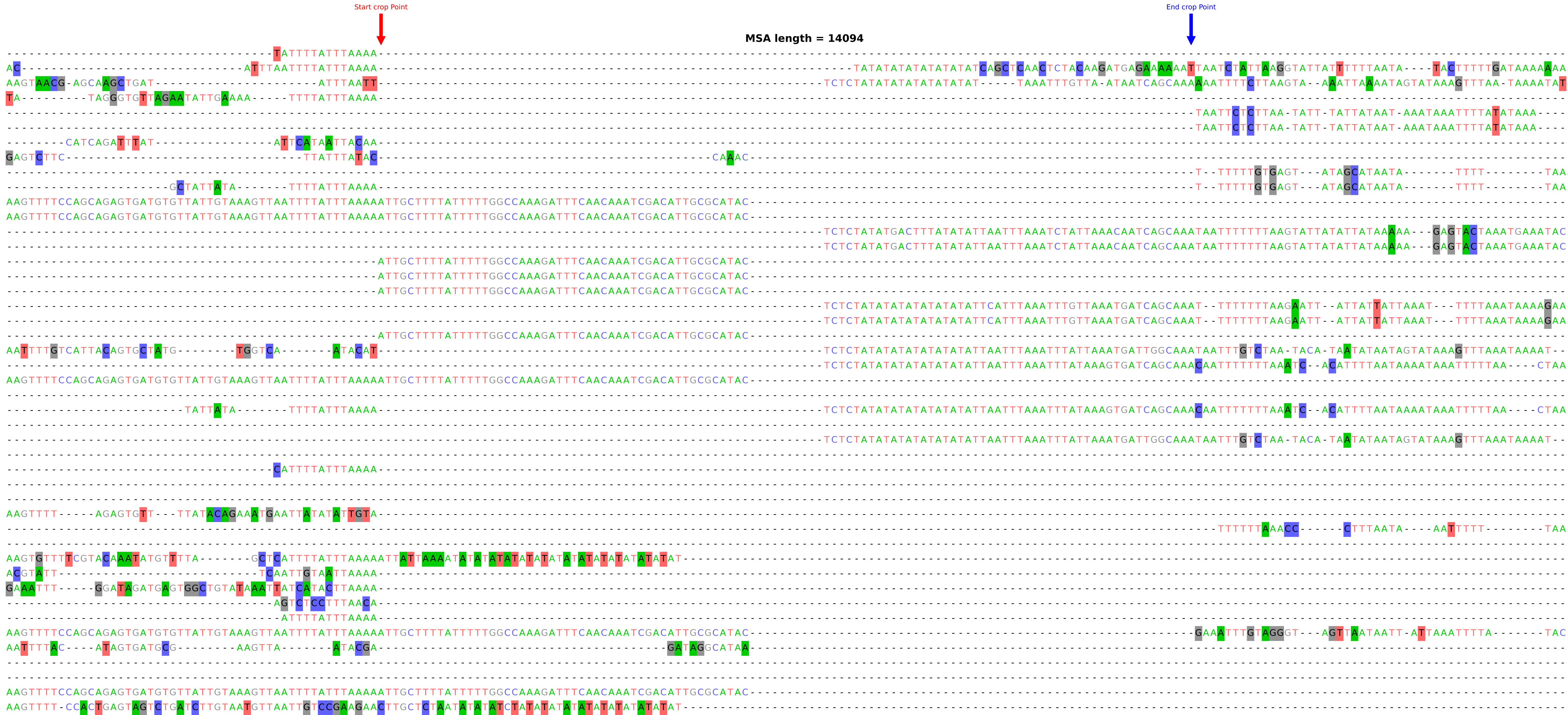
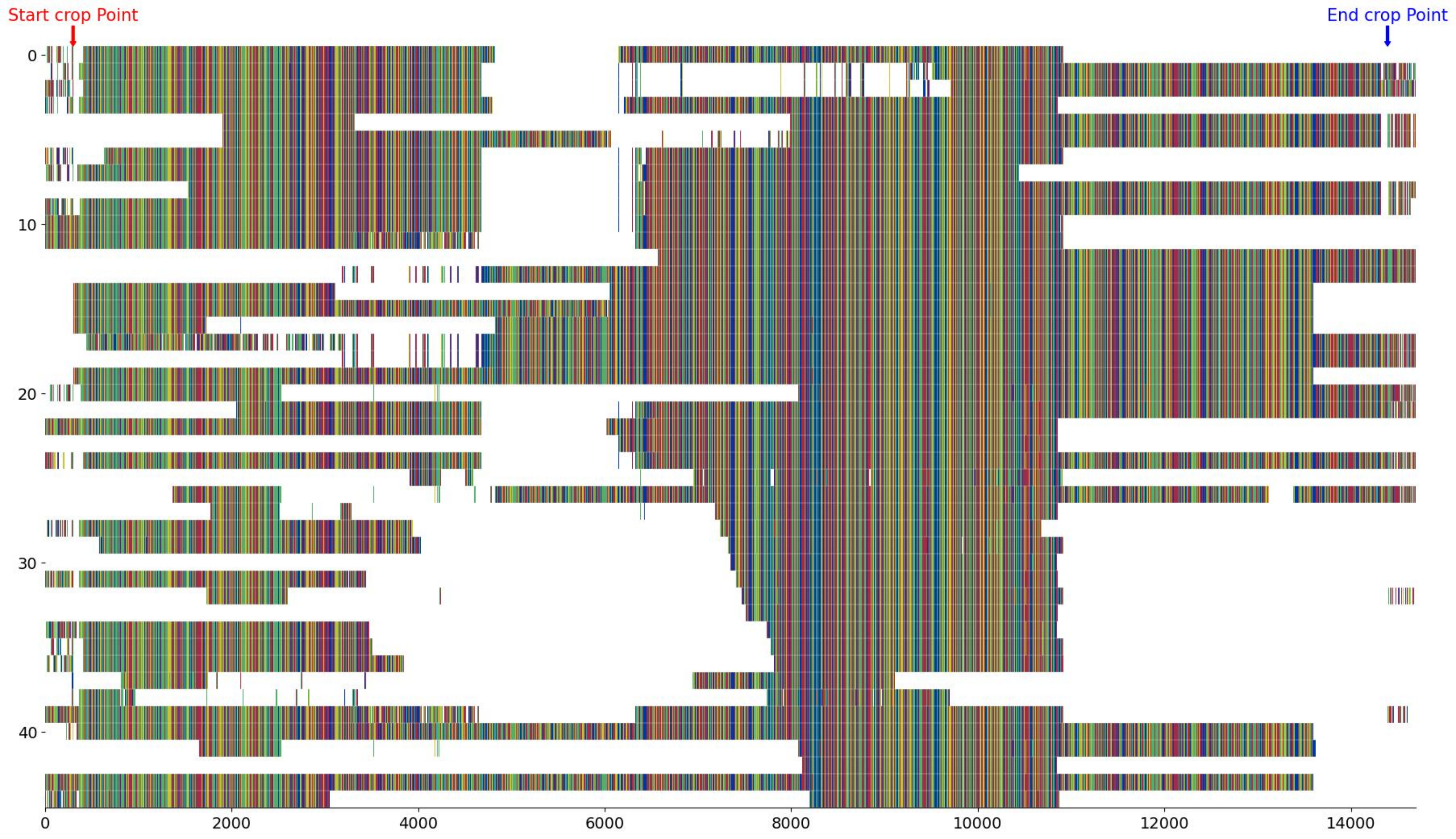
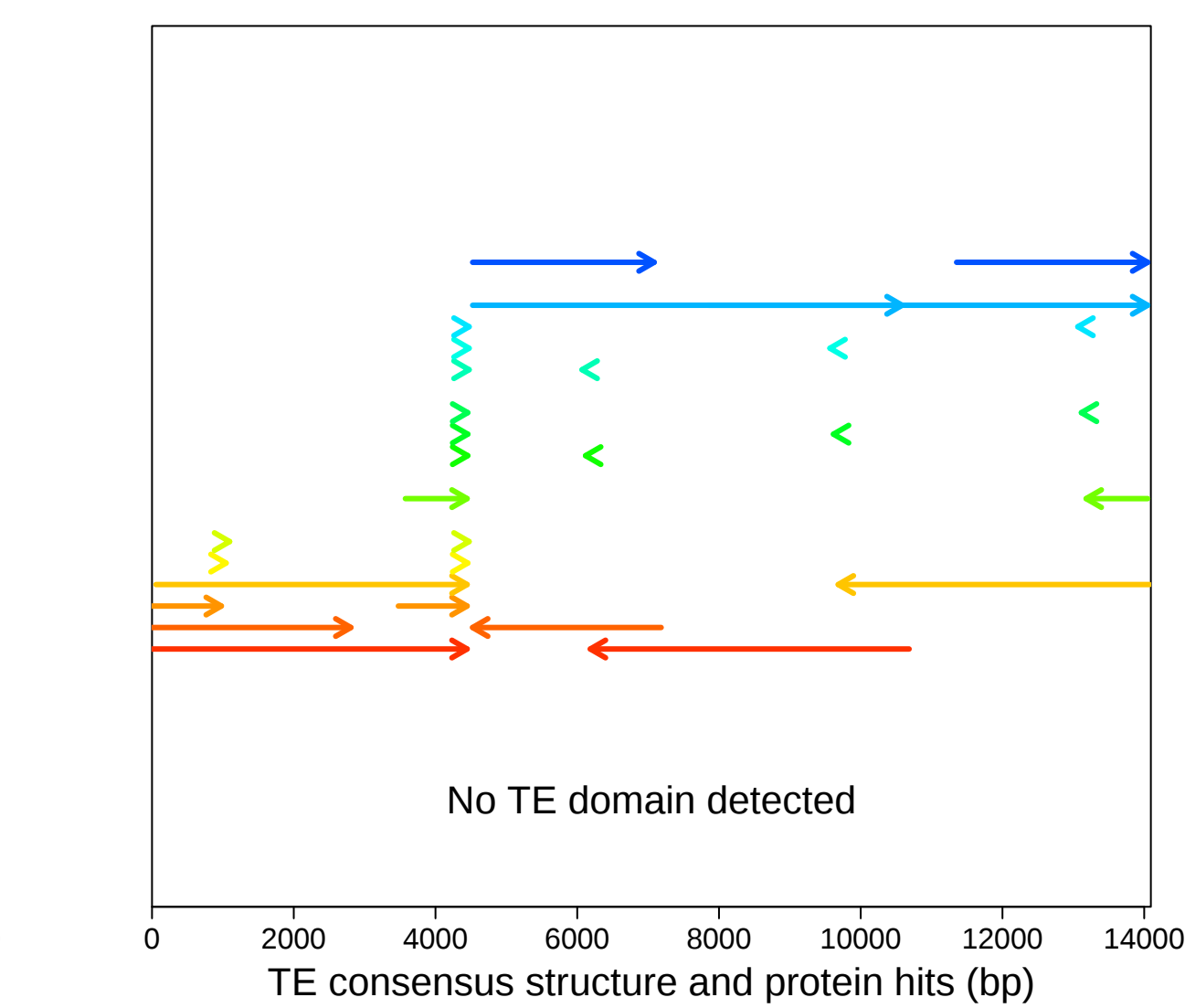
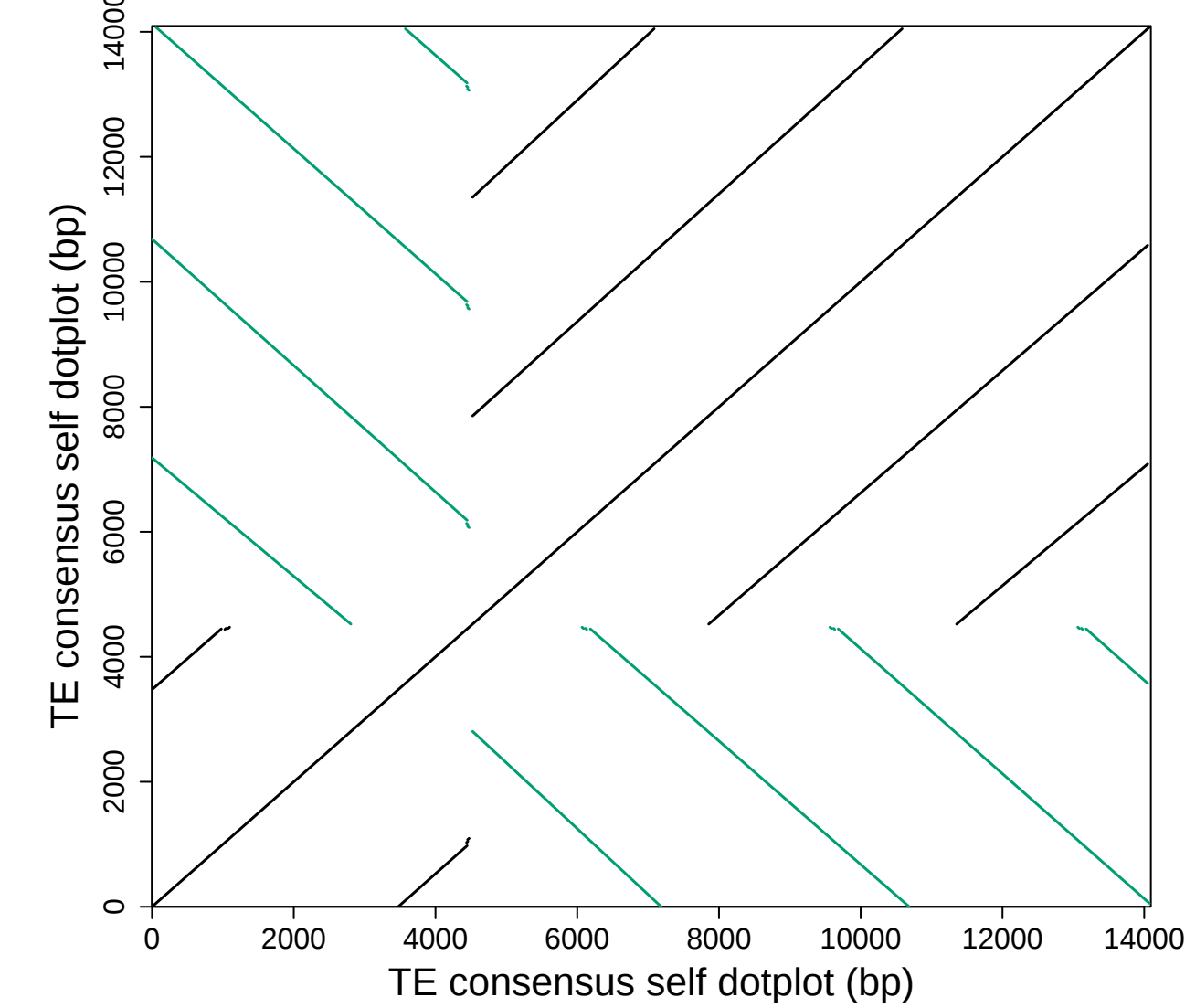
