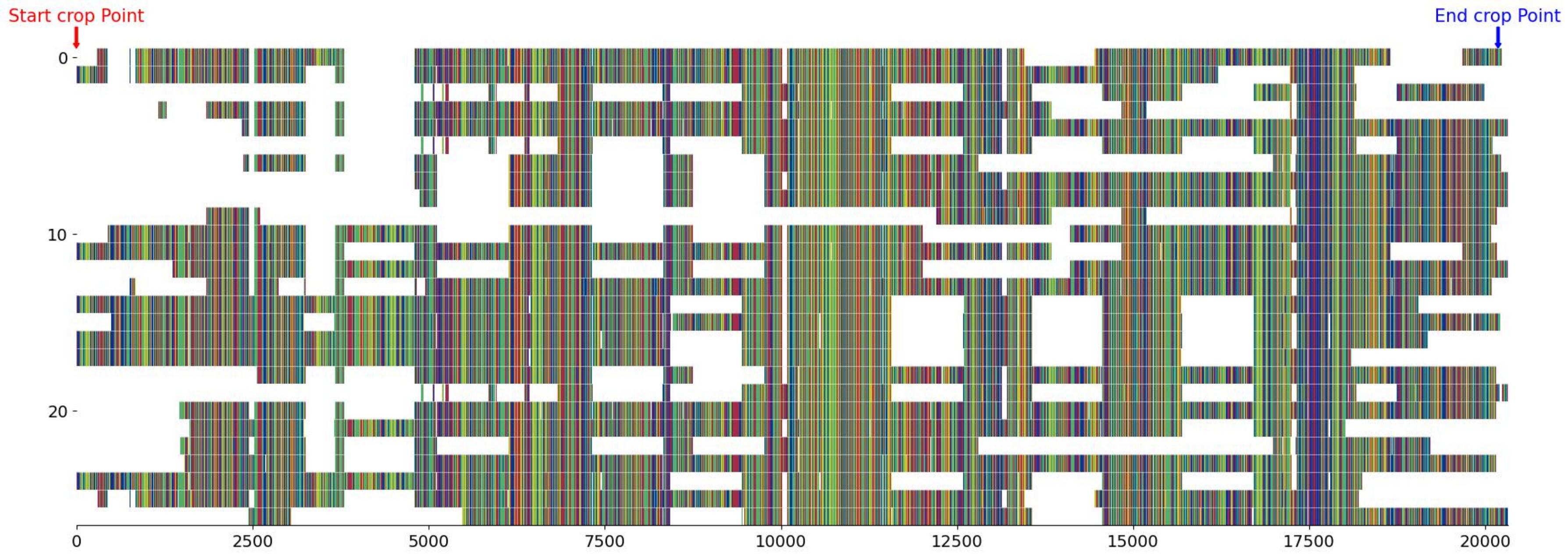
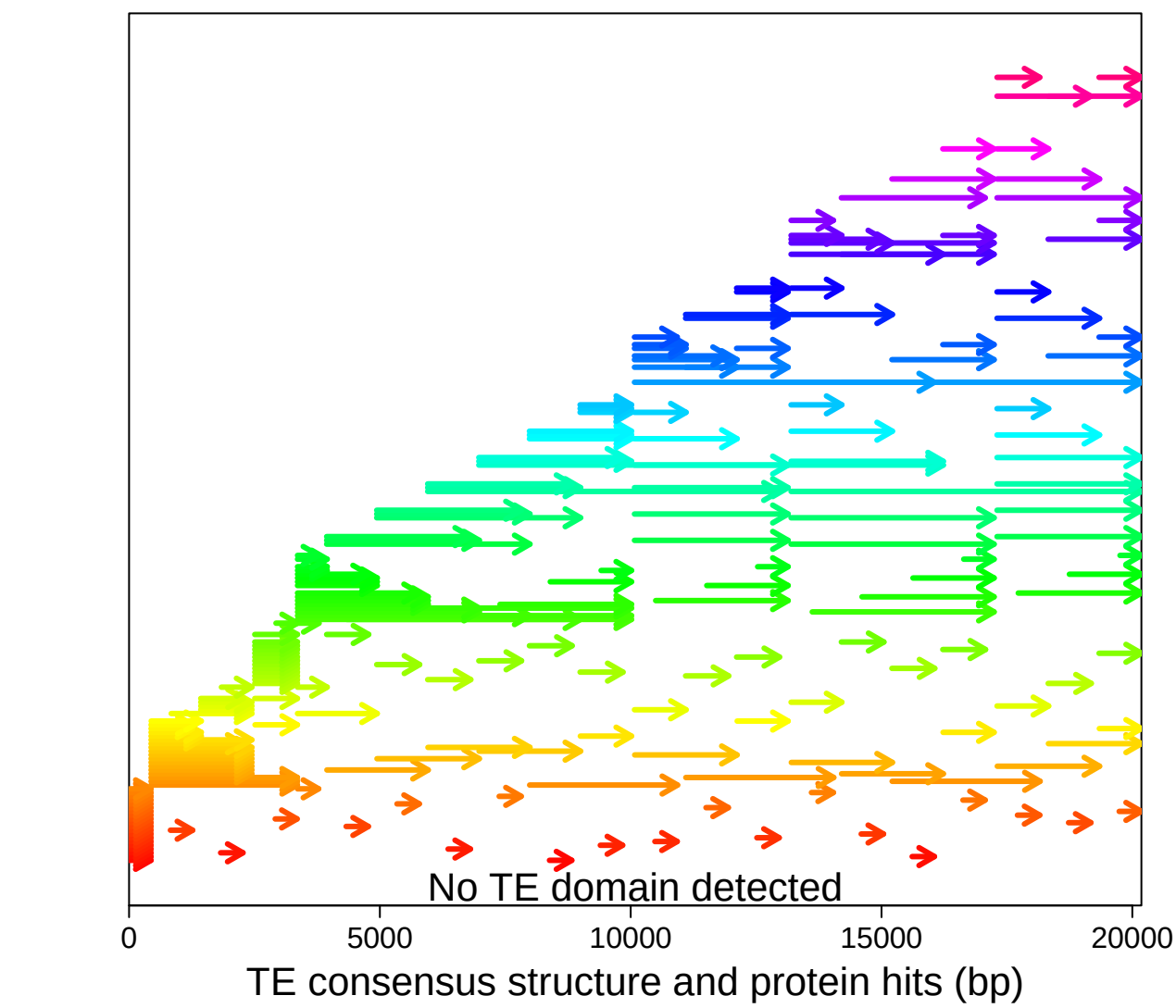
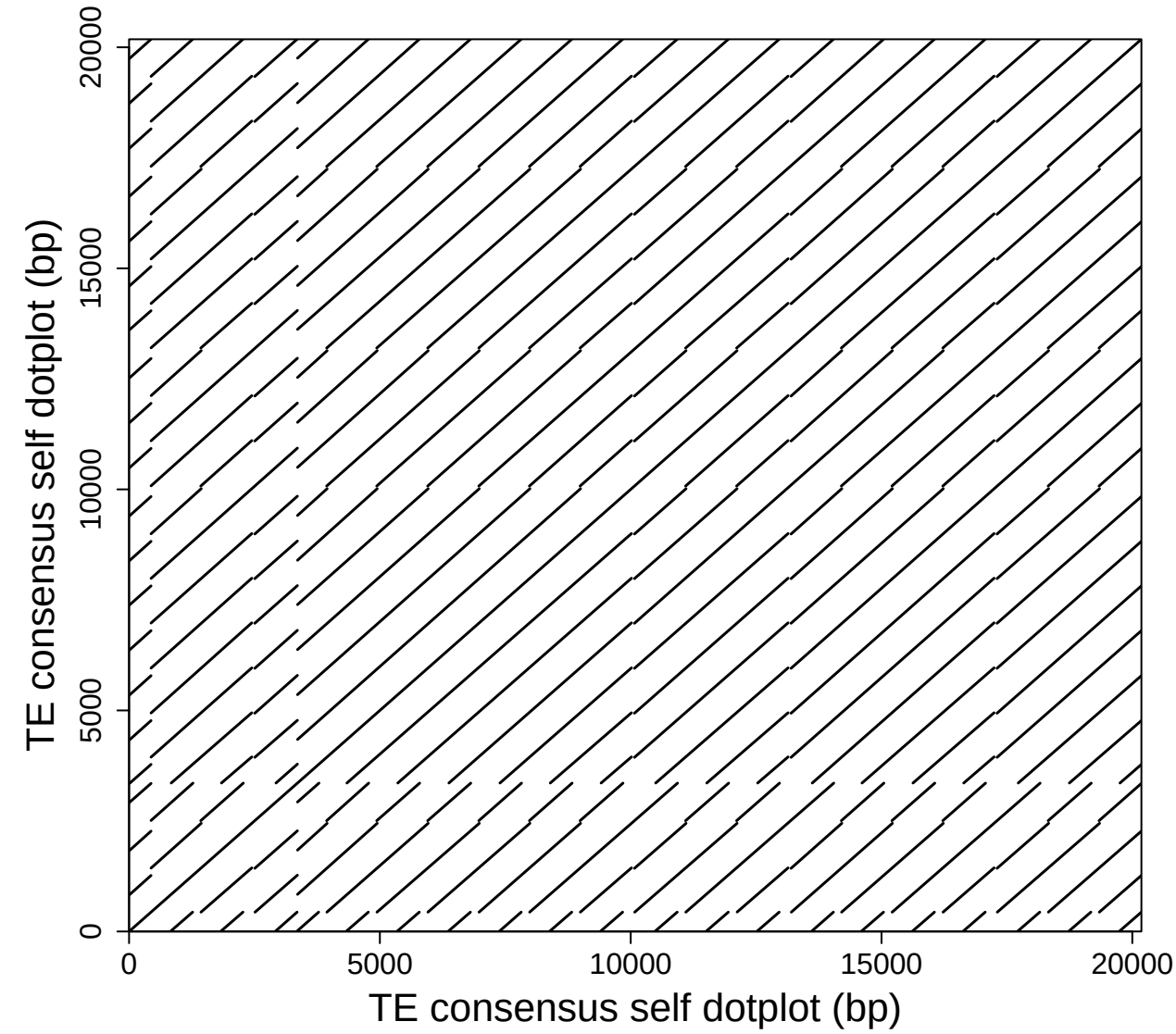
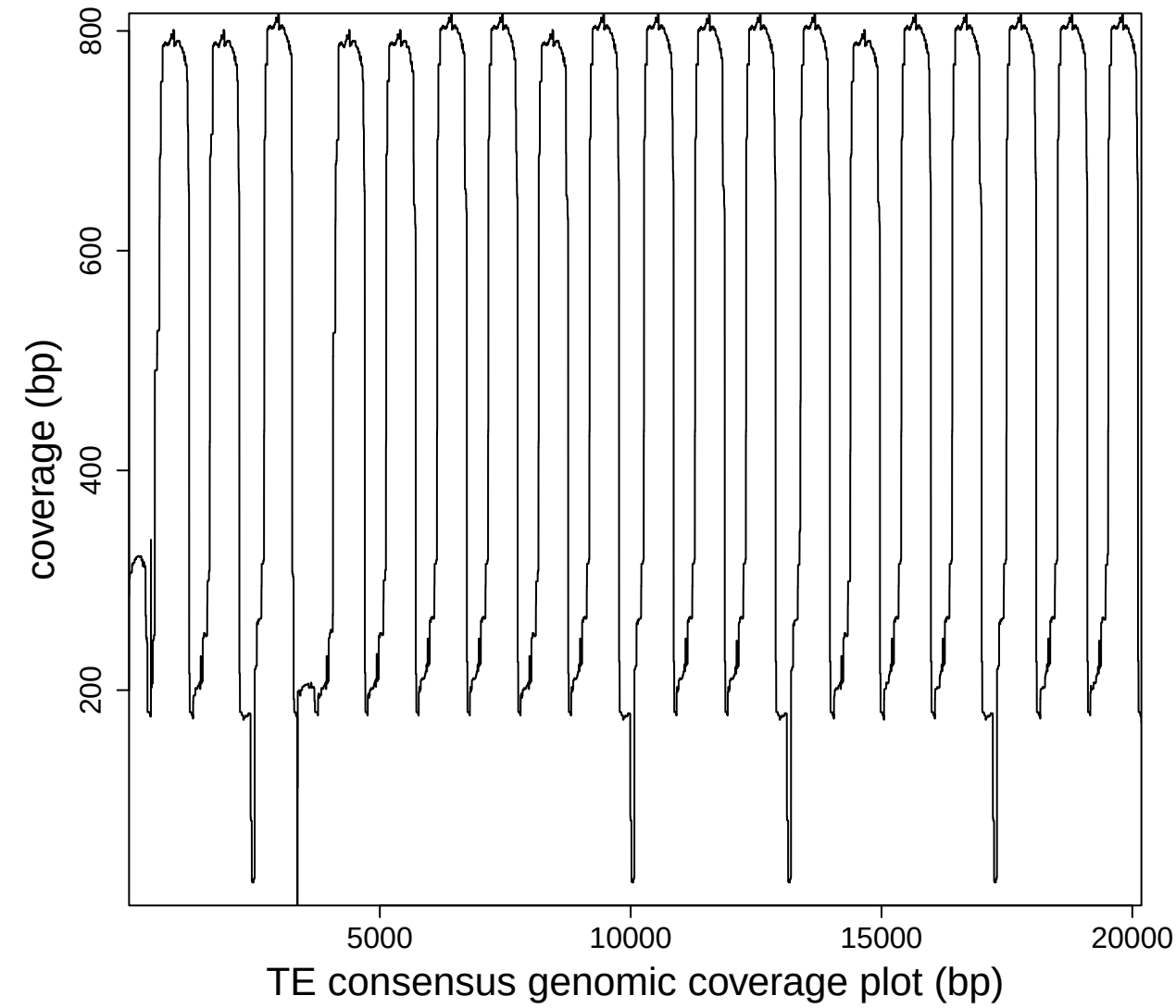
