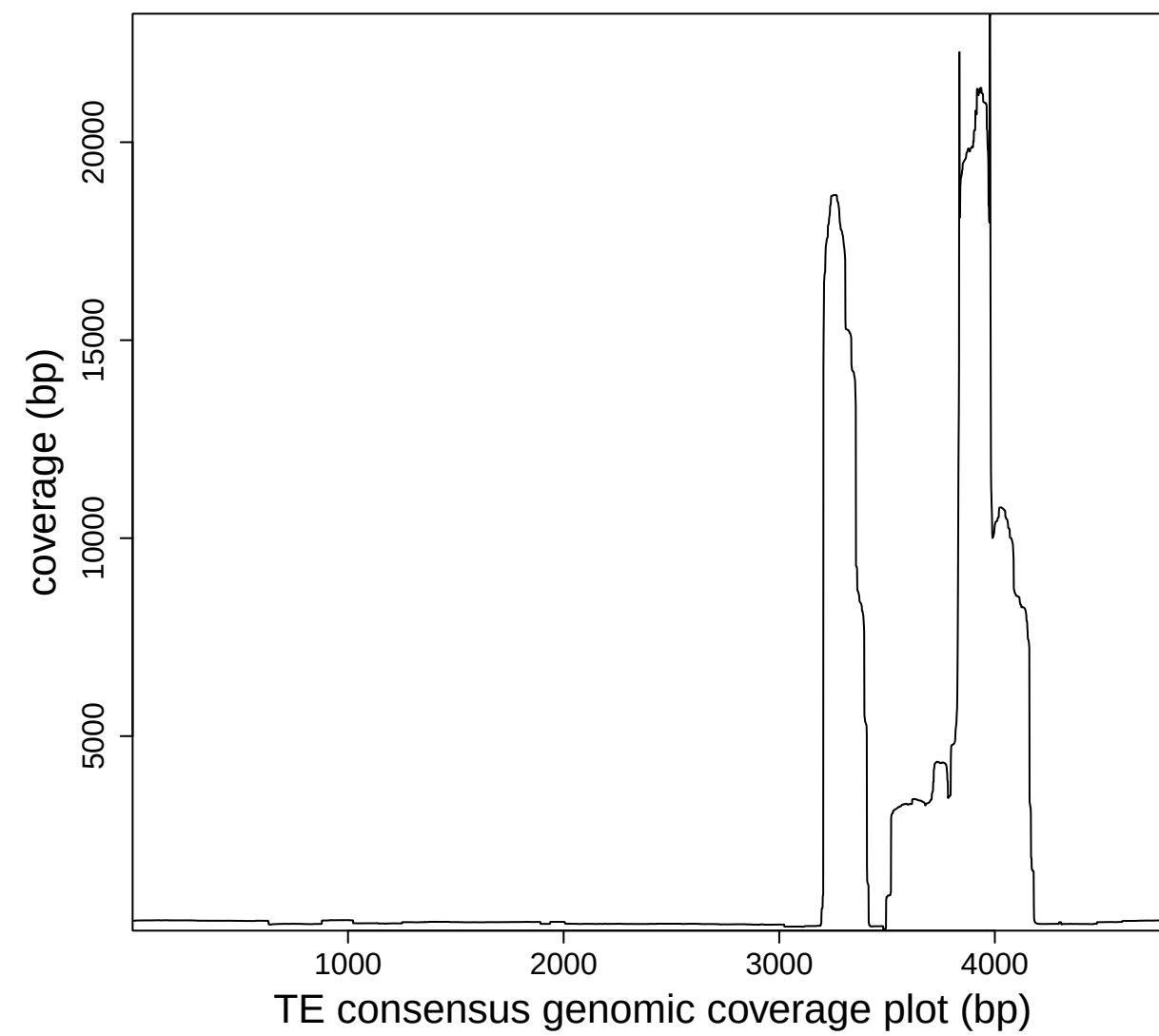
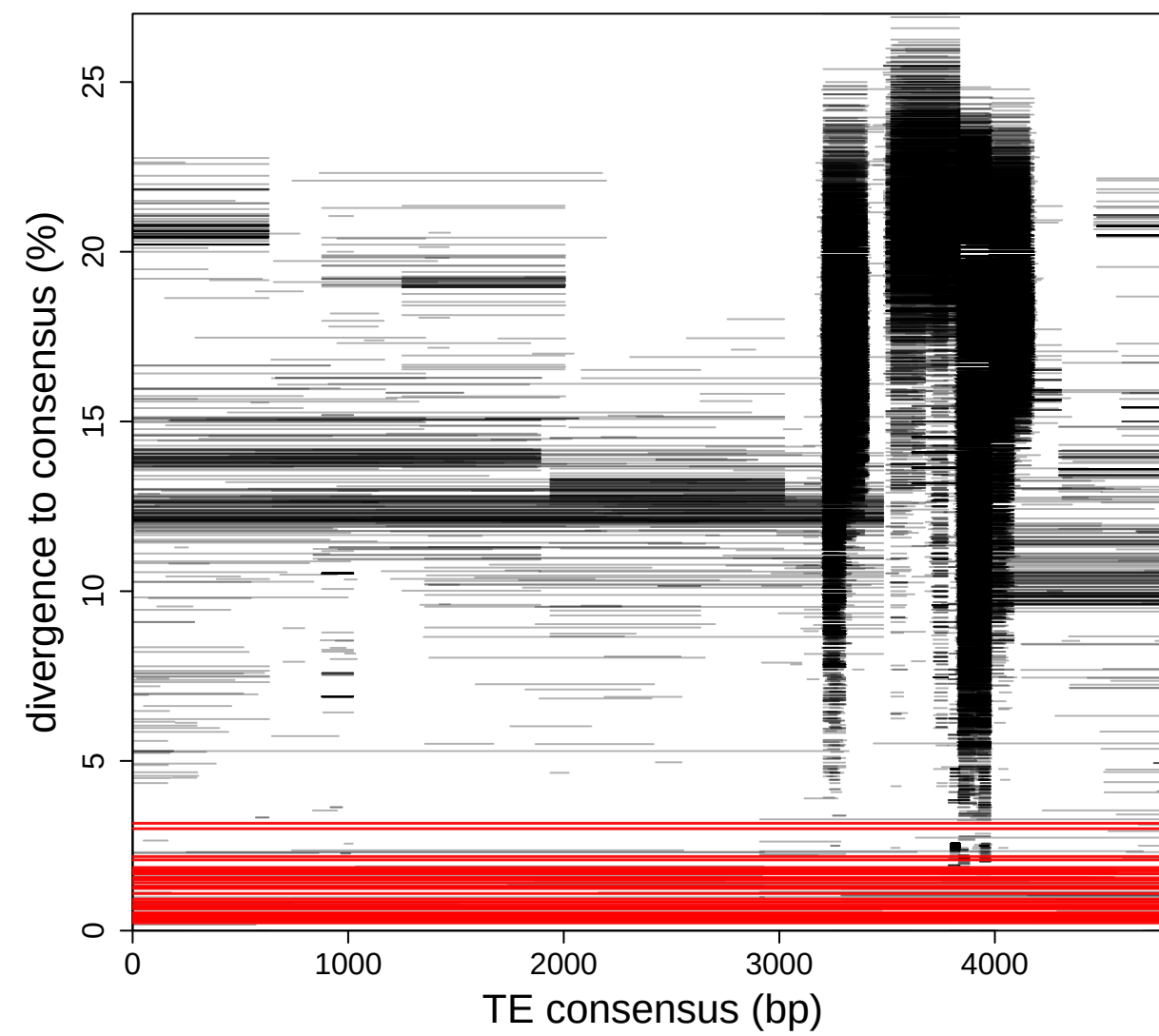
Start crop Point

End crop Point



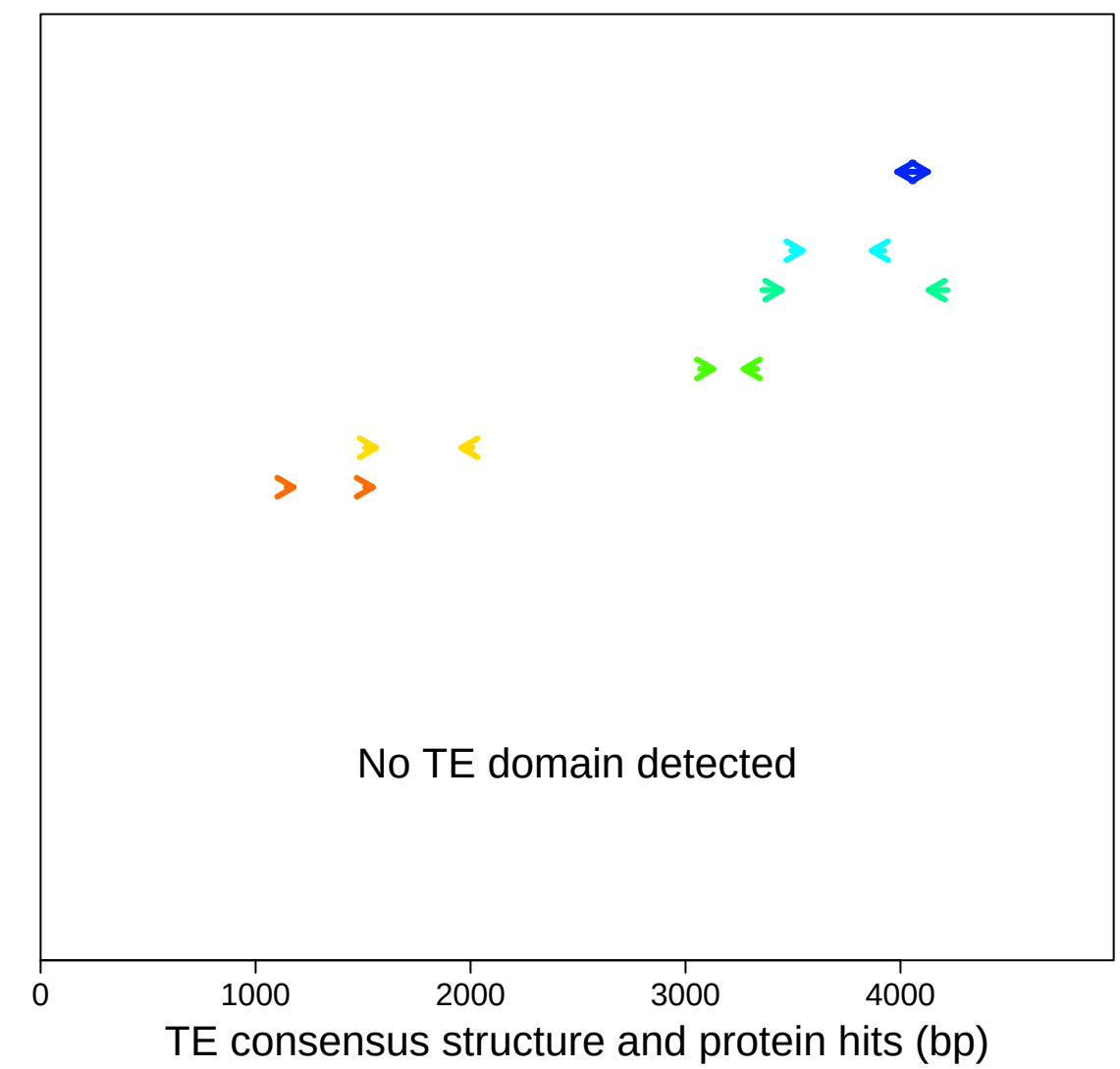