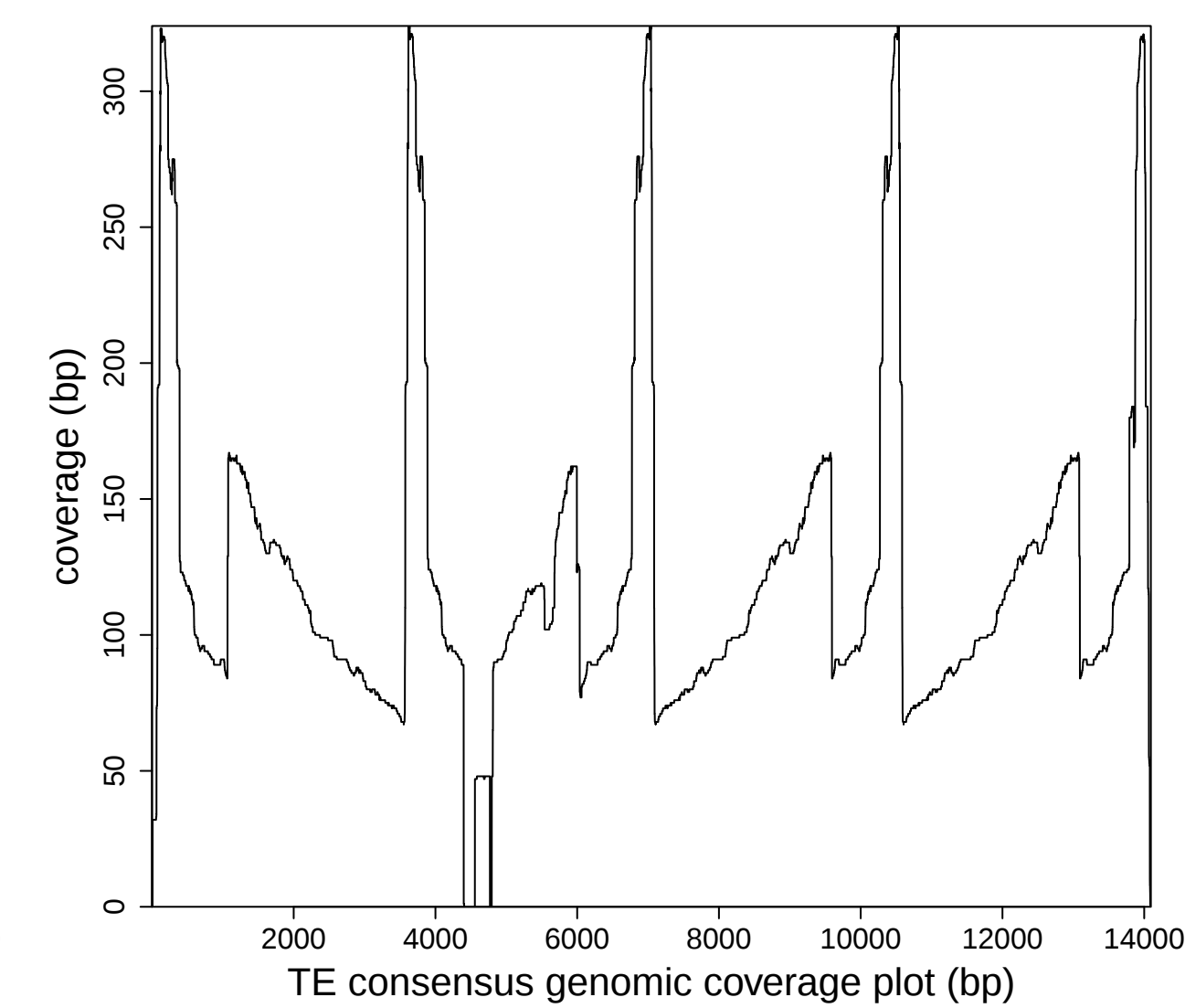
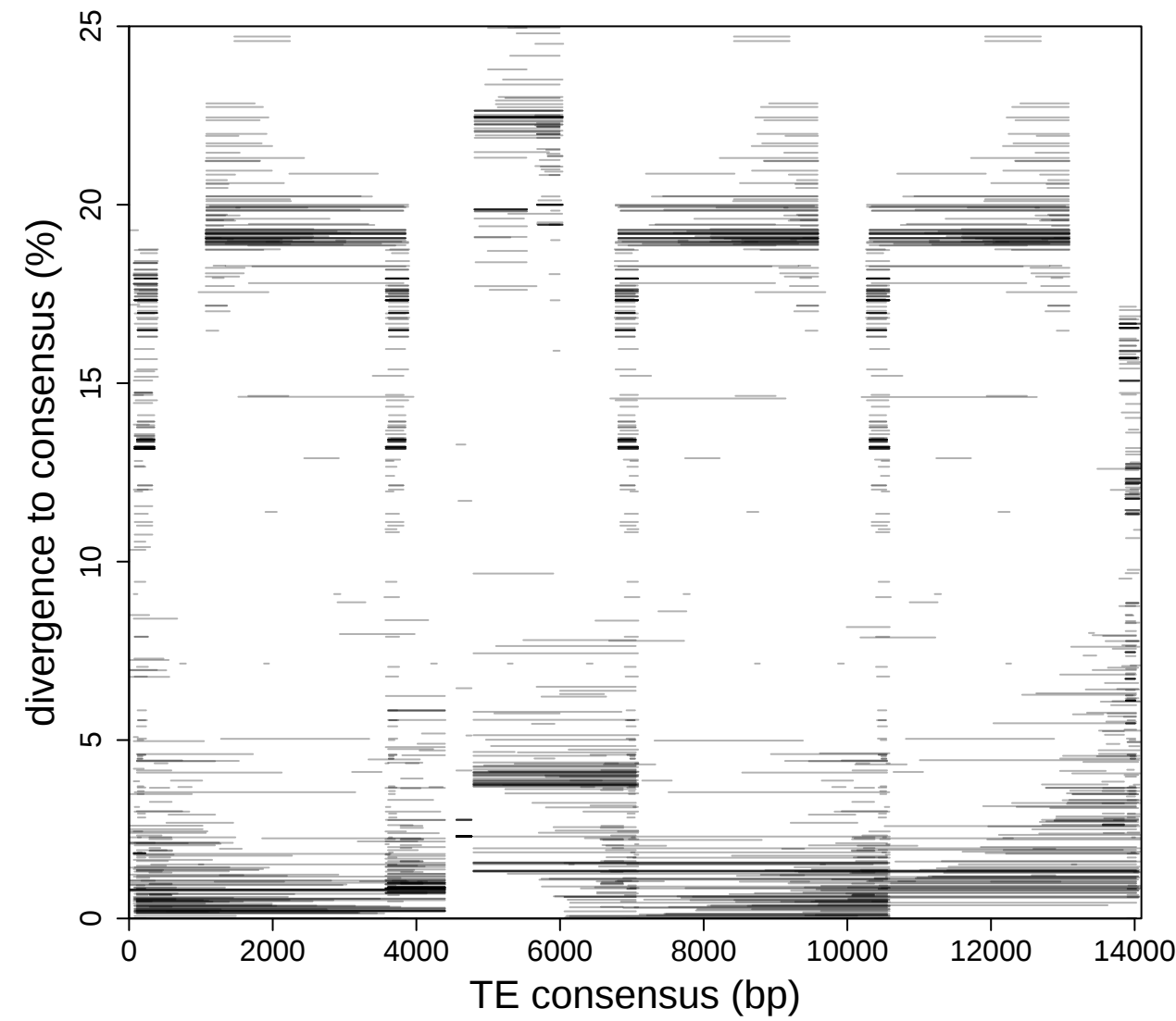






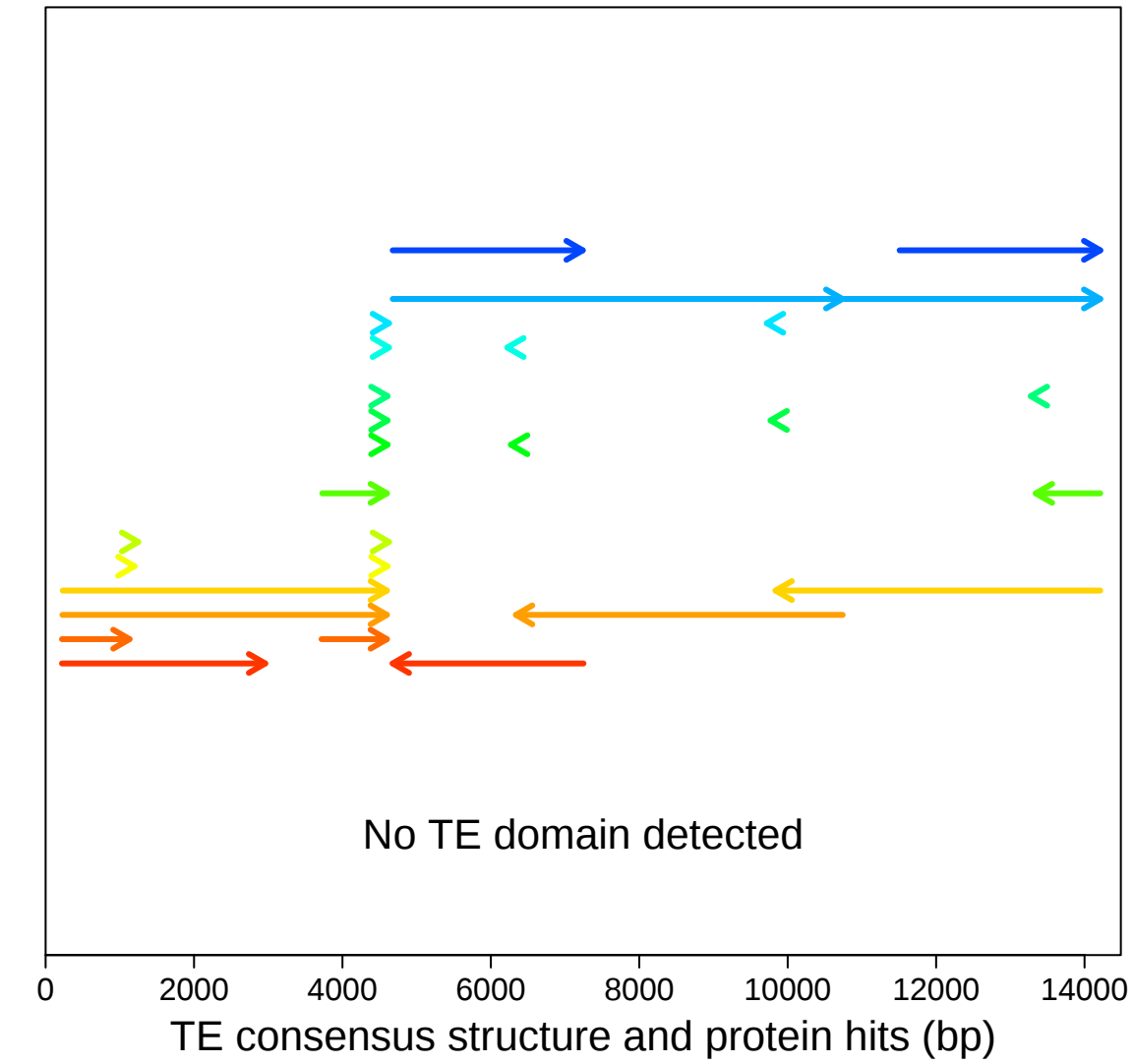
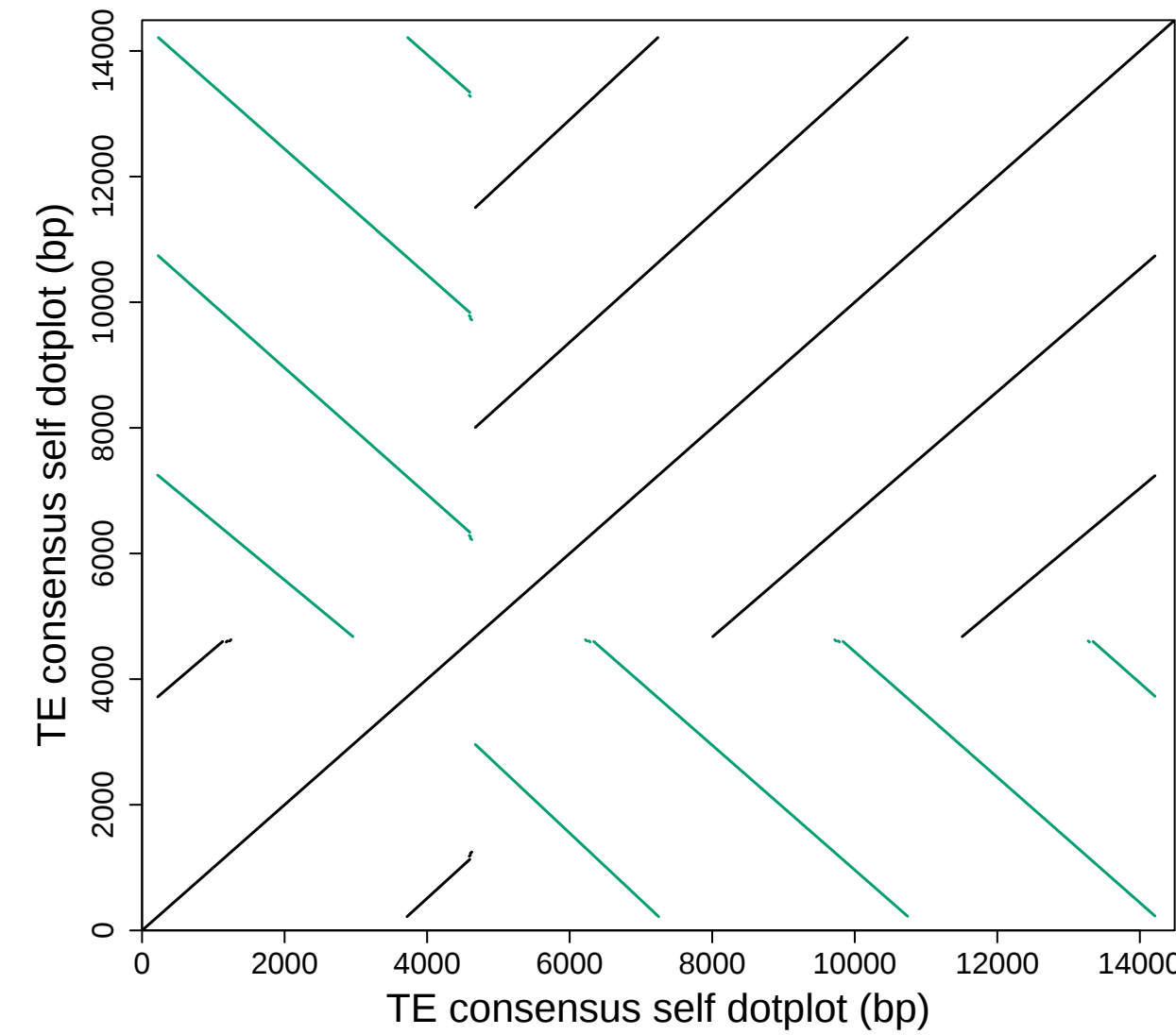
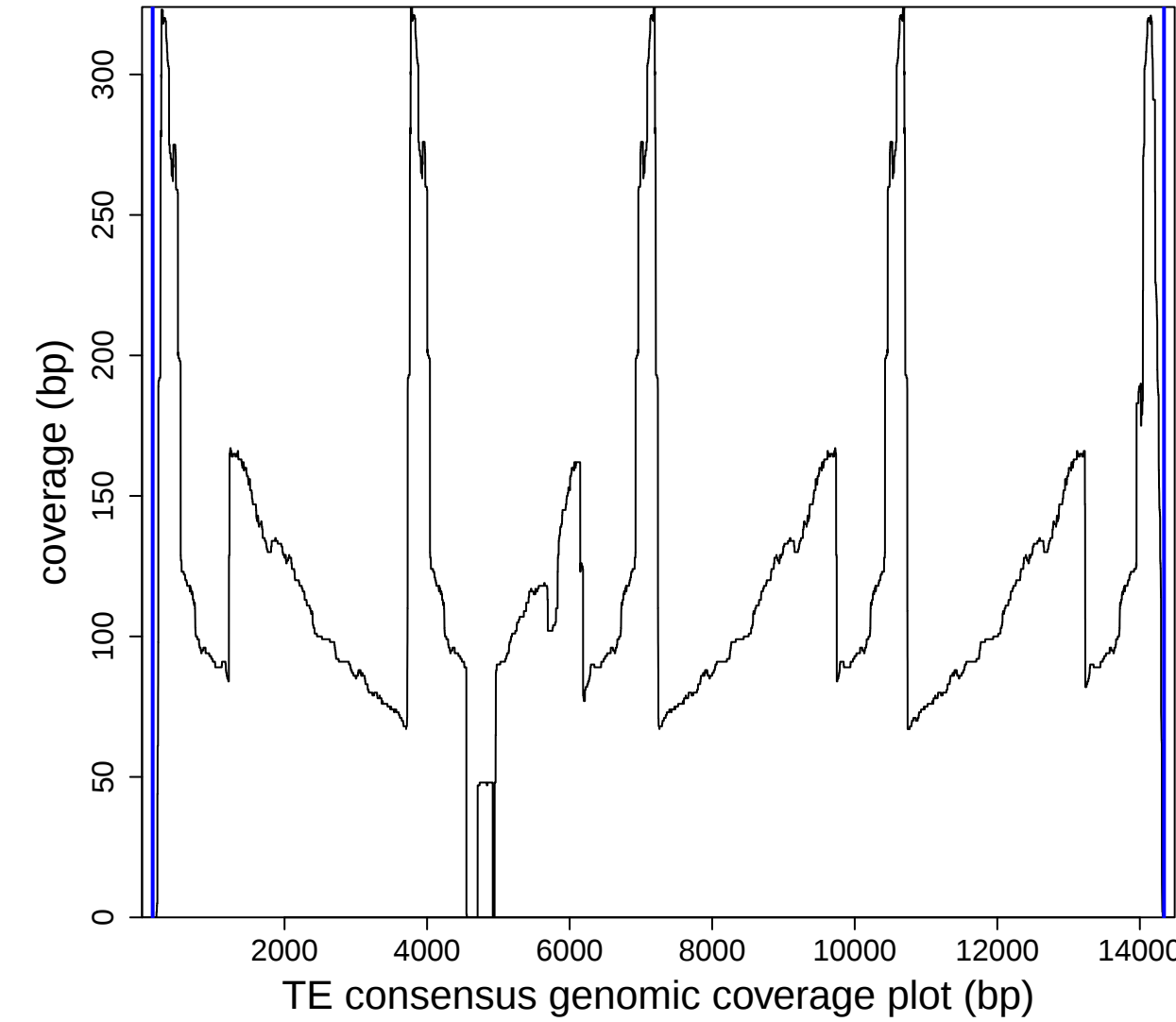