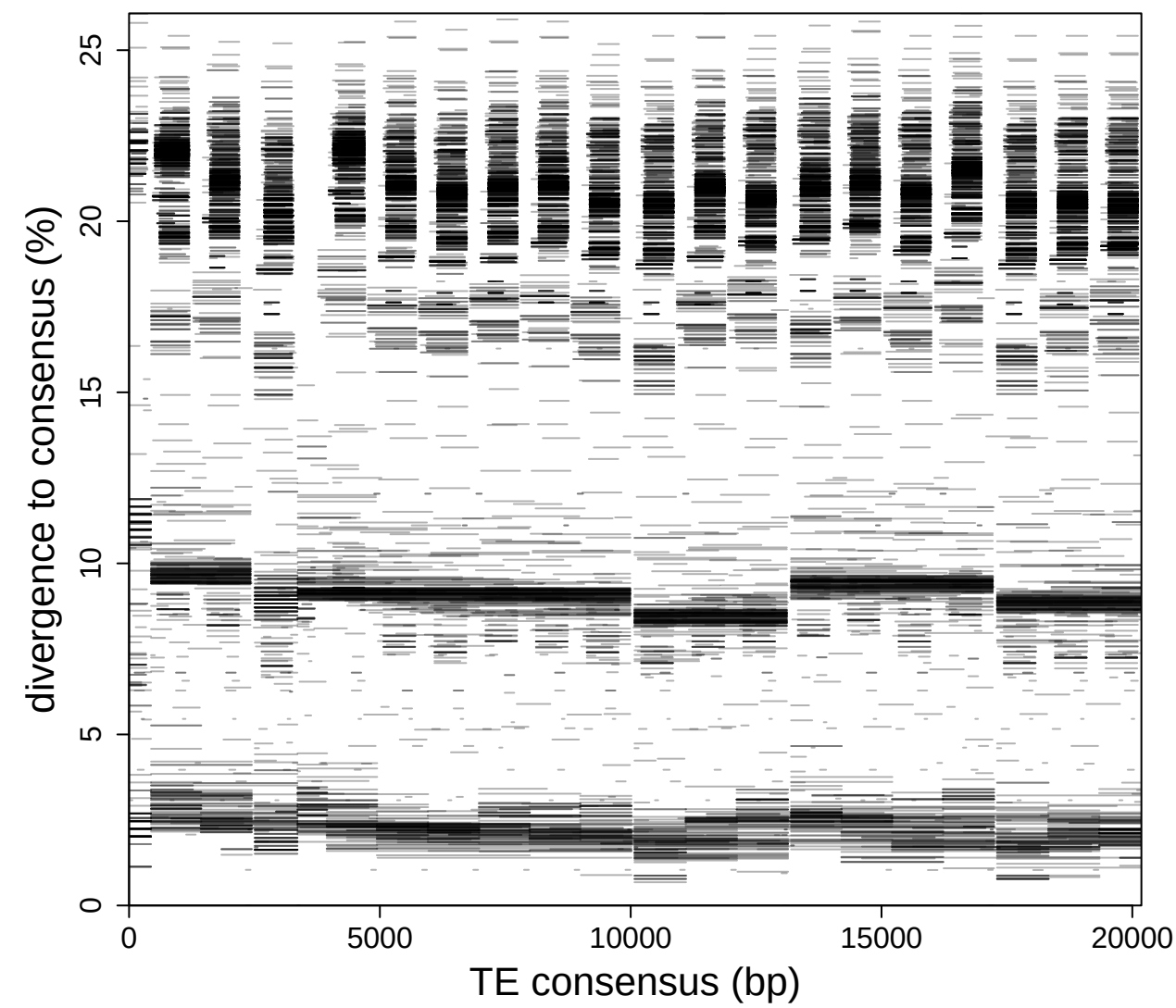


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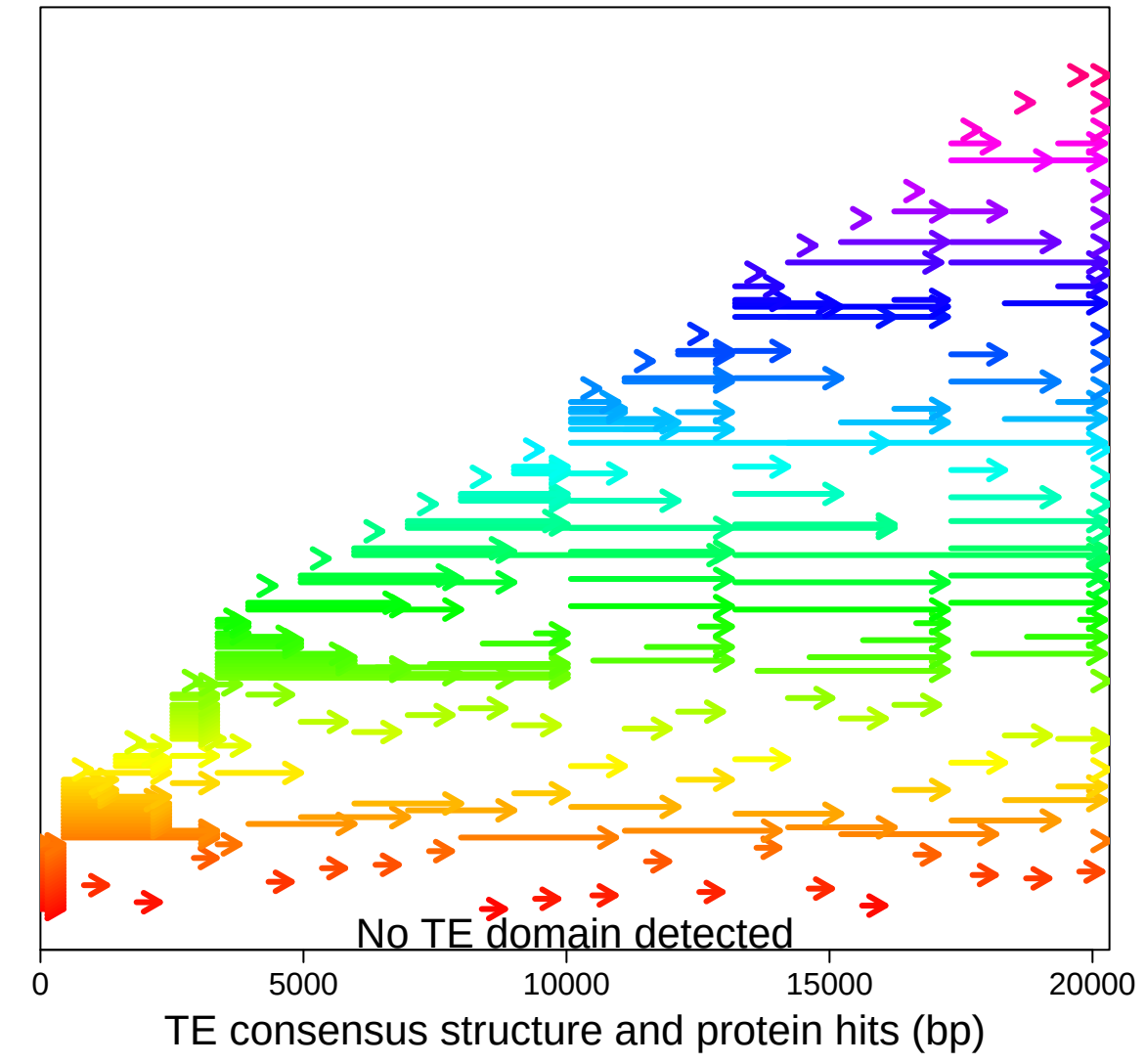
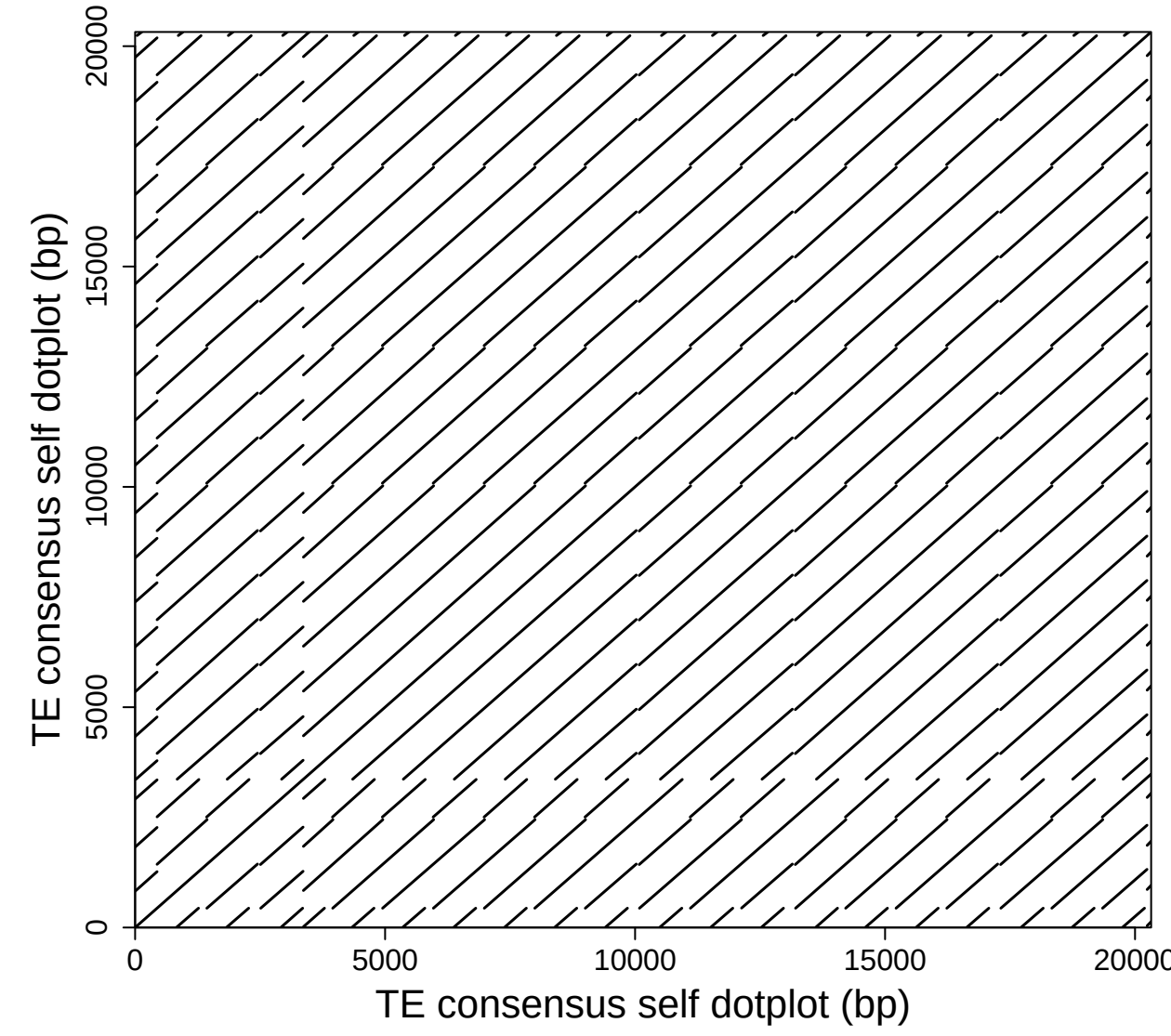
