

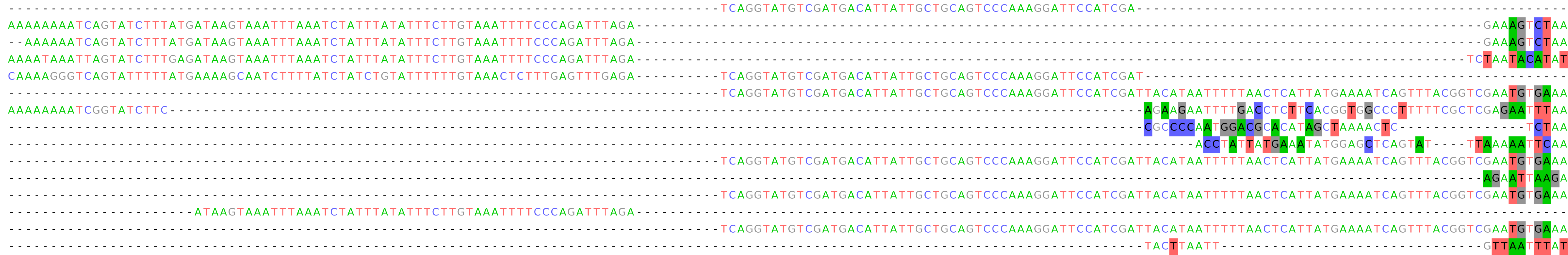
Start crop Point

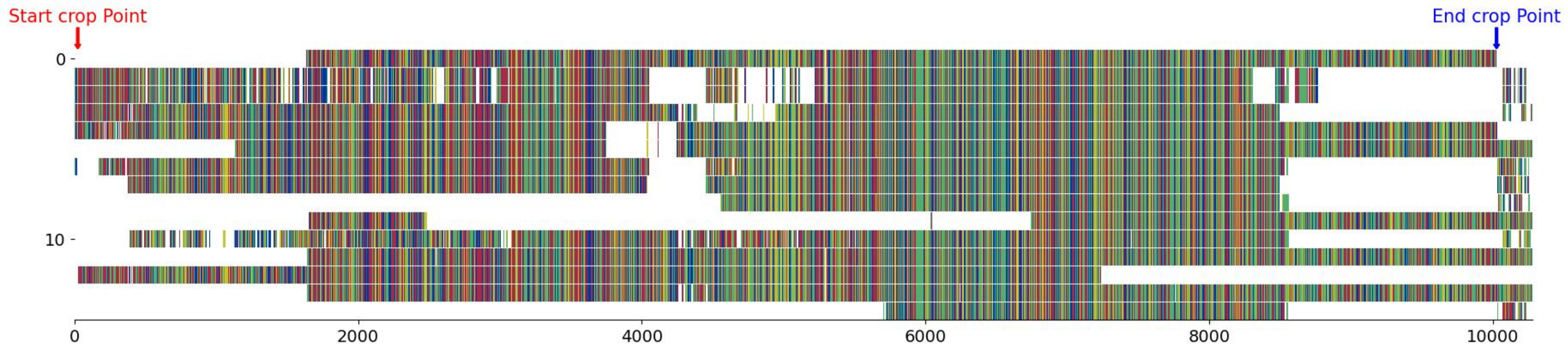