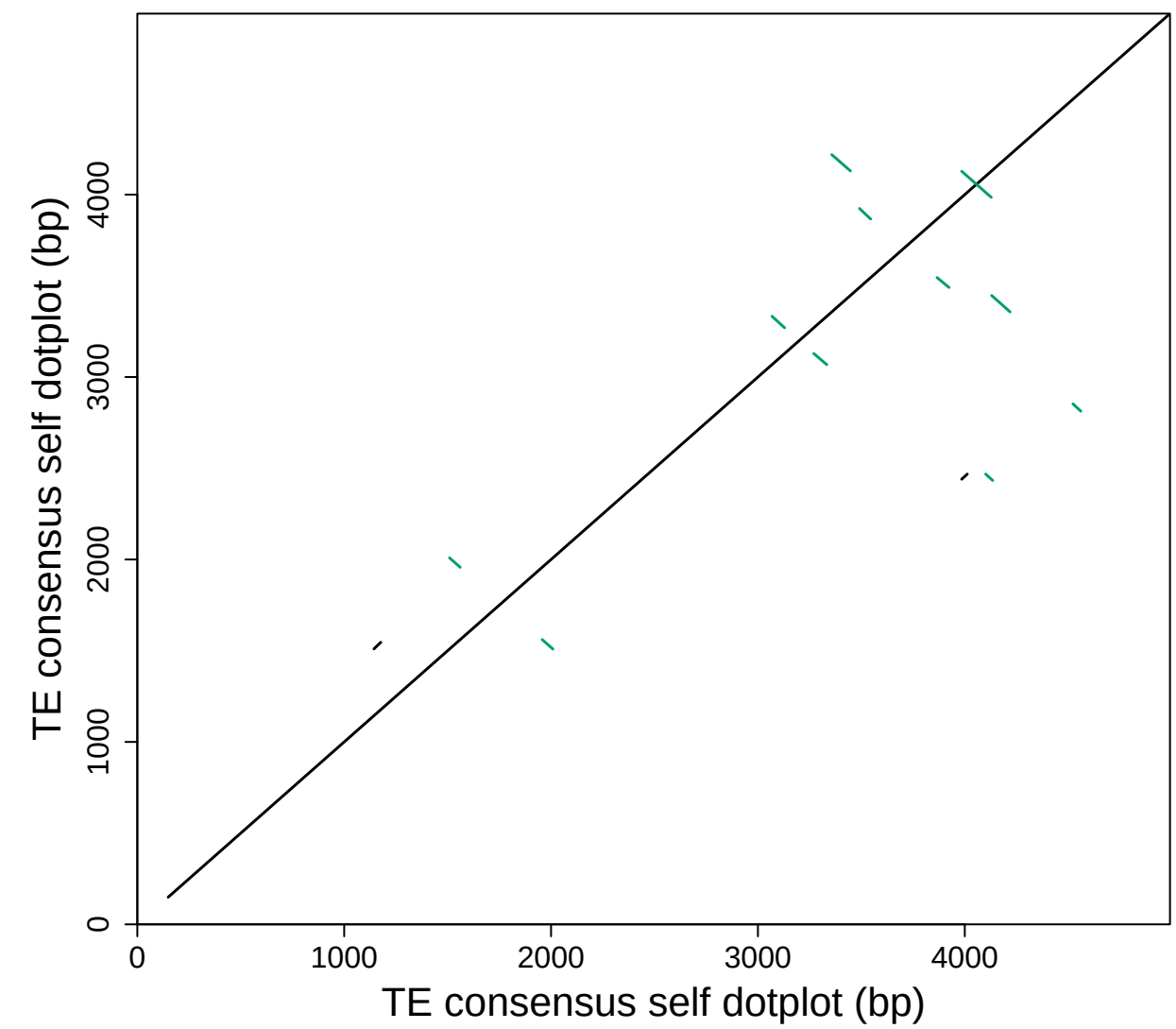
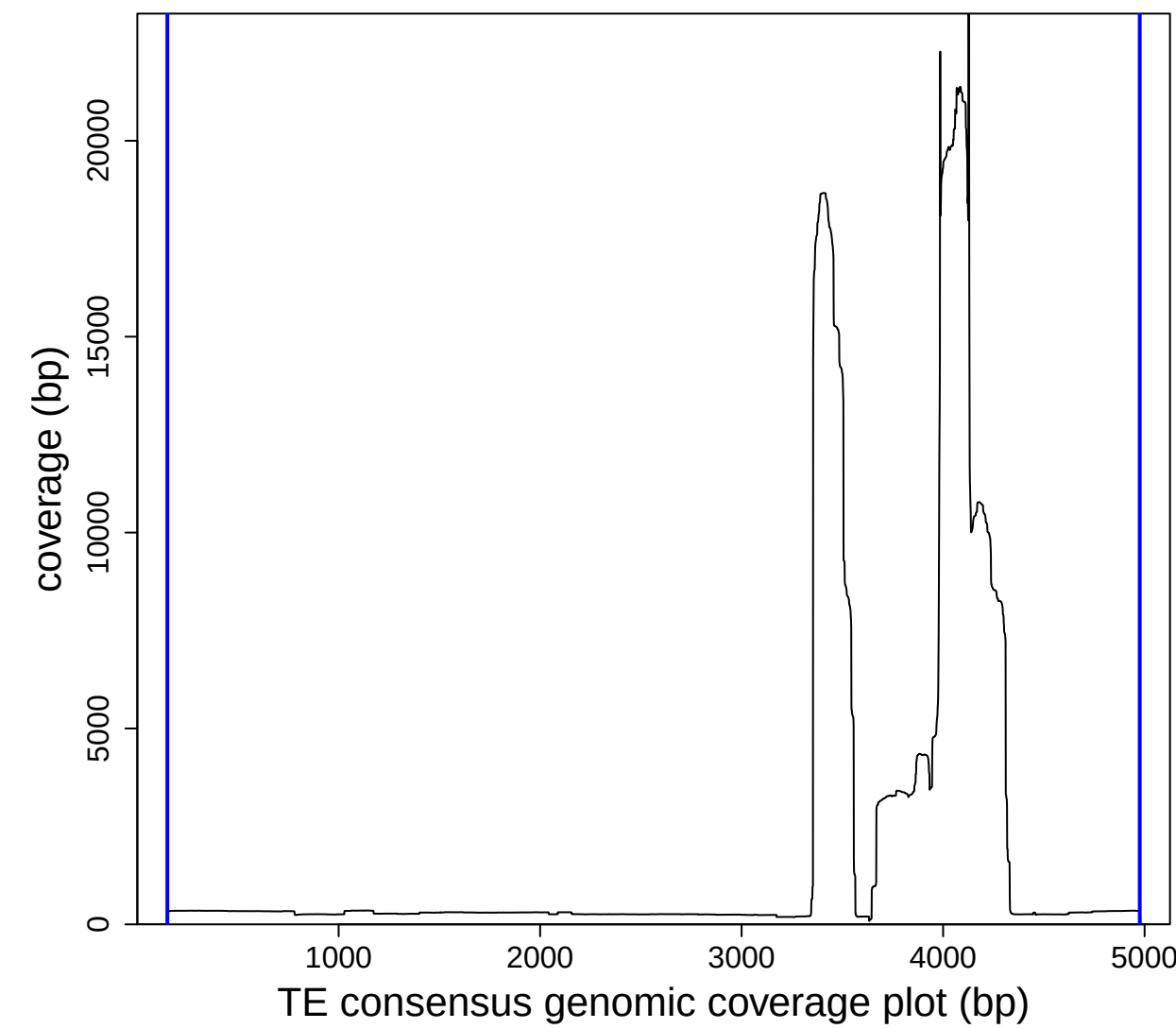
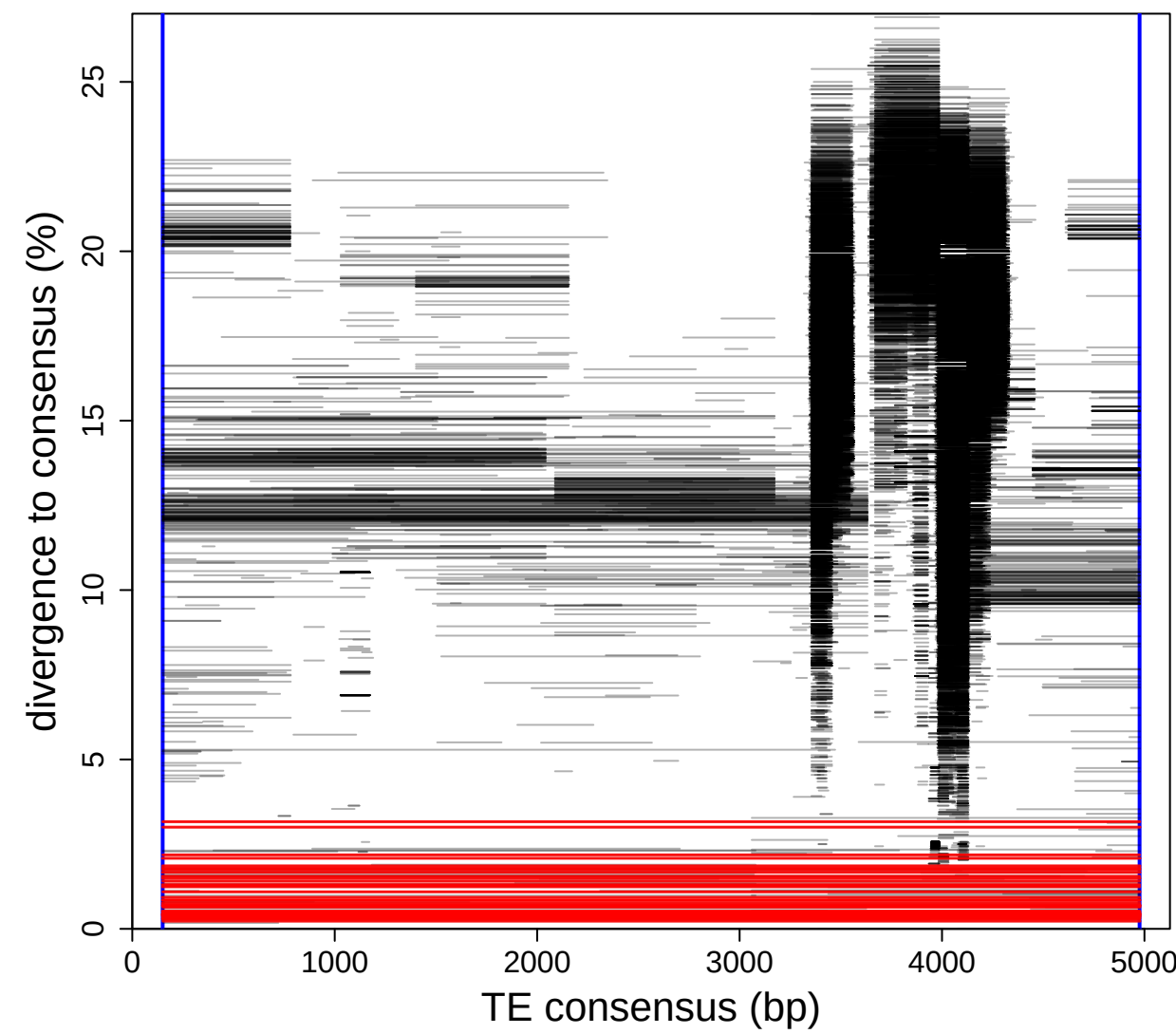