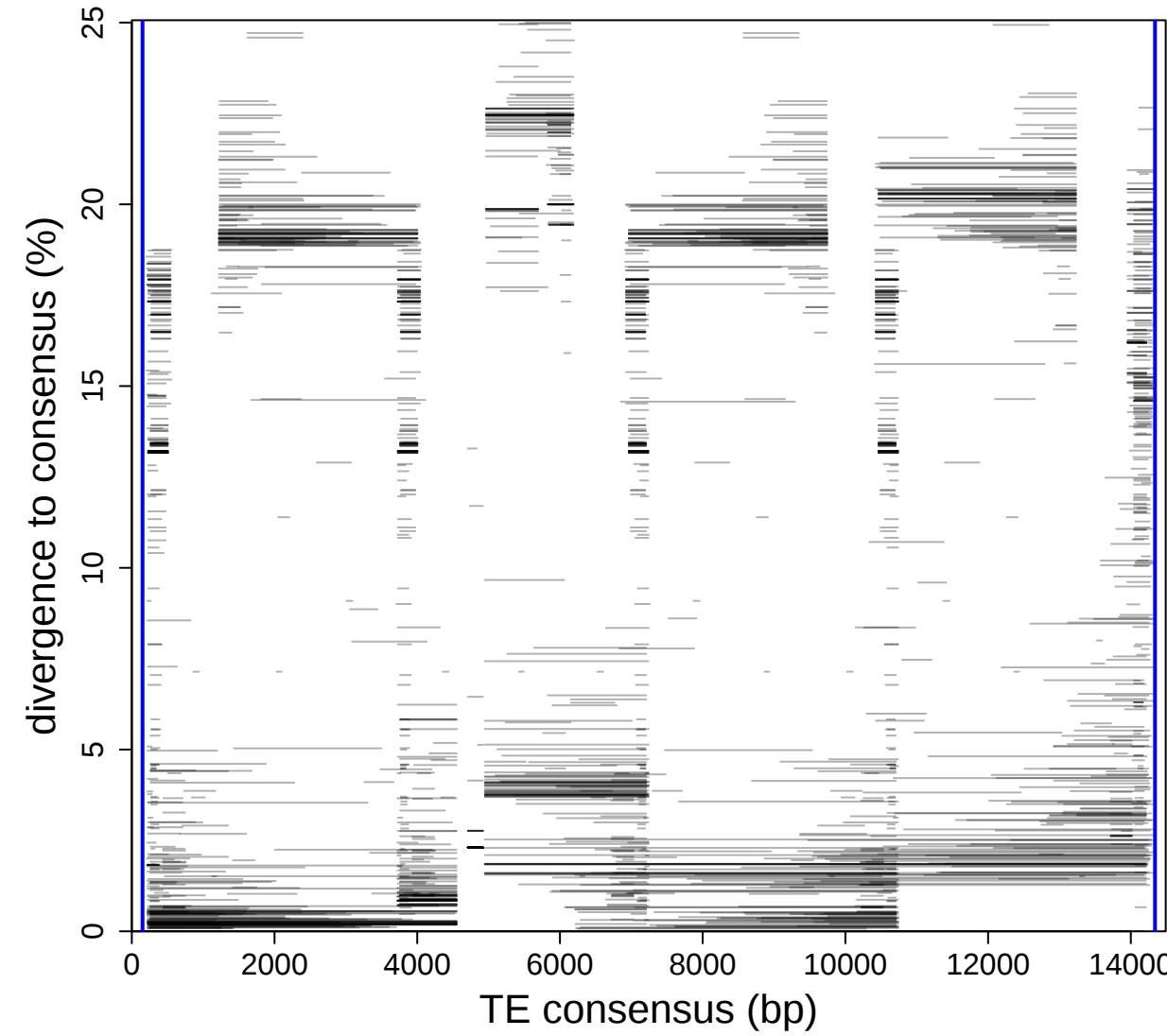
bed\_uf.bed\_g\_1.bed\_fm\_1.bed\_0\_0\_bcIn.fa\_aln.fa\_cl.fa\_gs.fa\_c  
size: 14094bp; fragments: 2100; full length: 0 ( $\geq 12684.6$ bp)