End crop Point



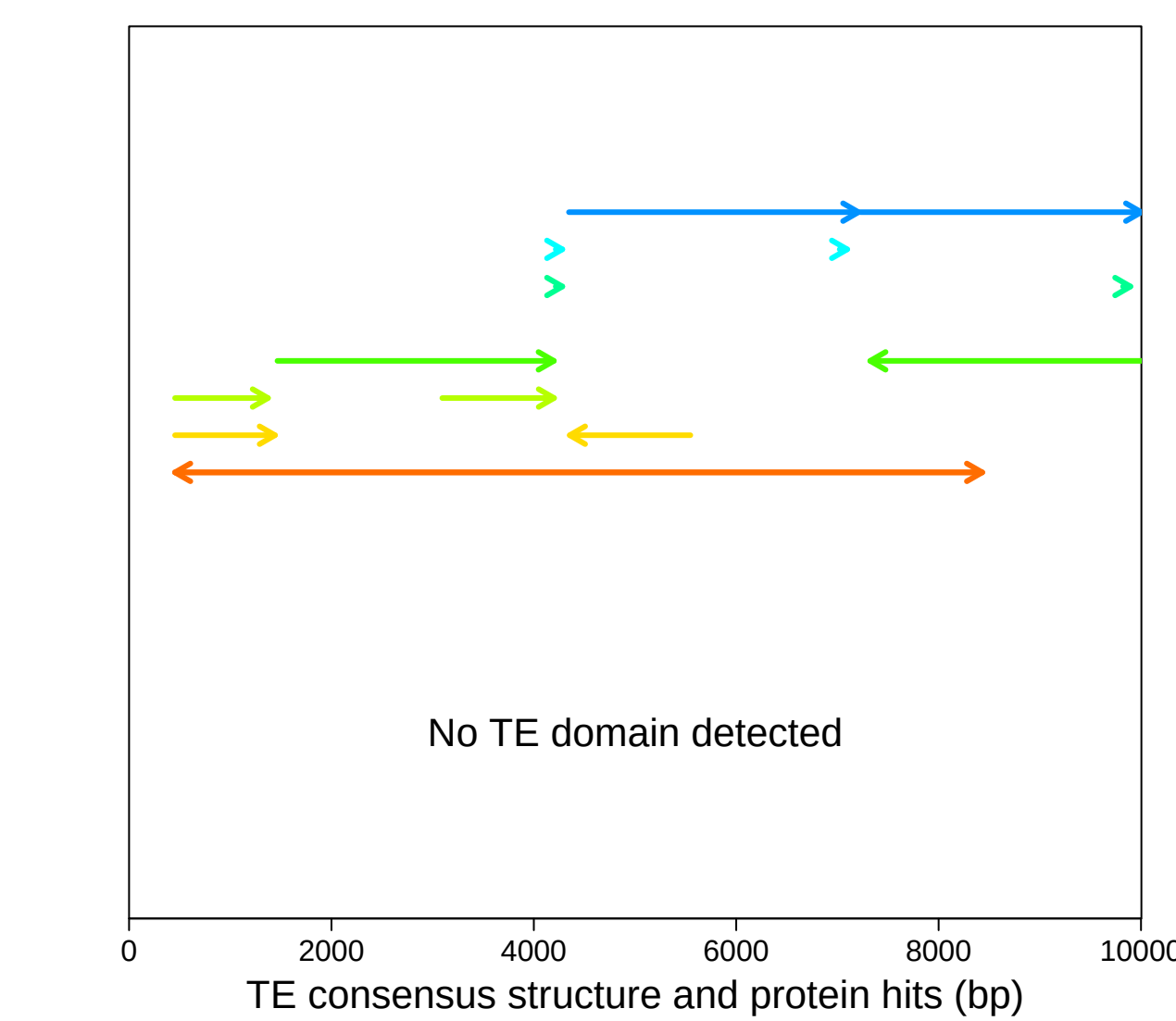