l.b.bed uf.bed g_2.bed fm_1.bed 0 0 bcln.fa aln.fa cl.fa gs.fa
size: 4826bp; fragments: 58388; full length: 44 (>=4343.4bp)

After TEtrimmer 4826 bp

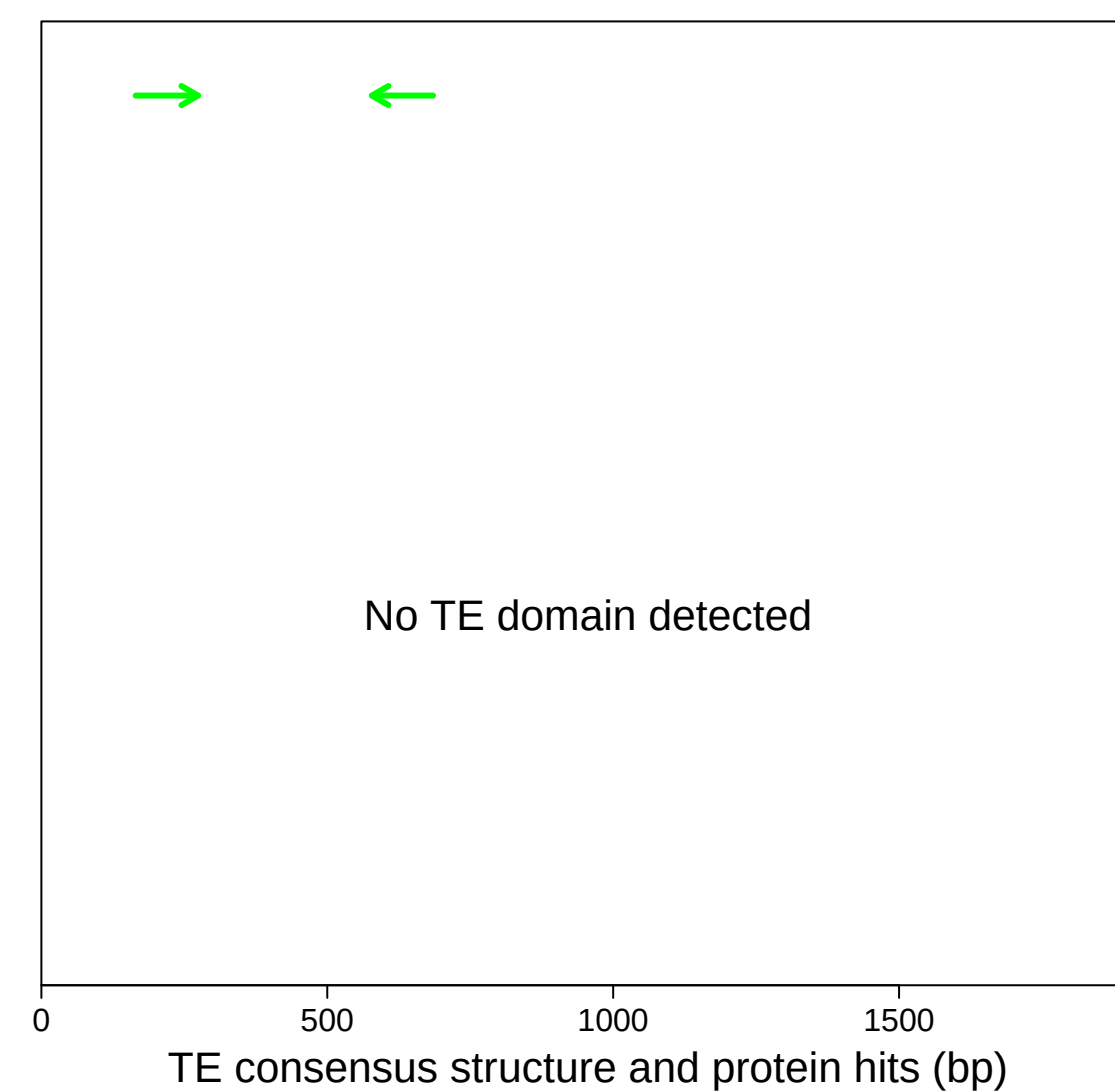
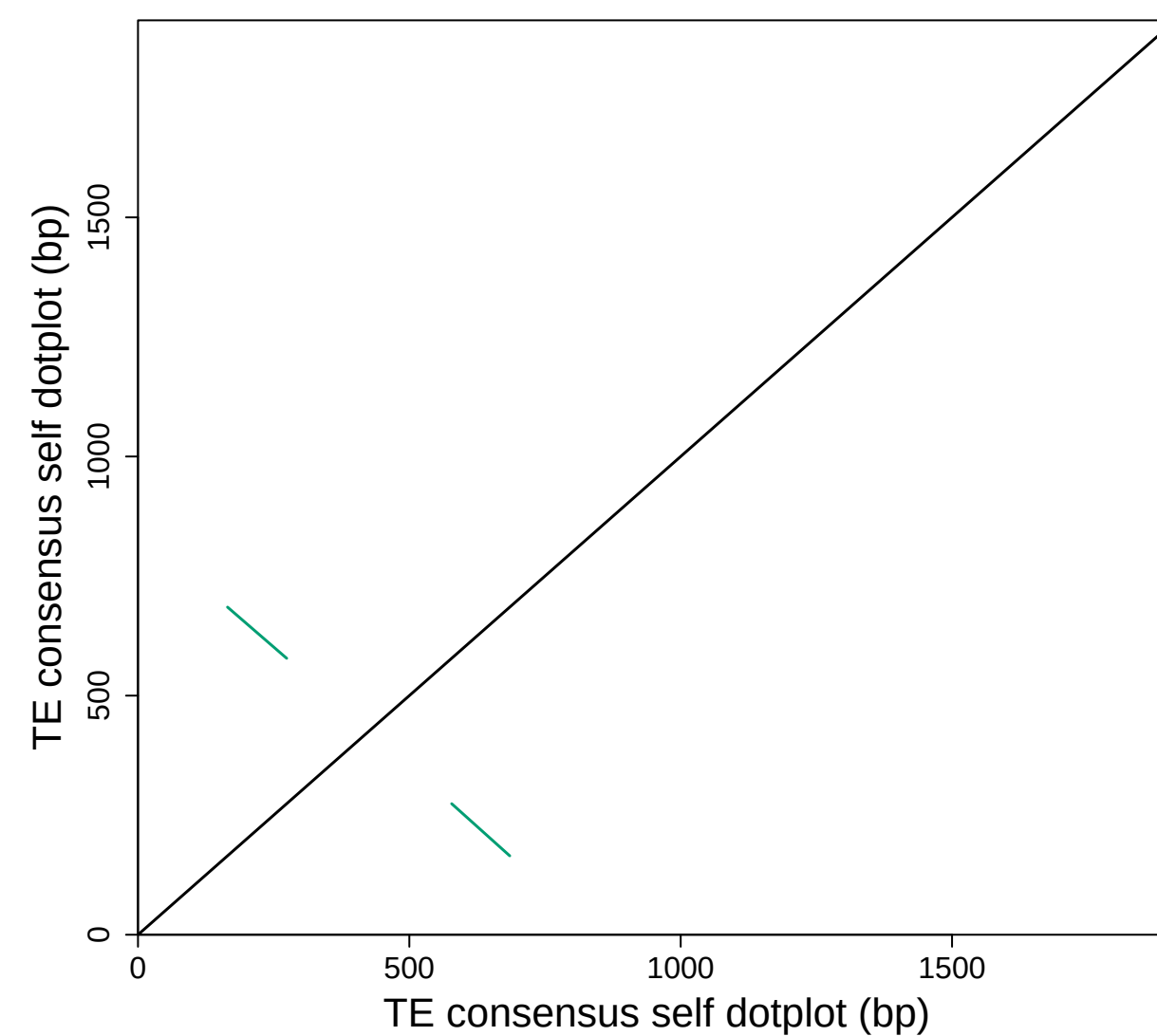
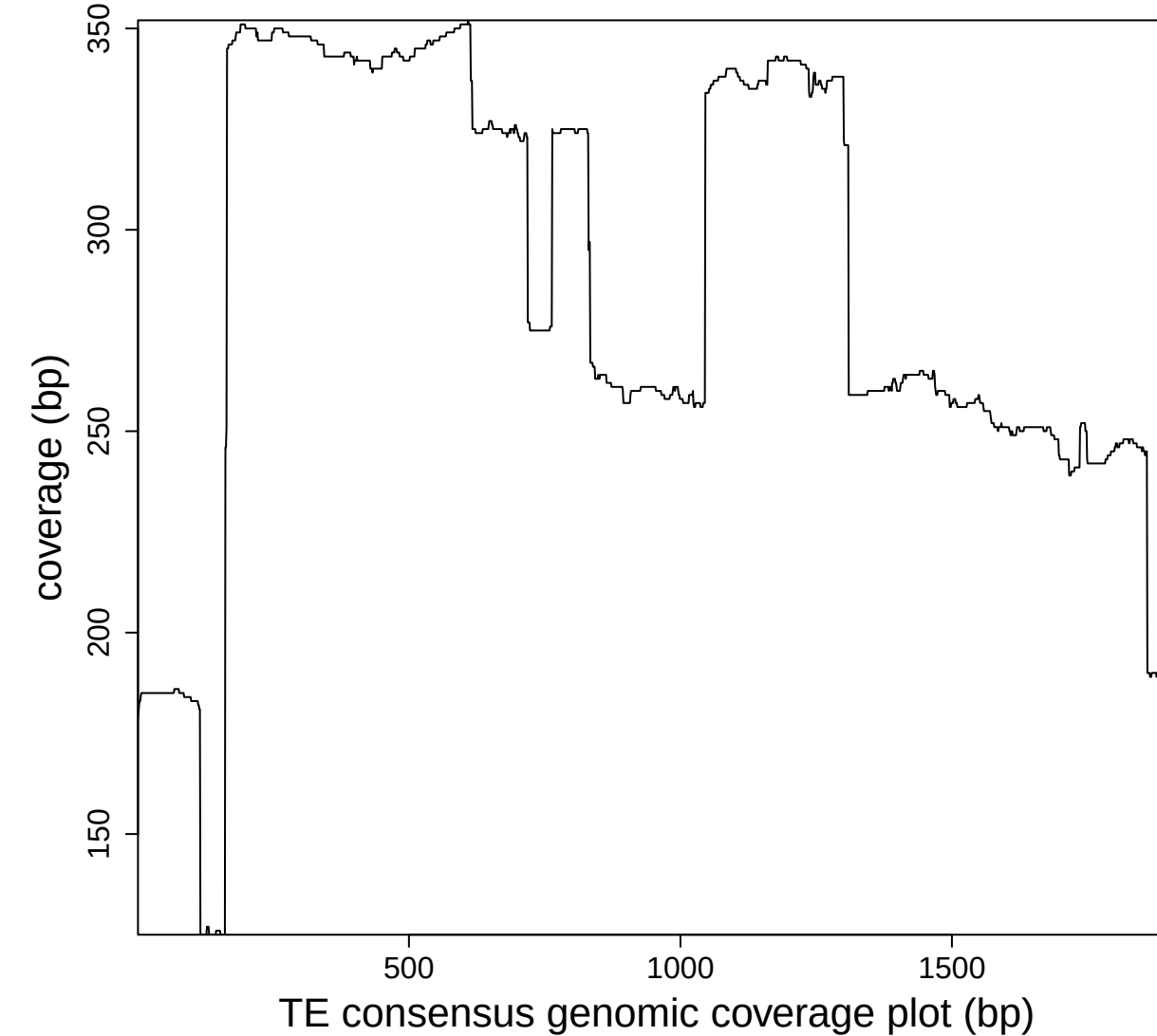