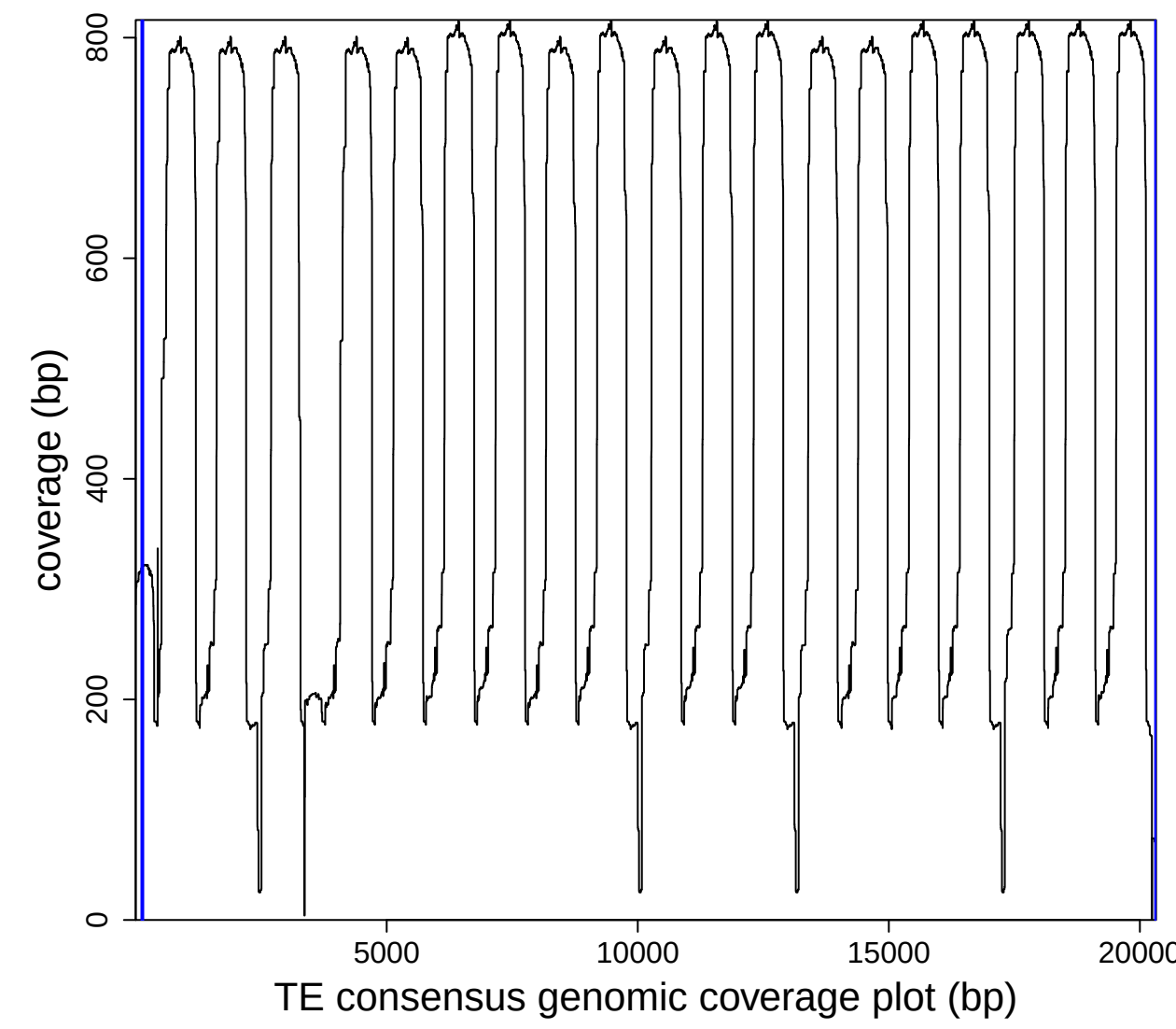
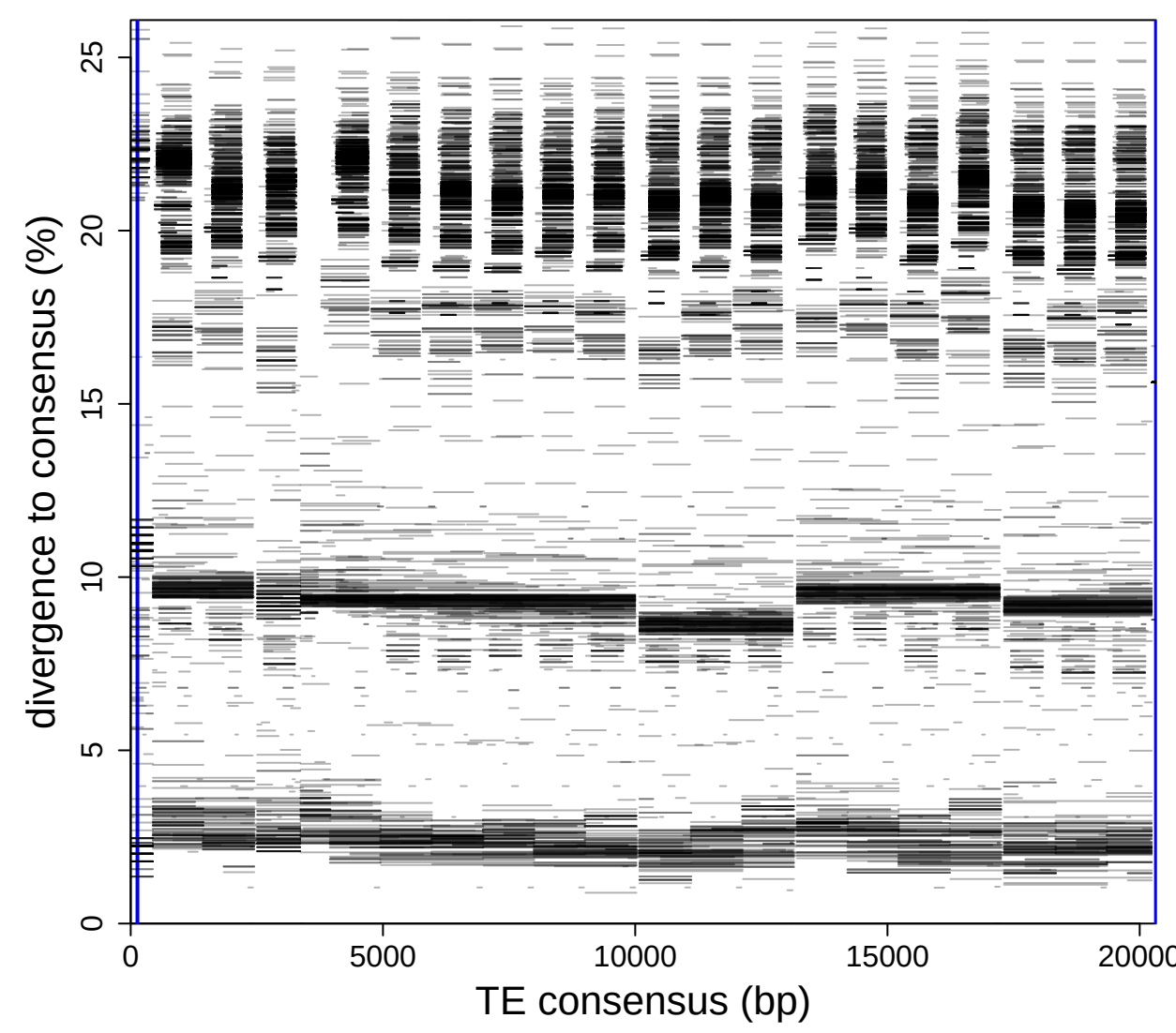
bed uf.bed g 2.bed fm 1.bed 0 0 bcln.fa aln.fa cl.fa gs.fa ce