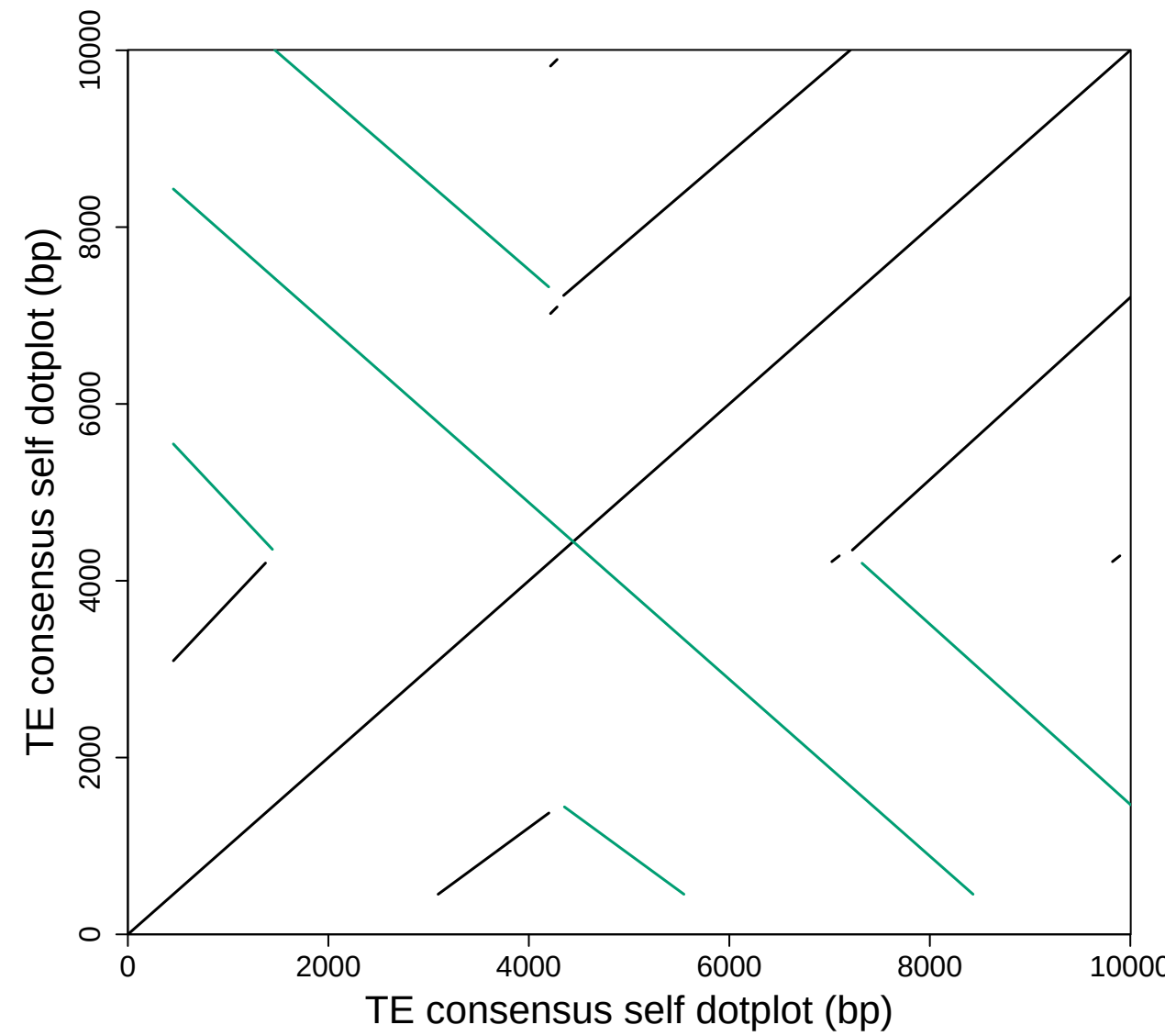
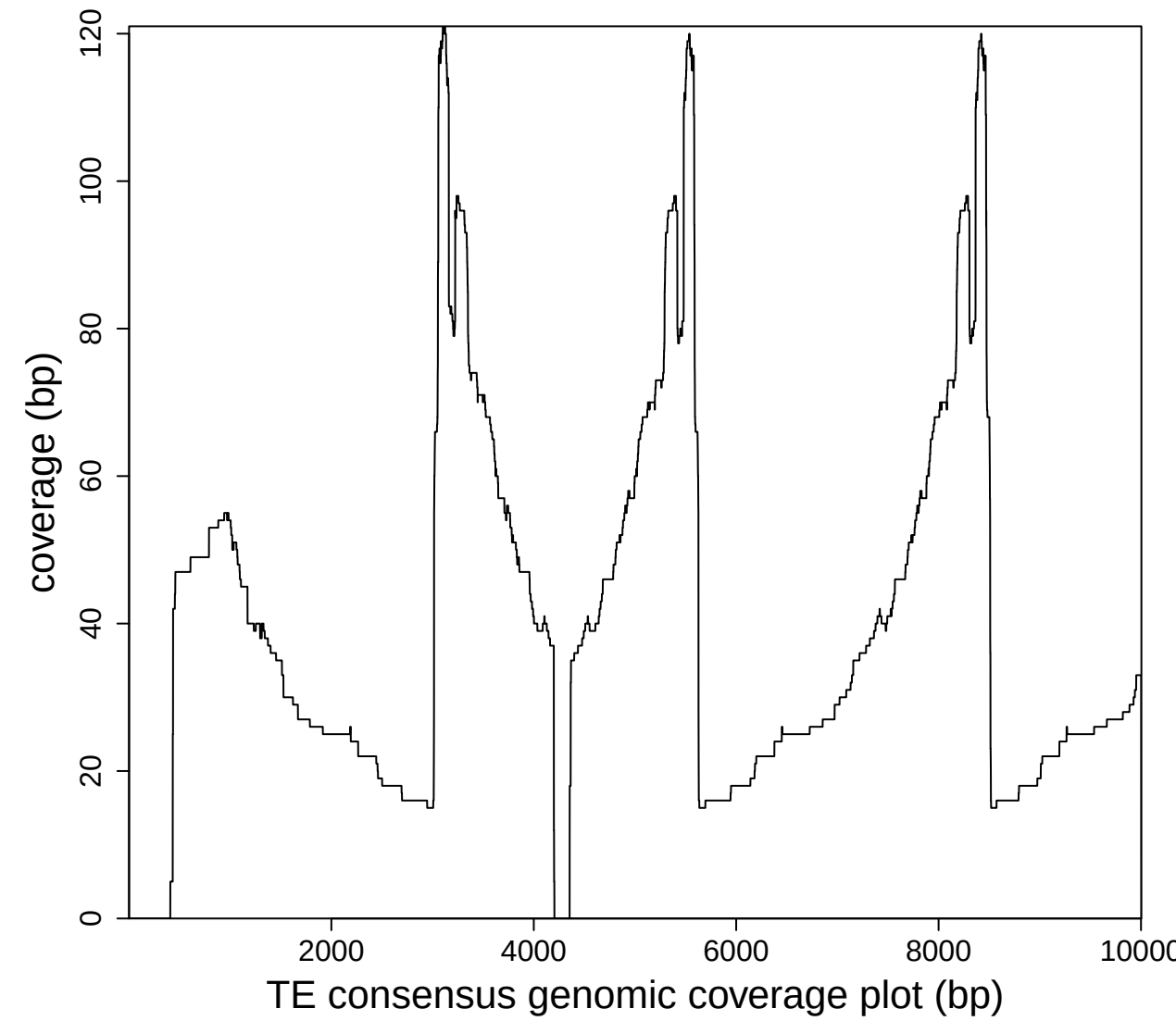