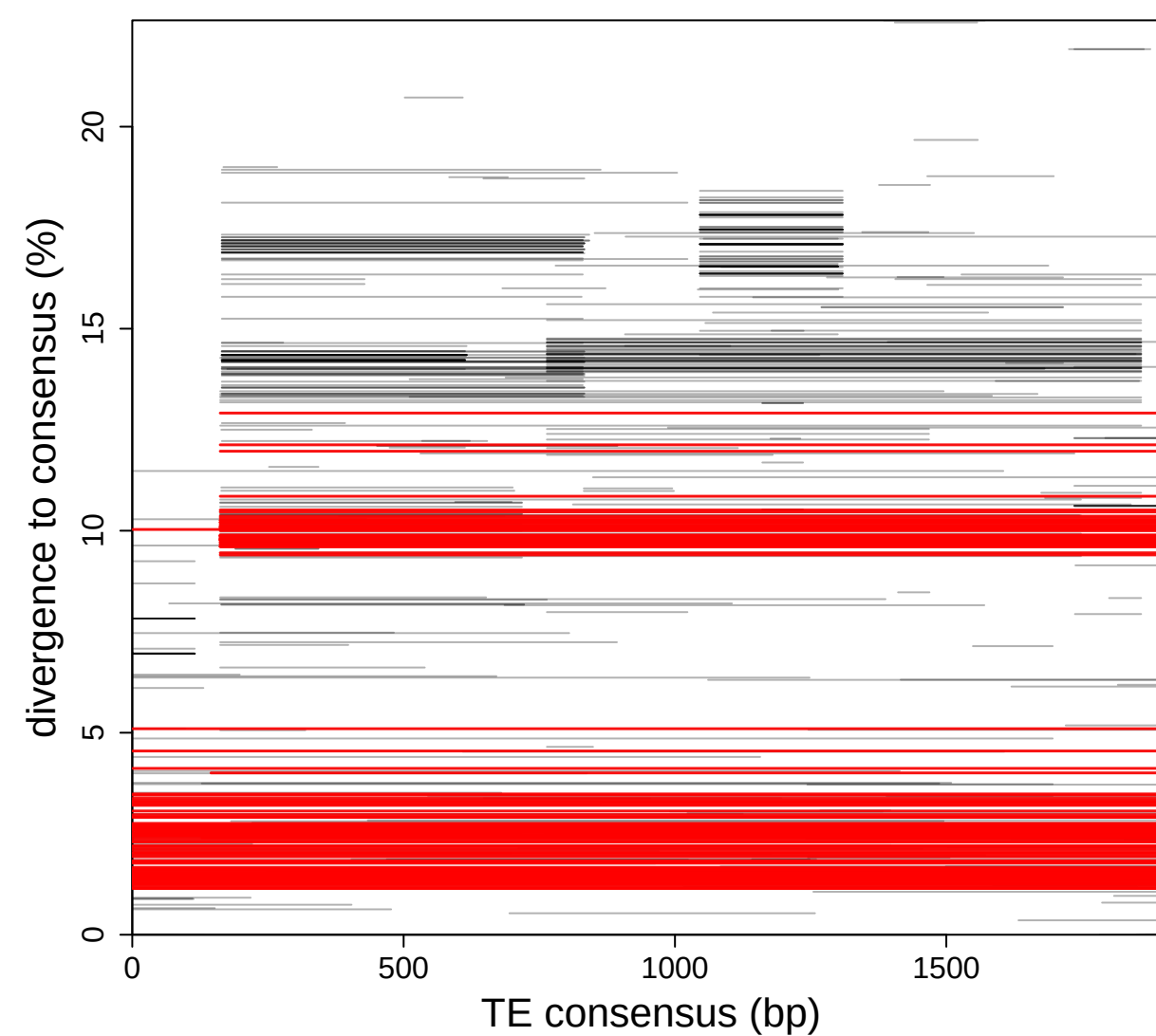


2.fasta.b.bed uf.bed g 2.bed fm 1.bed 0 0 bcln.fa aln.fa cl.fa
size: 5126bp; fragments: 58388; full