size: 20181bp; fragments: 14656; full length: 0 (>=18162.9bp)

After TEtrimmer 20181 bp

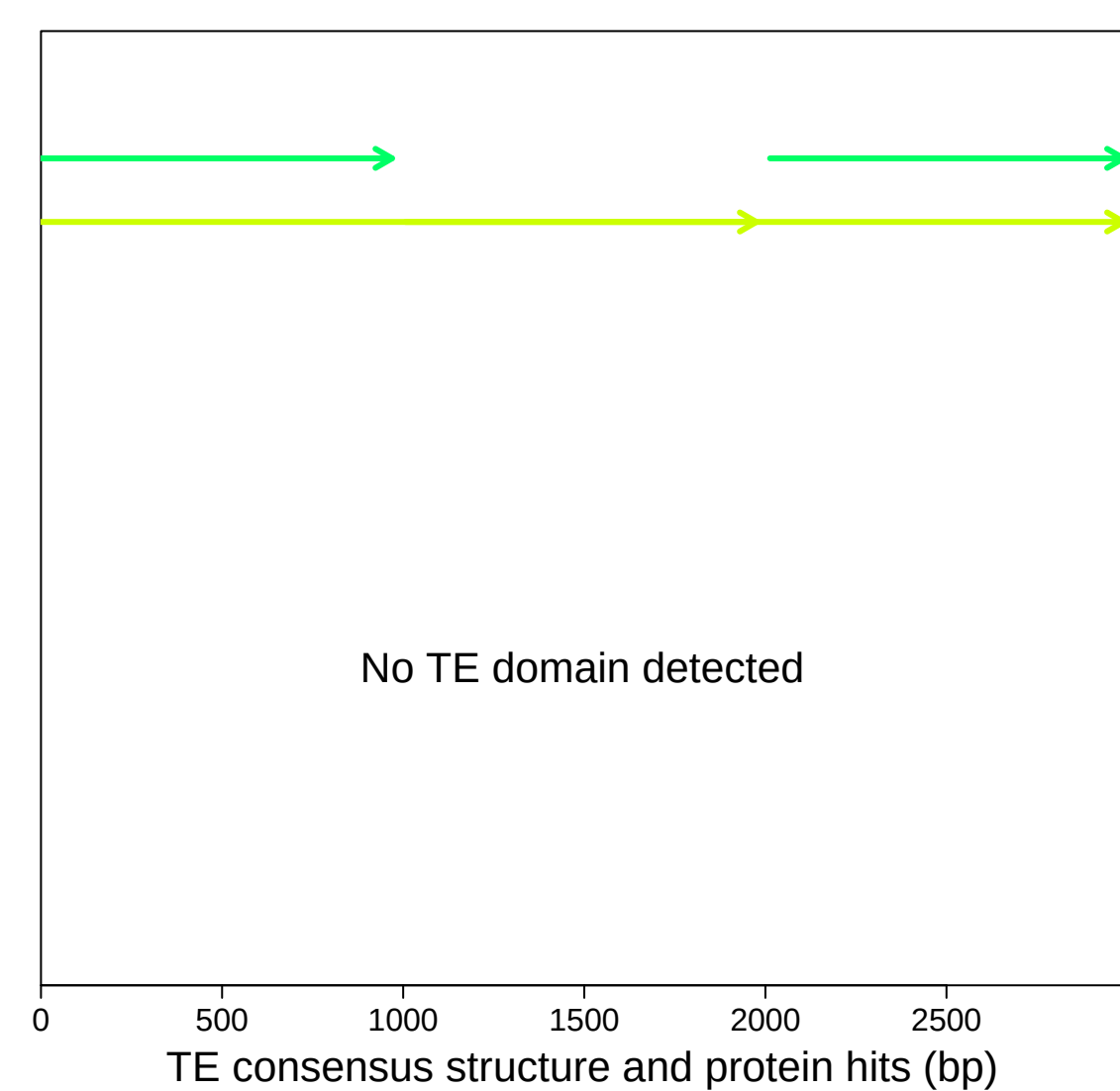
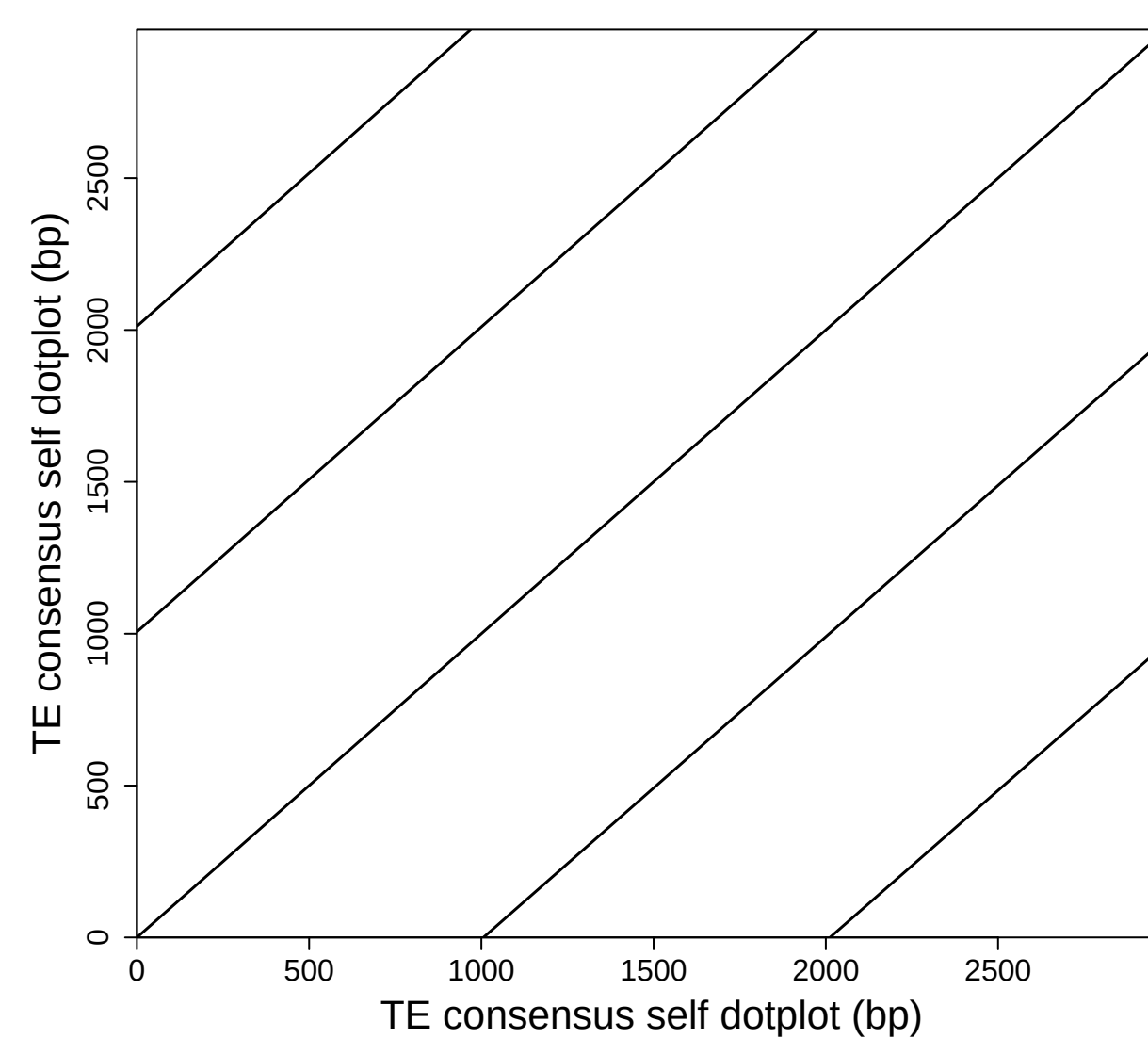
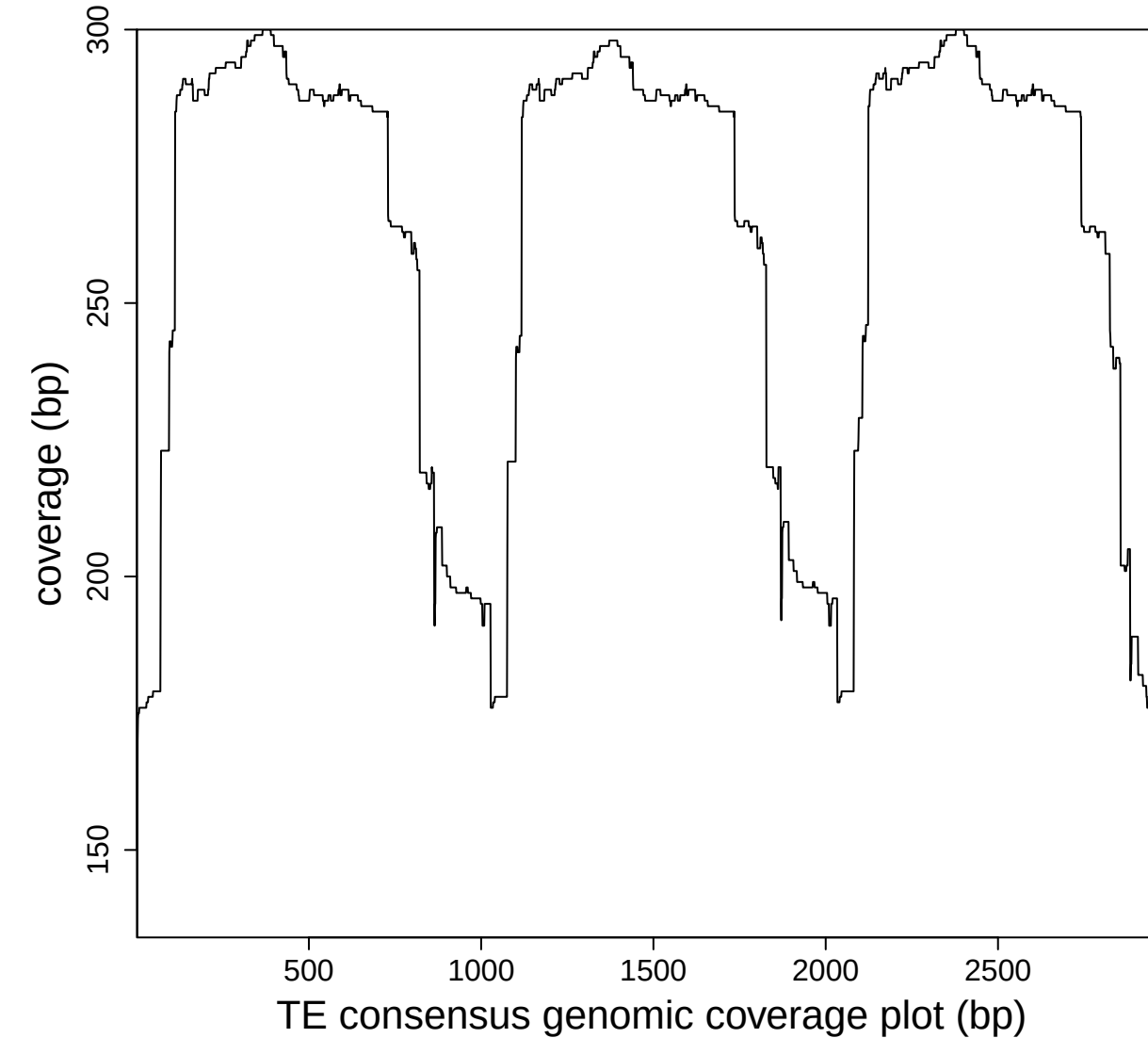