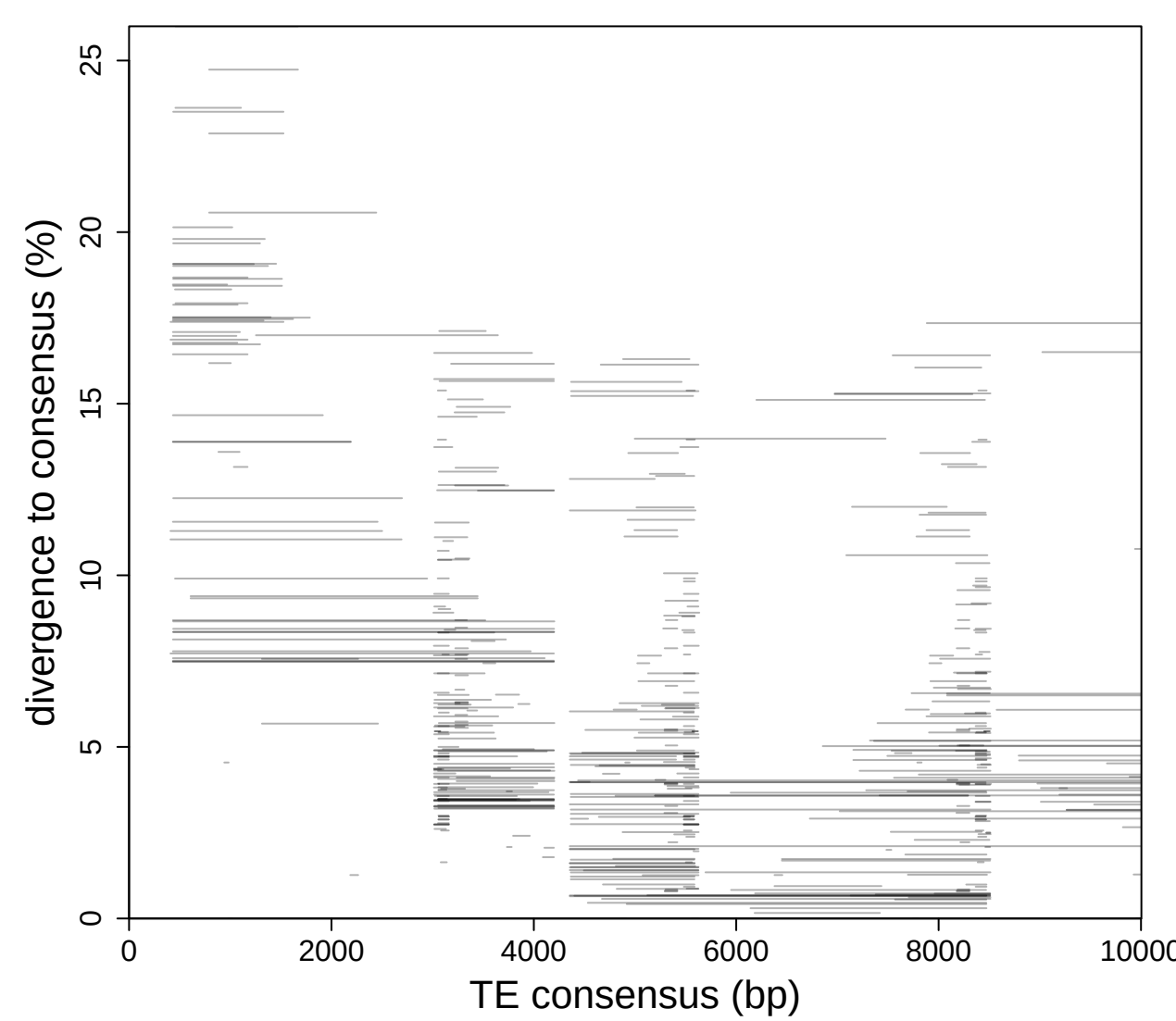
MSA length = 10004





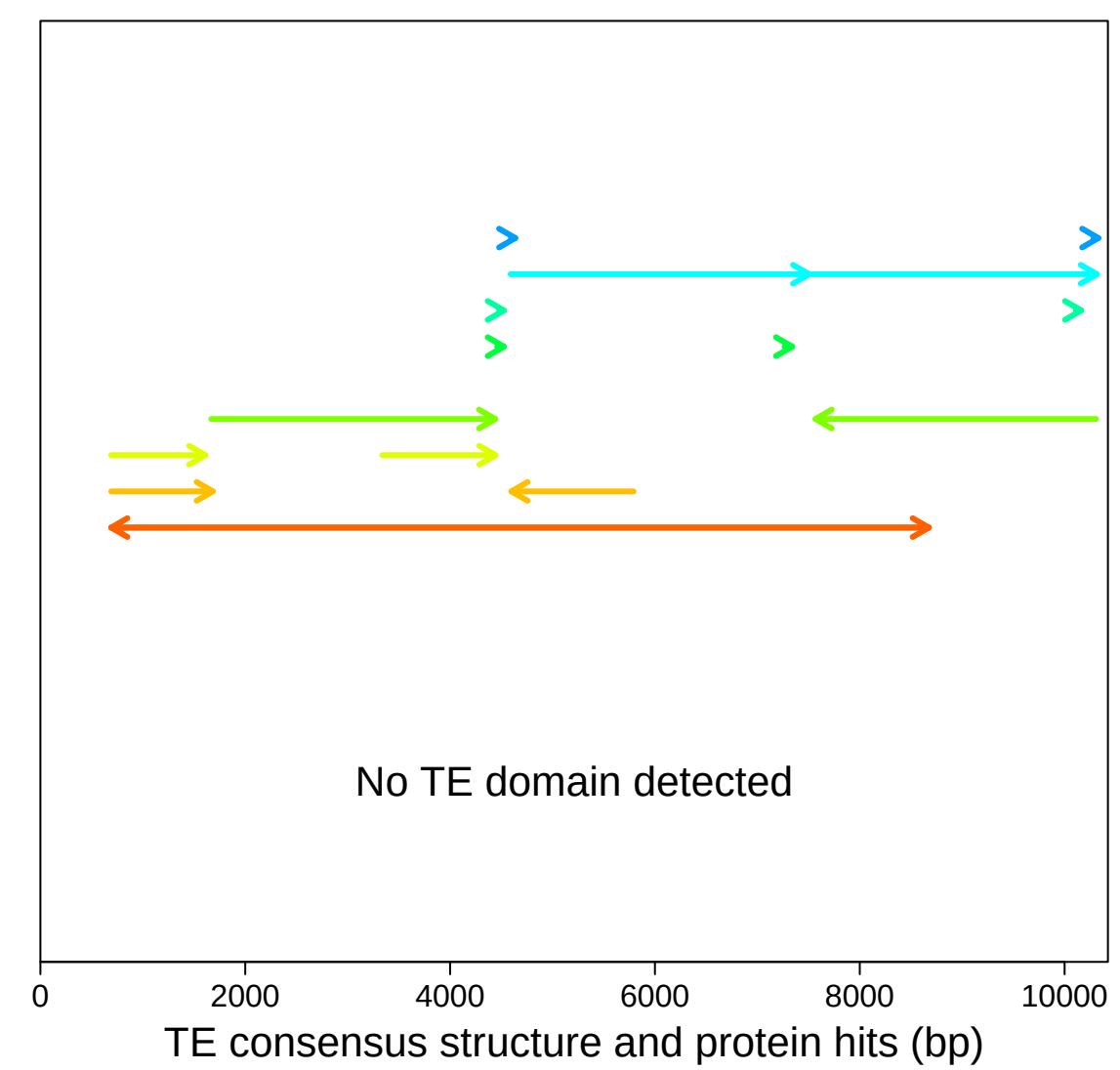