length: 39 (>=4613.4bp)

After TEtrimmer Extended plot Blue lines are boundaries

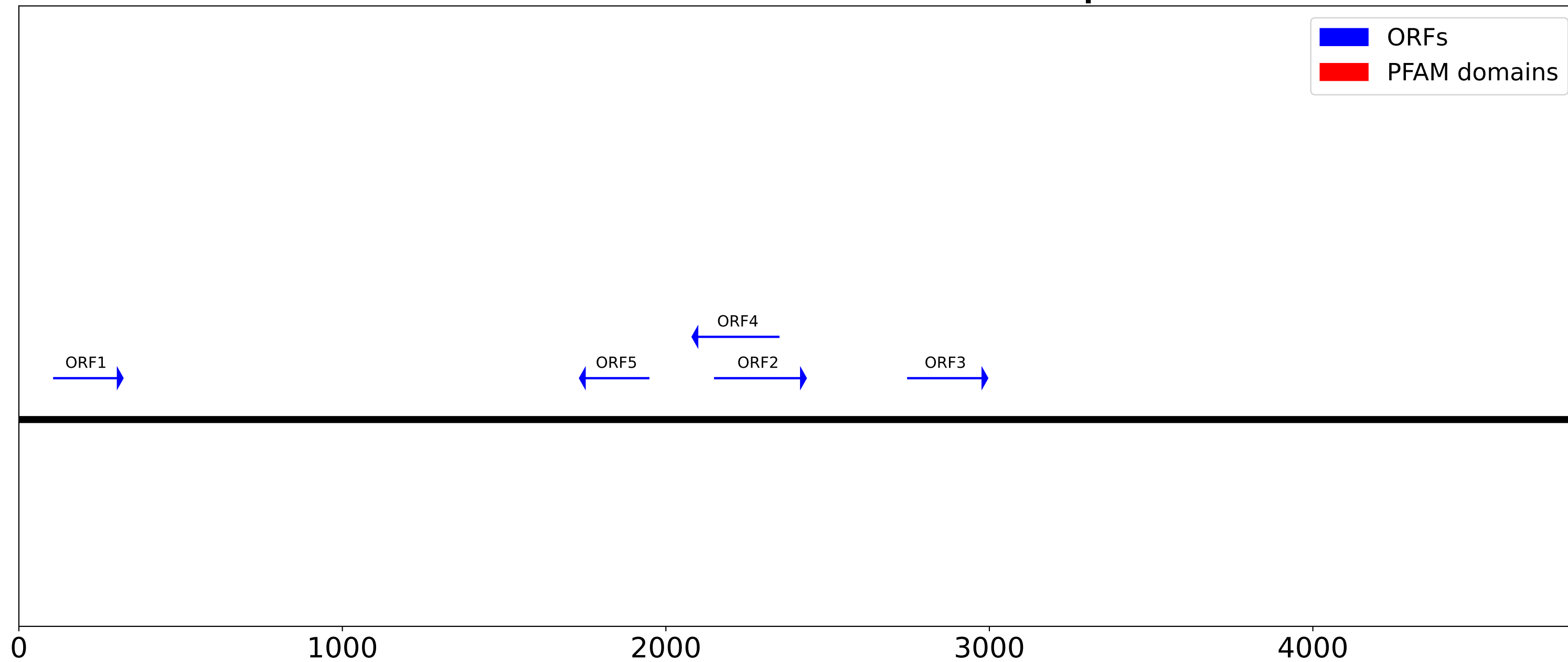


TE: rnd_1_family_642