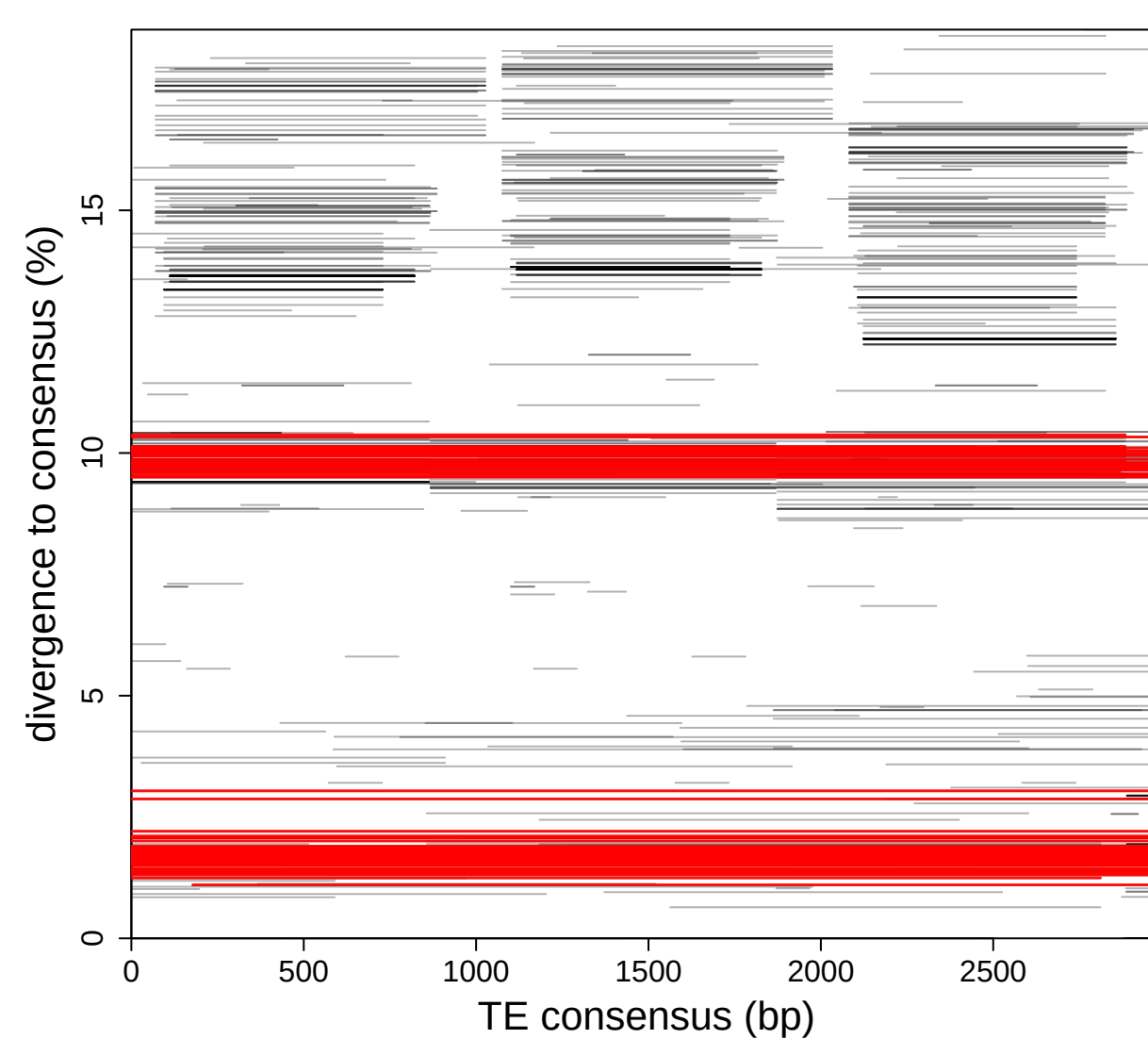


_316.fasta.b.bed uf.bed g 2.bed fm 1.bed 0 0 bcln.fa aln.fa c
size: 20322bp; fragments: 14668; full length: 0 (>=18289.8bp)