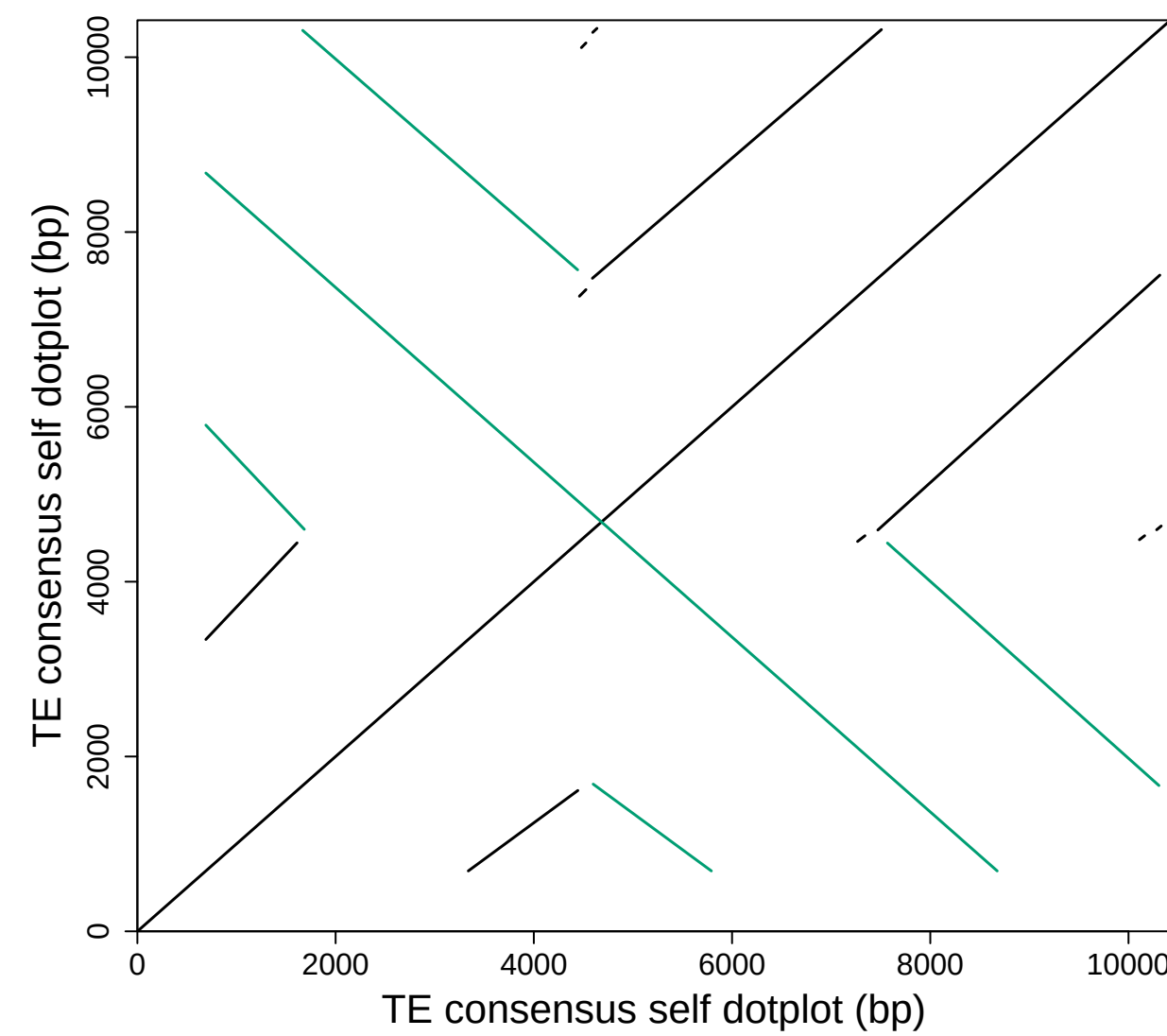
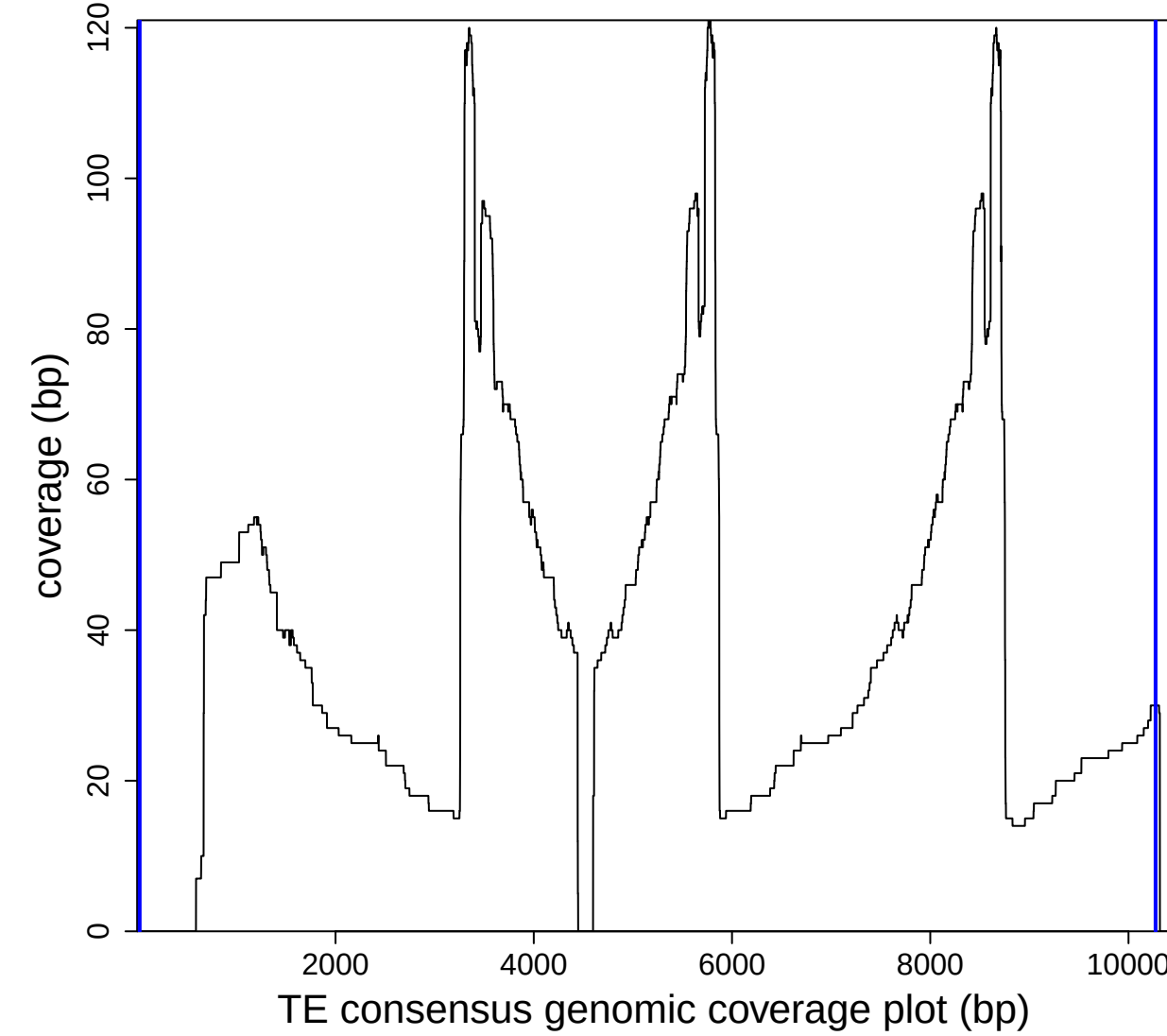