size: 1912bp; fragments: 677; full length: 132 (≥ 1720.8 bp)

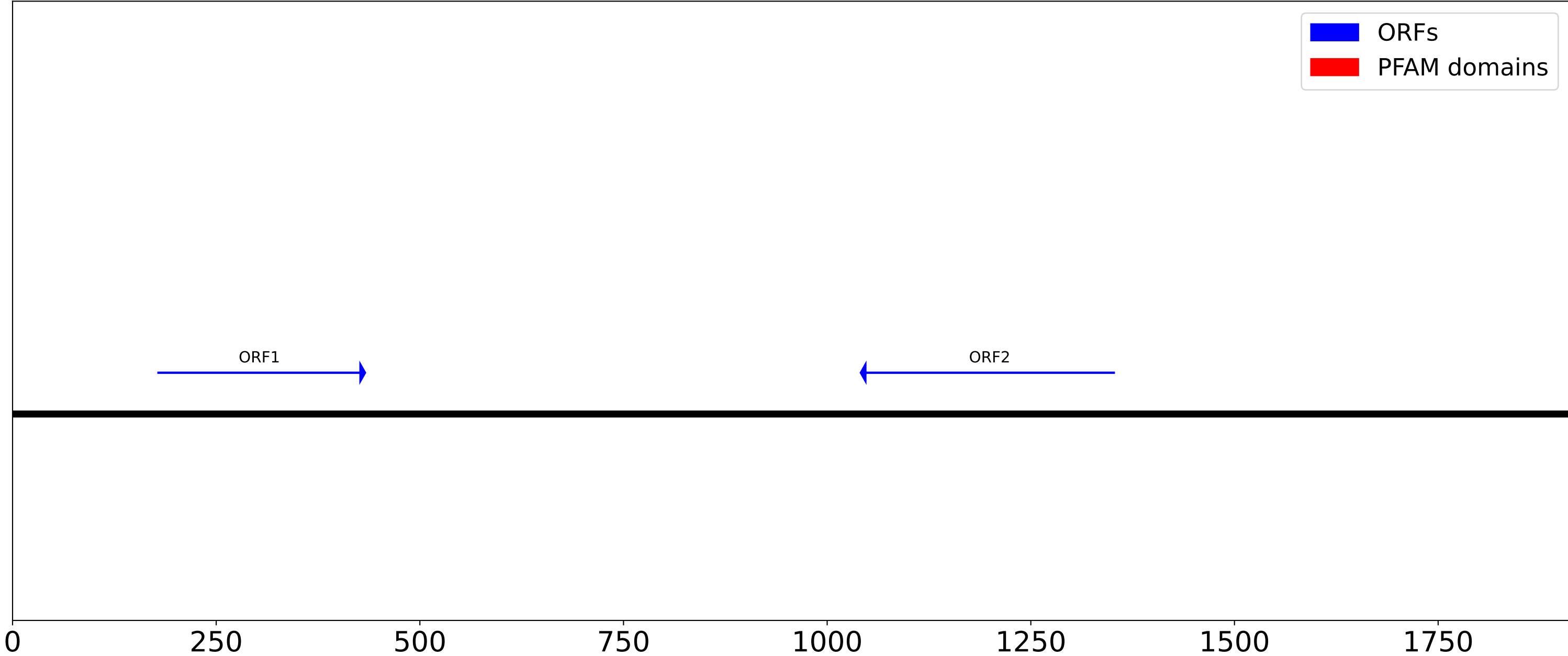


Before TEtrimmer 1912 bp