### After TEtrimmer 14094 bp



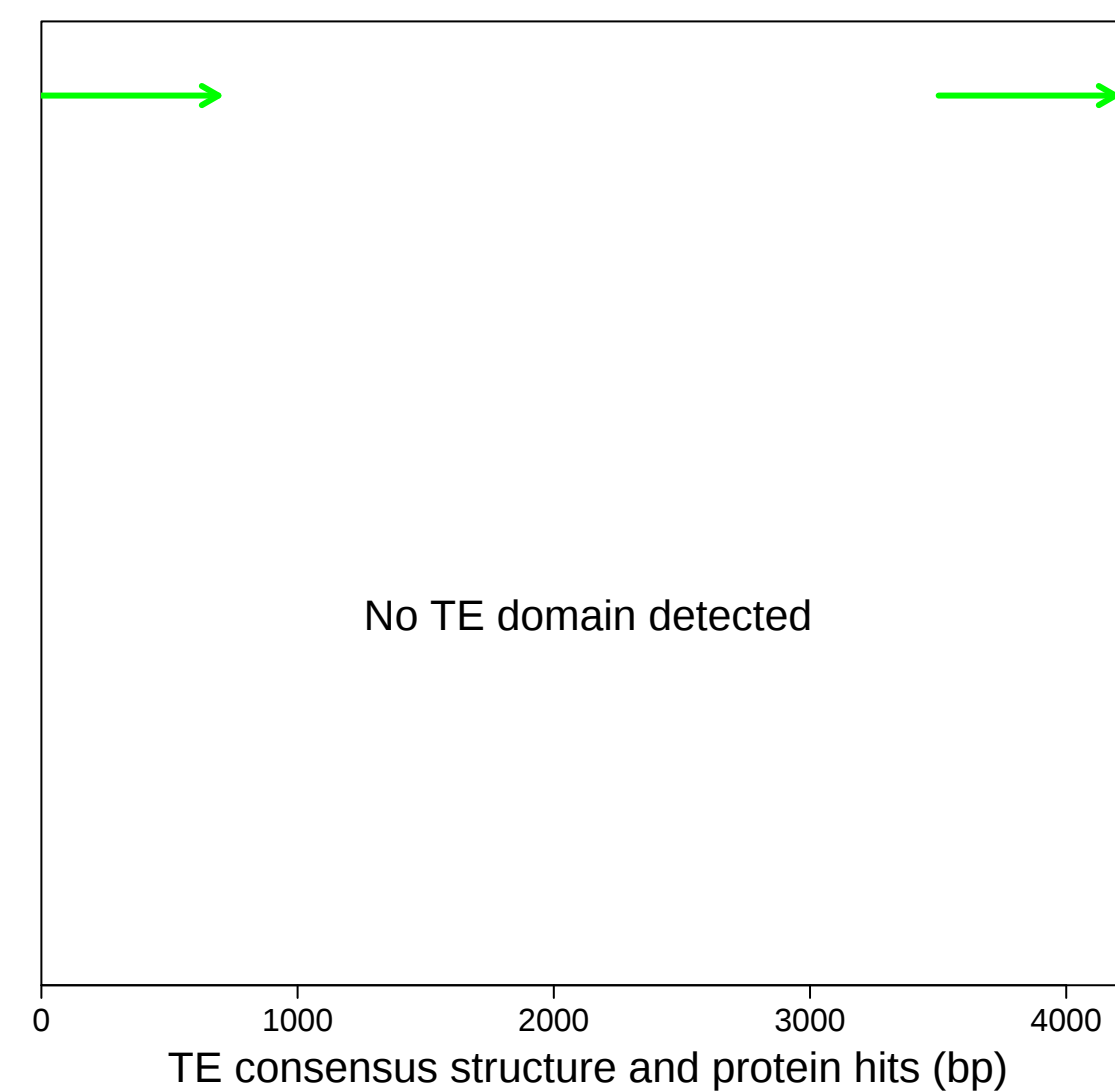
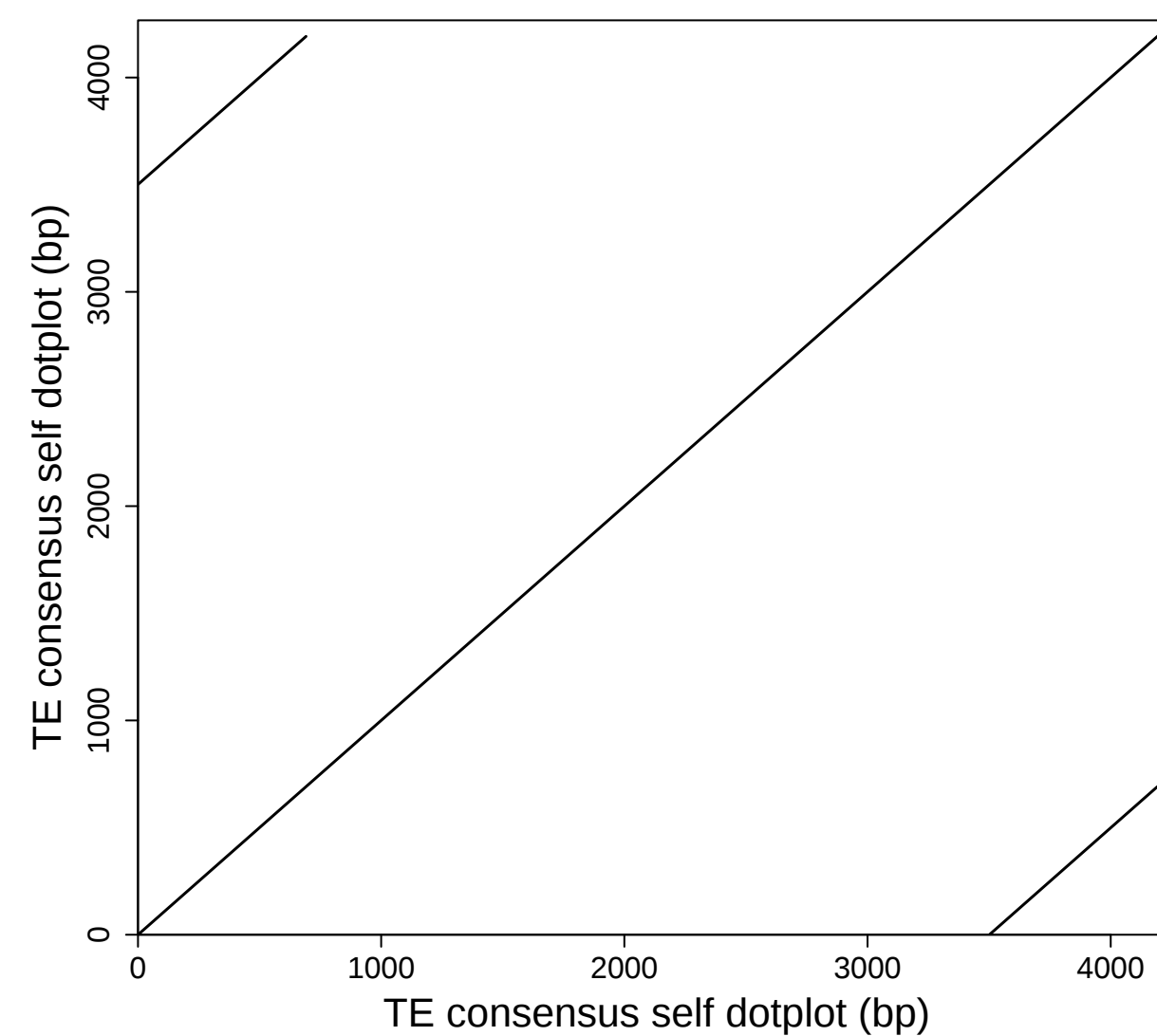
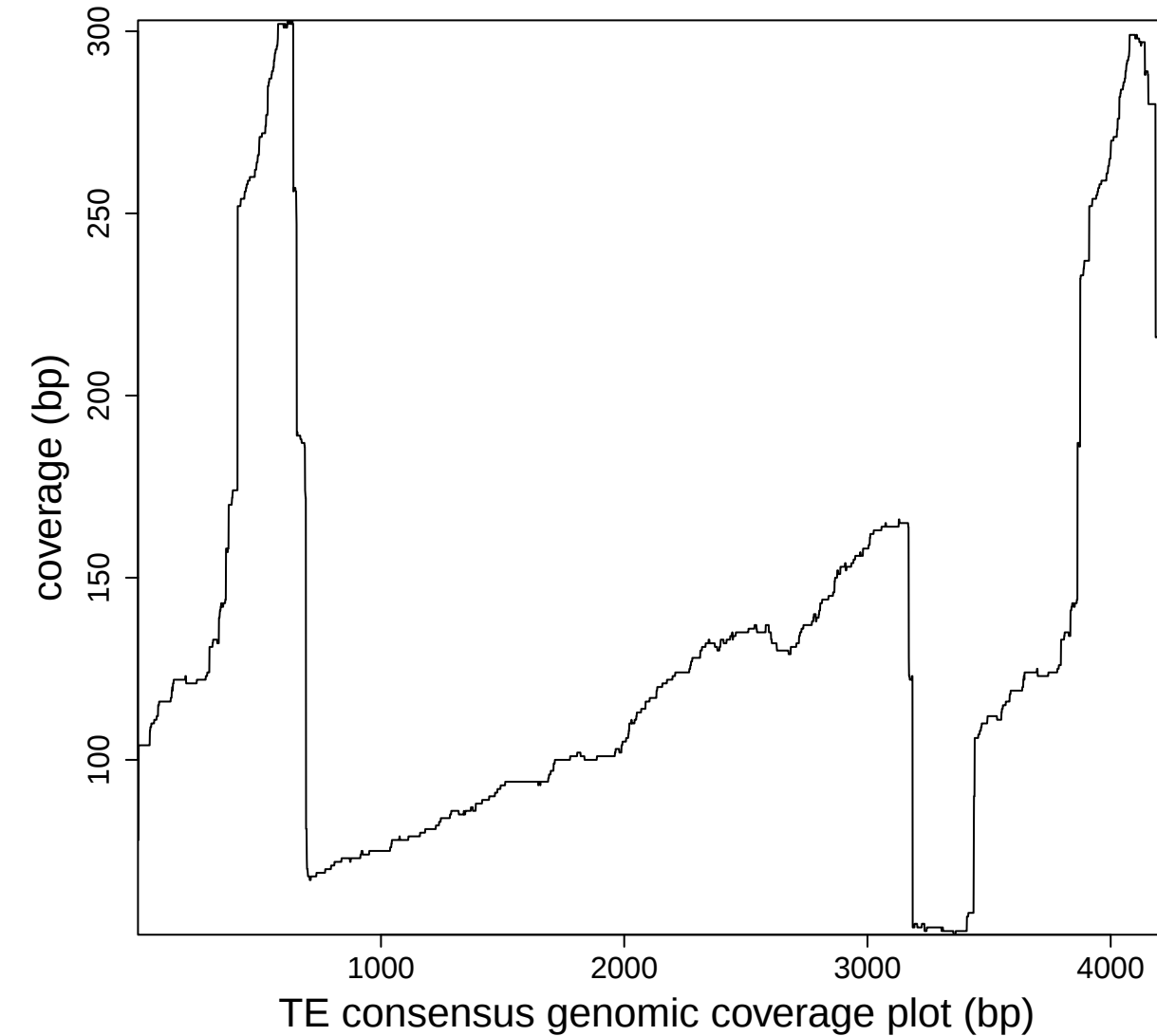