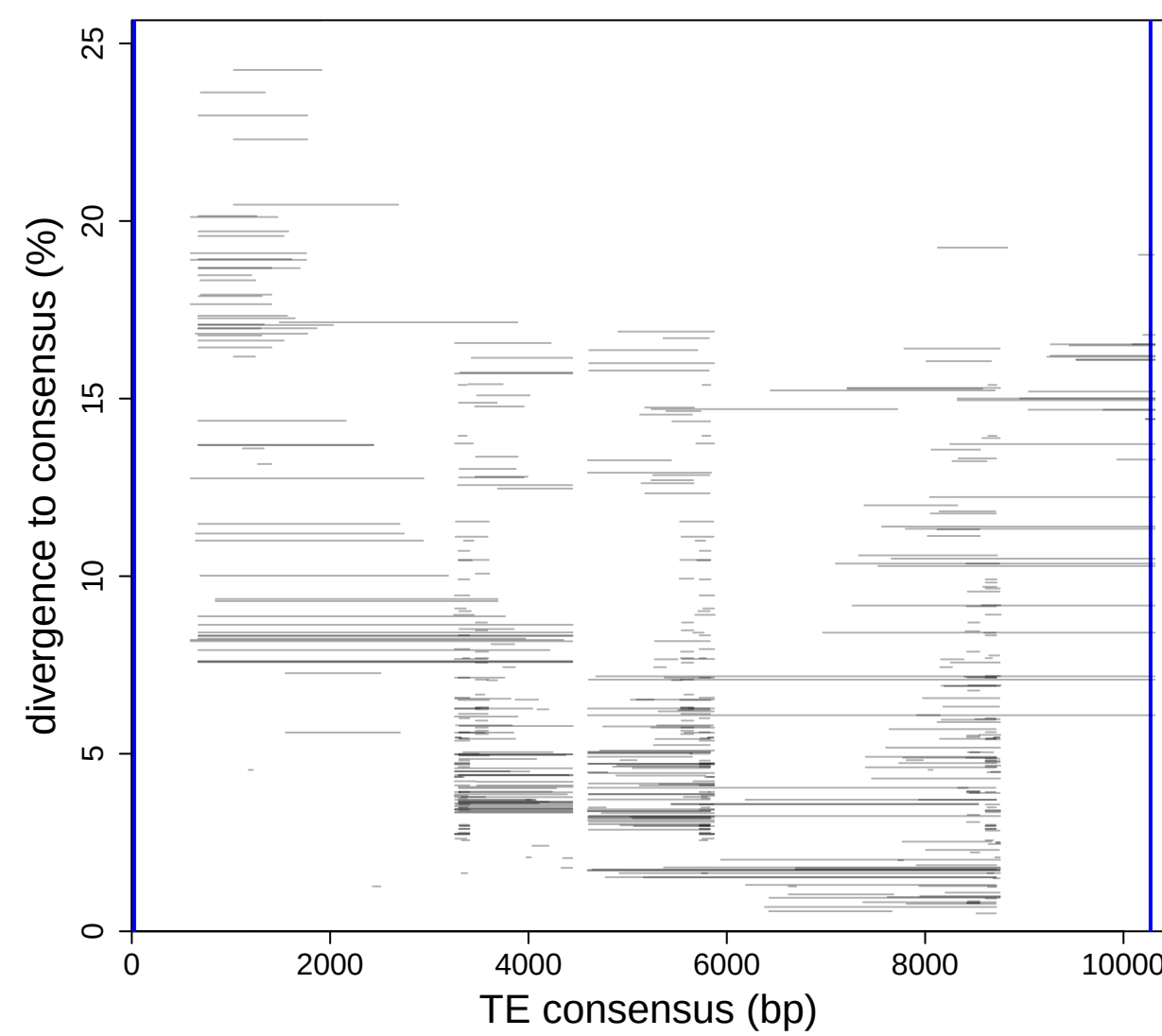
bed uf.bed g 3.bed fm 1.bed 0 0 bcln.fa aln.fa cl.fa gs.fa ce