After TEtrimmer Extended plot Blue lines are boundaries

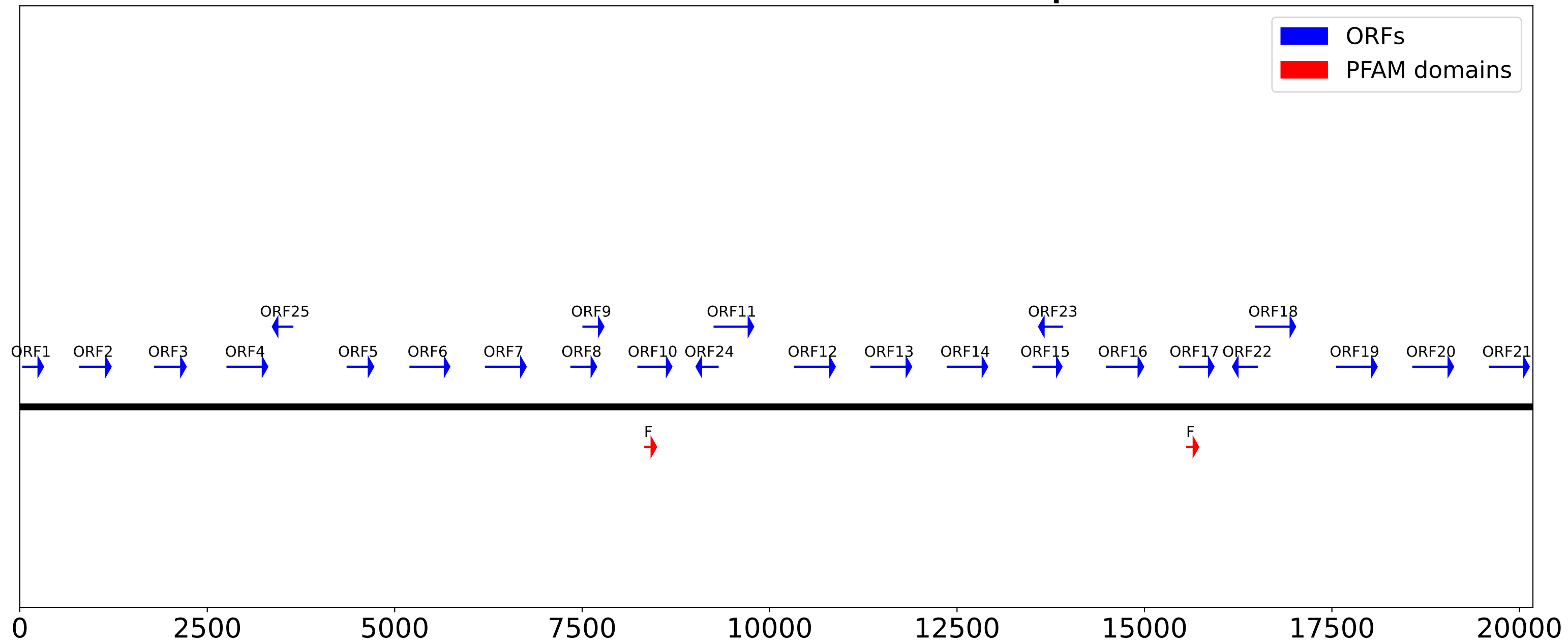


TE: rnd_1_family_316
size: 2989bp; fragments: 729; full length: 93 (>=2690.1bp)