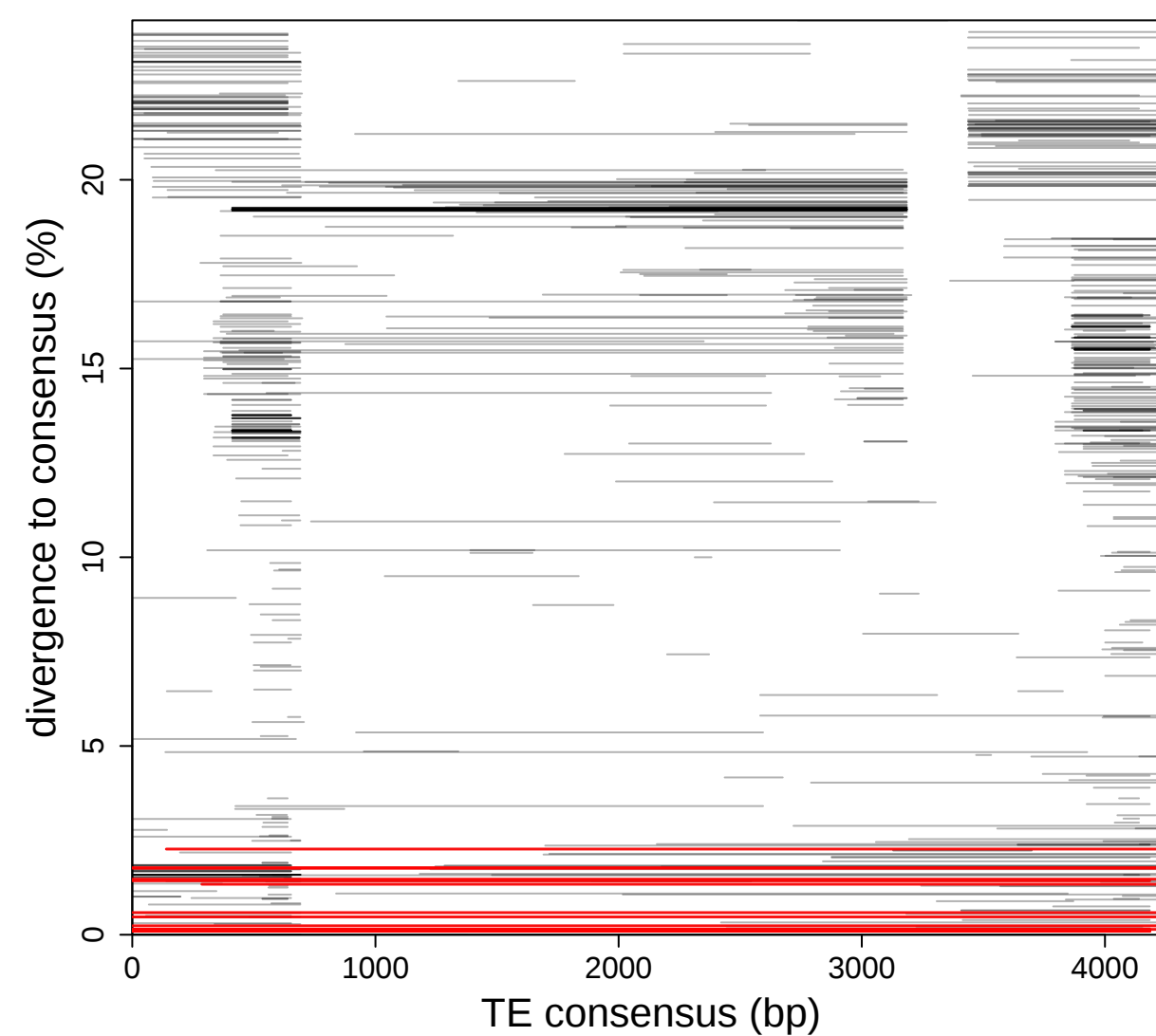
47.fasta.b.bed\_uf.bed\_g\_1.bed\_fm\_1.bed\_0\_0\_bcln.fa\_aln.fa\_cl.f  
size: 14489bp; fragments: 2100; full length: 0 (>=13040.1bp)