After TEtrimmer ORF and PFAM domain plot



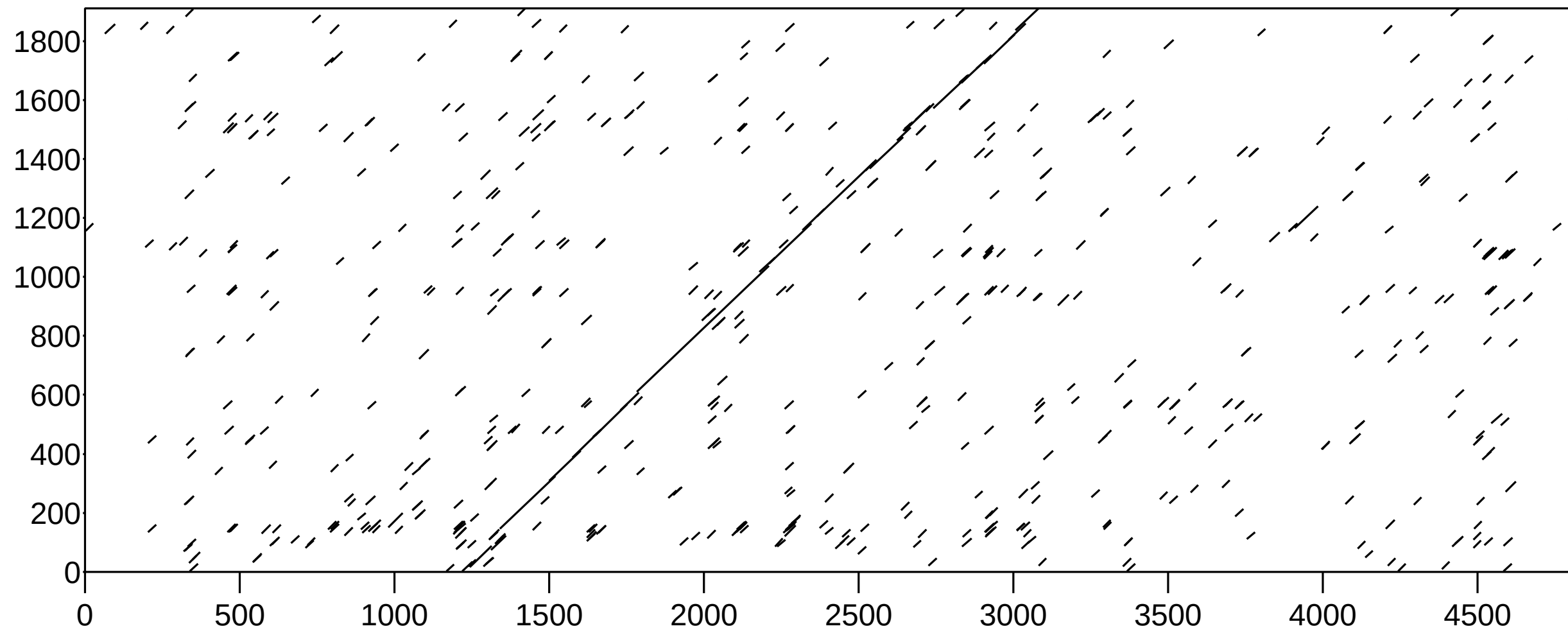
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)