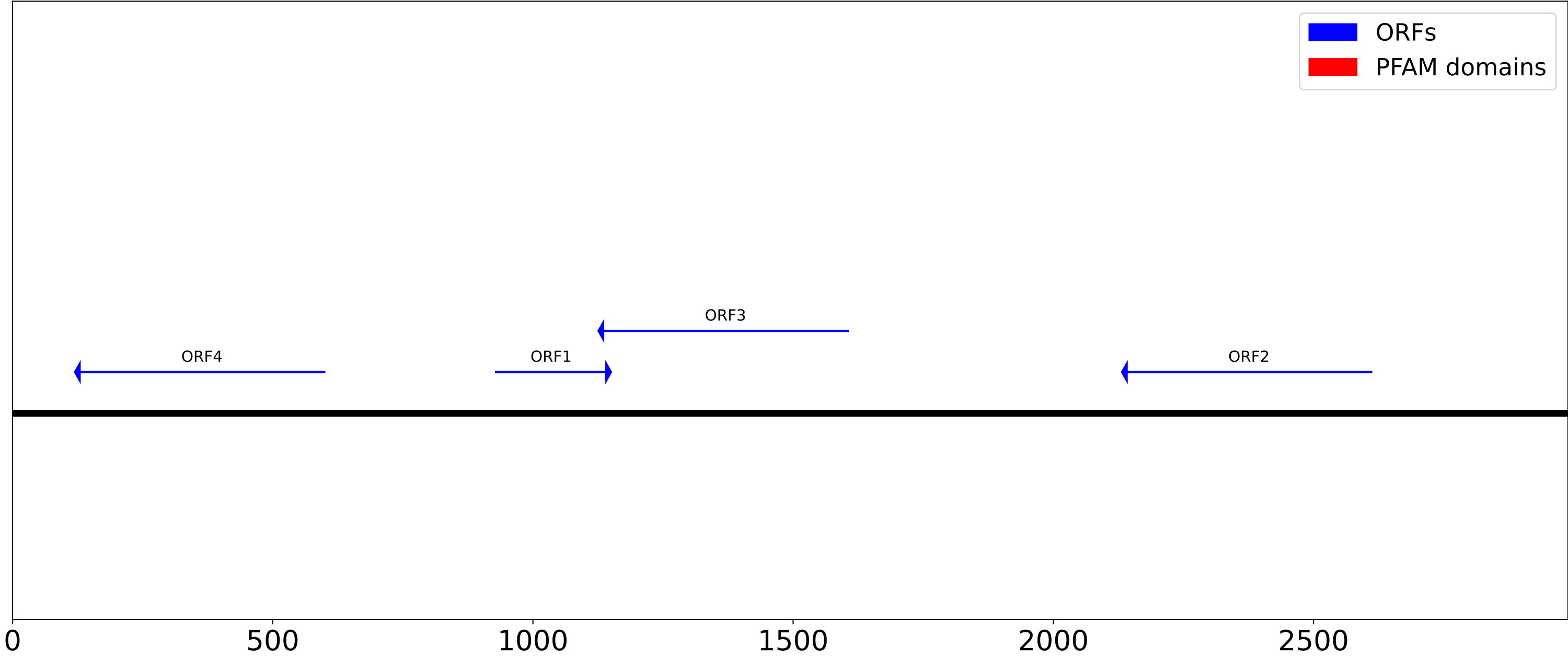


Before TETrimmer 2989 bp

After TEtrimmer ORF and PFAM domain plot



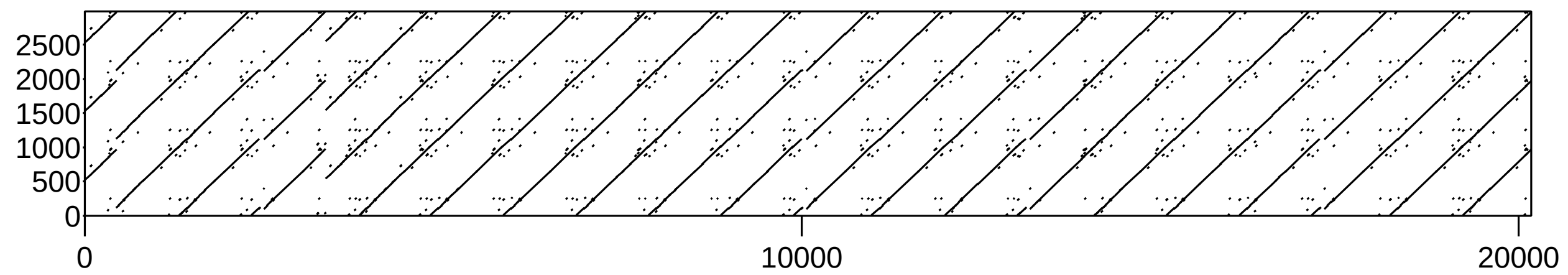
Before TEtrimmer ORF and PFAM domain plot



Dotplot

(window size = 25, threshold = 50.00 09/10/25)

Isensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)