## After TEtrimmer Extended plot Blue lines are boundaries

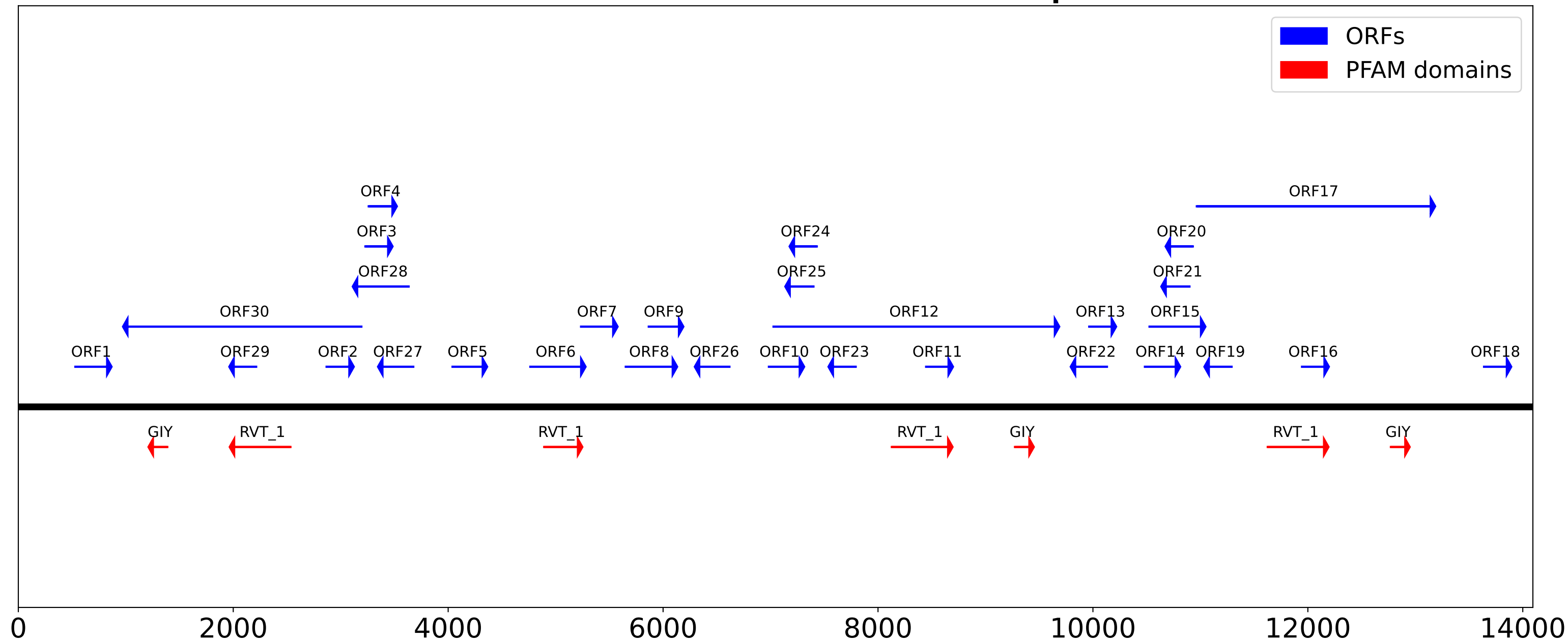


TE: rnd\_5\_family\_10547  
size: 4267bp; fragments: 723; full length: 16 ( $\geq 3840.3$ bp)

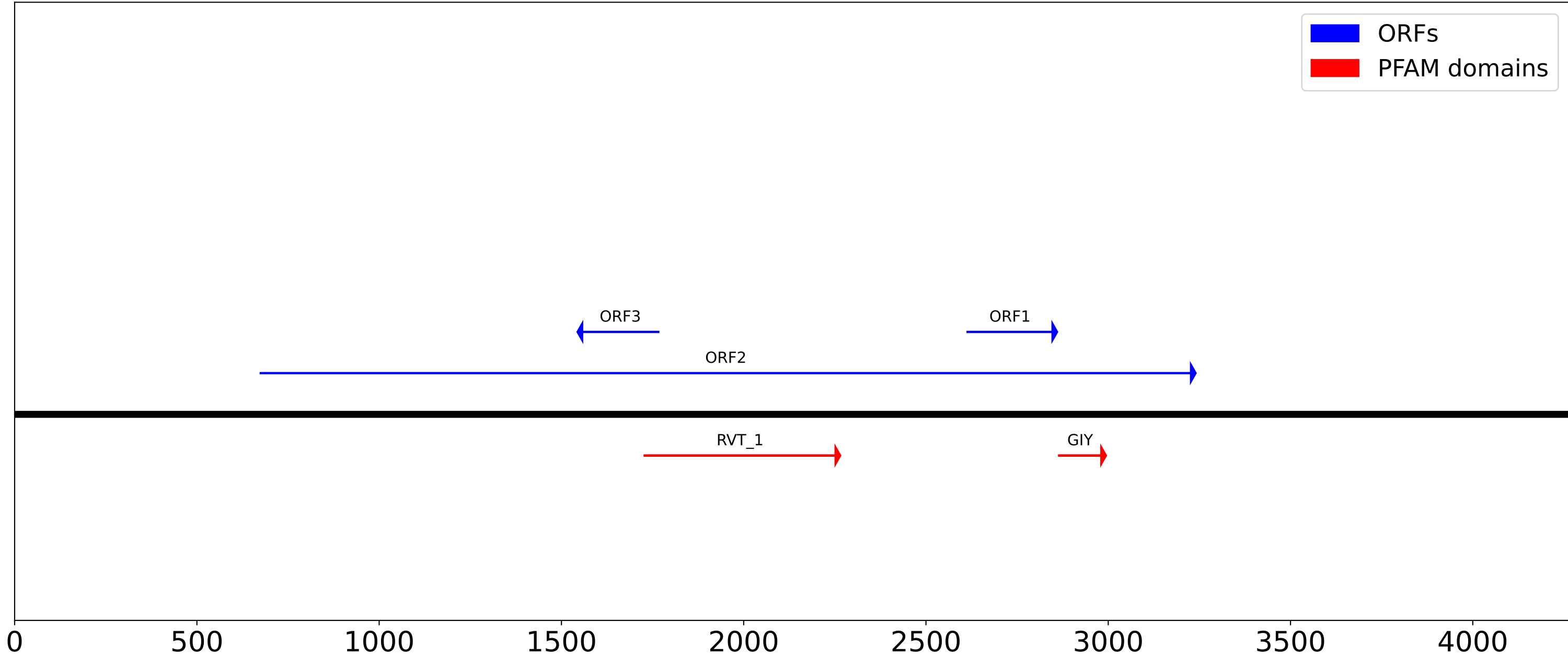
## Before TEtrimmer 4267 bp



## After TEtrimmer ORF and PFAM domain plot



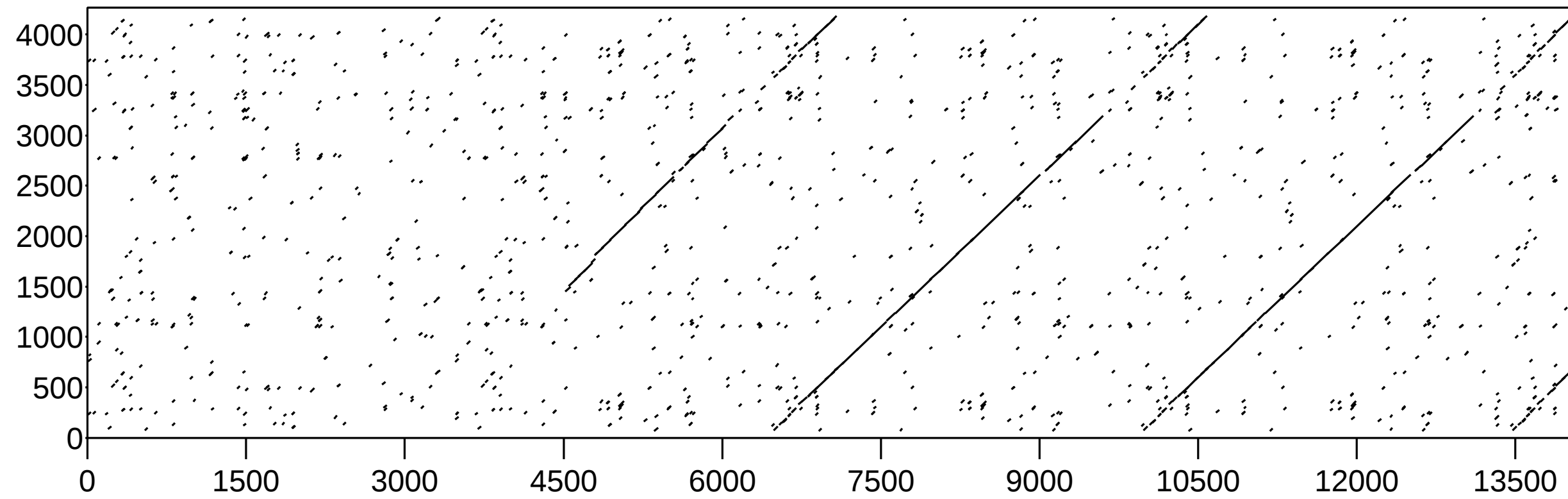
**Before TEtrimmer ORF and PFAM domain plot**



# Dotplot

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

: consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)