size: 10004bp; fragments: 529; full length: 0 (>=9003.6bp)

After TEtrimmer 10004 bp

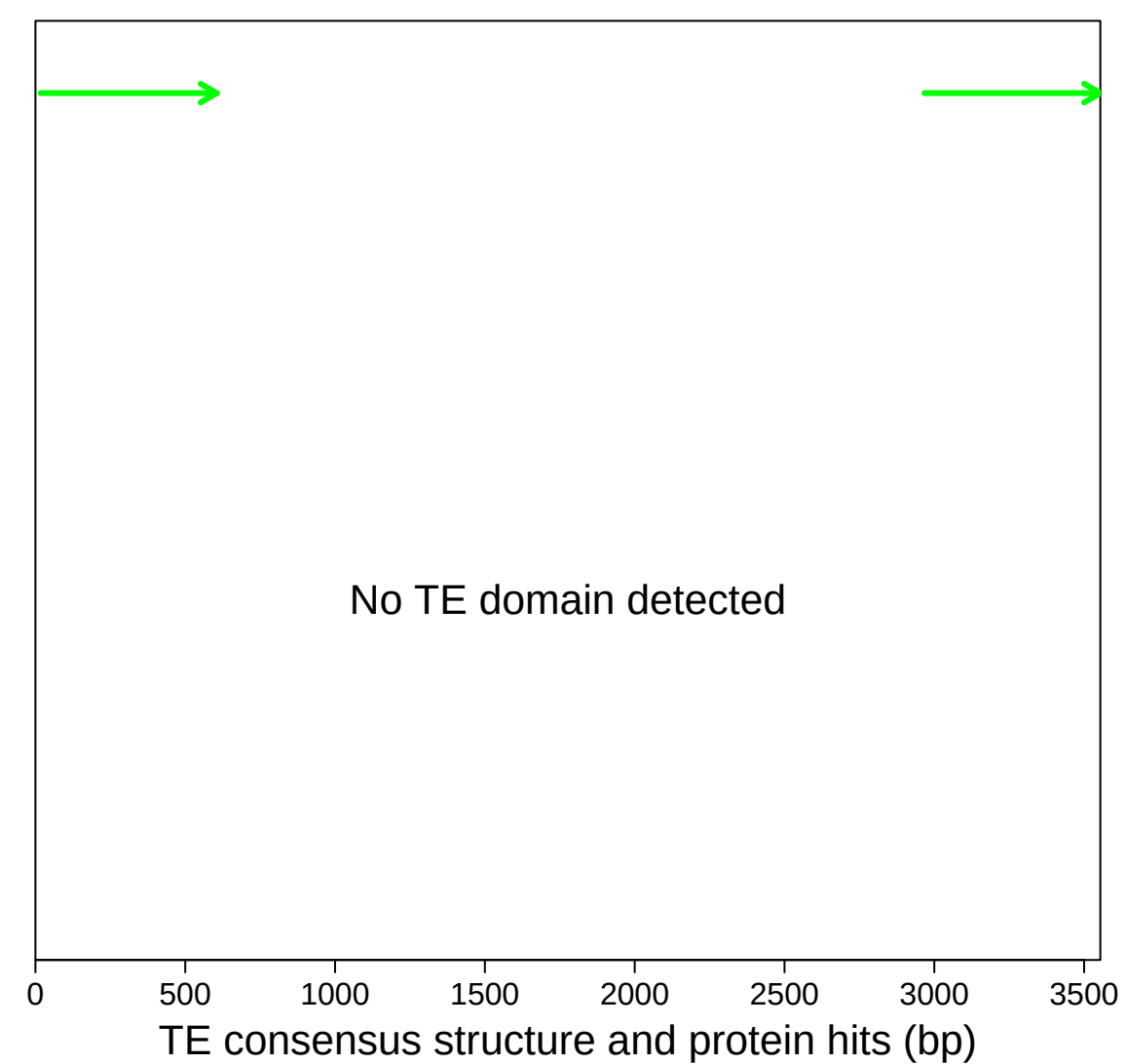
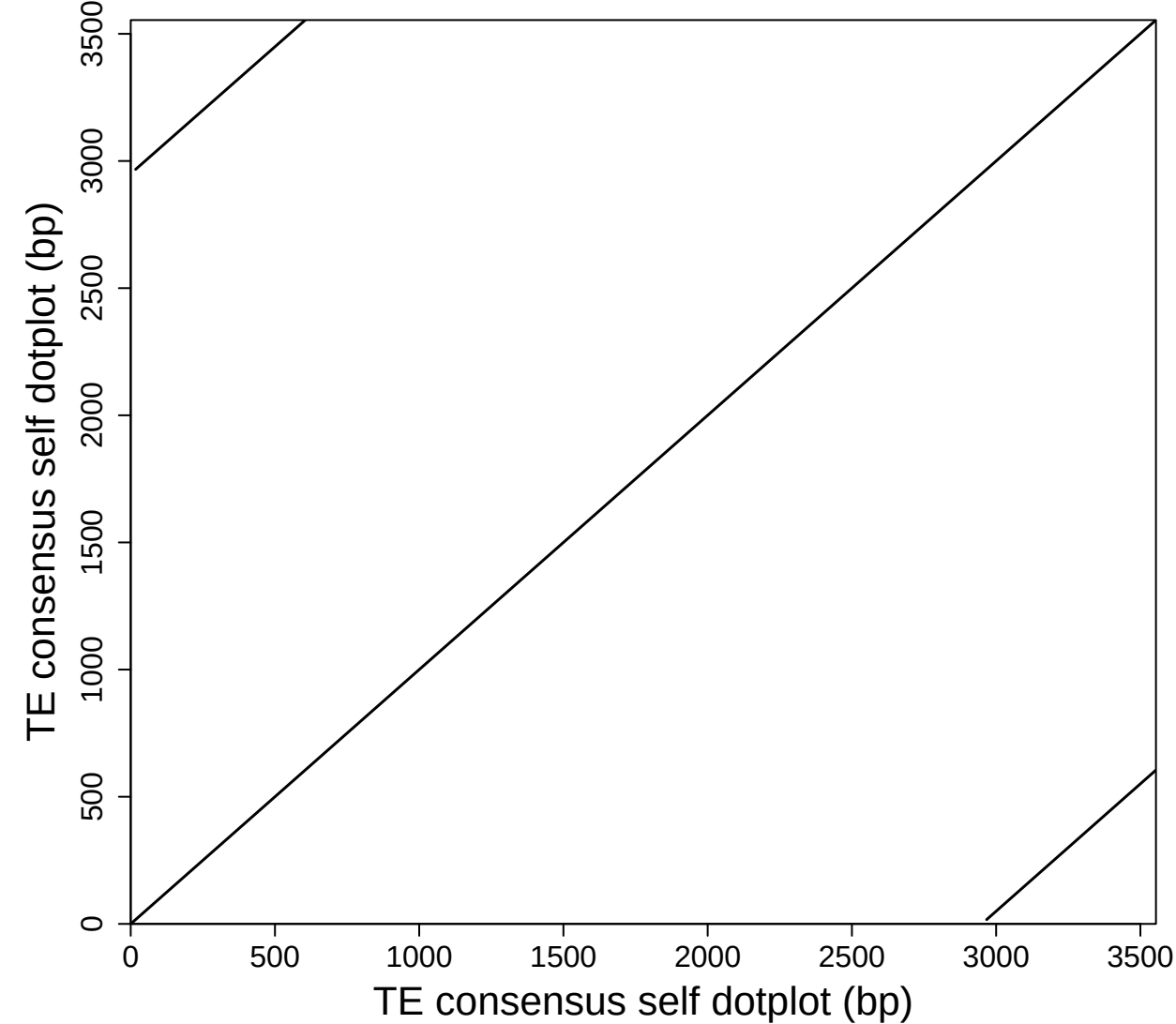
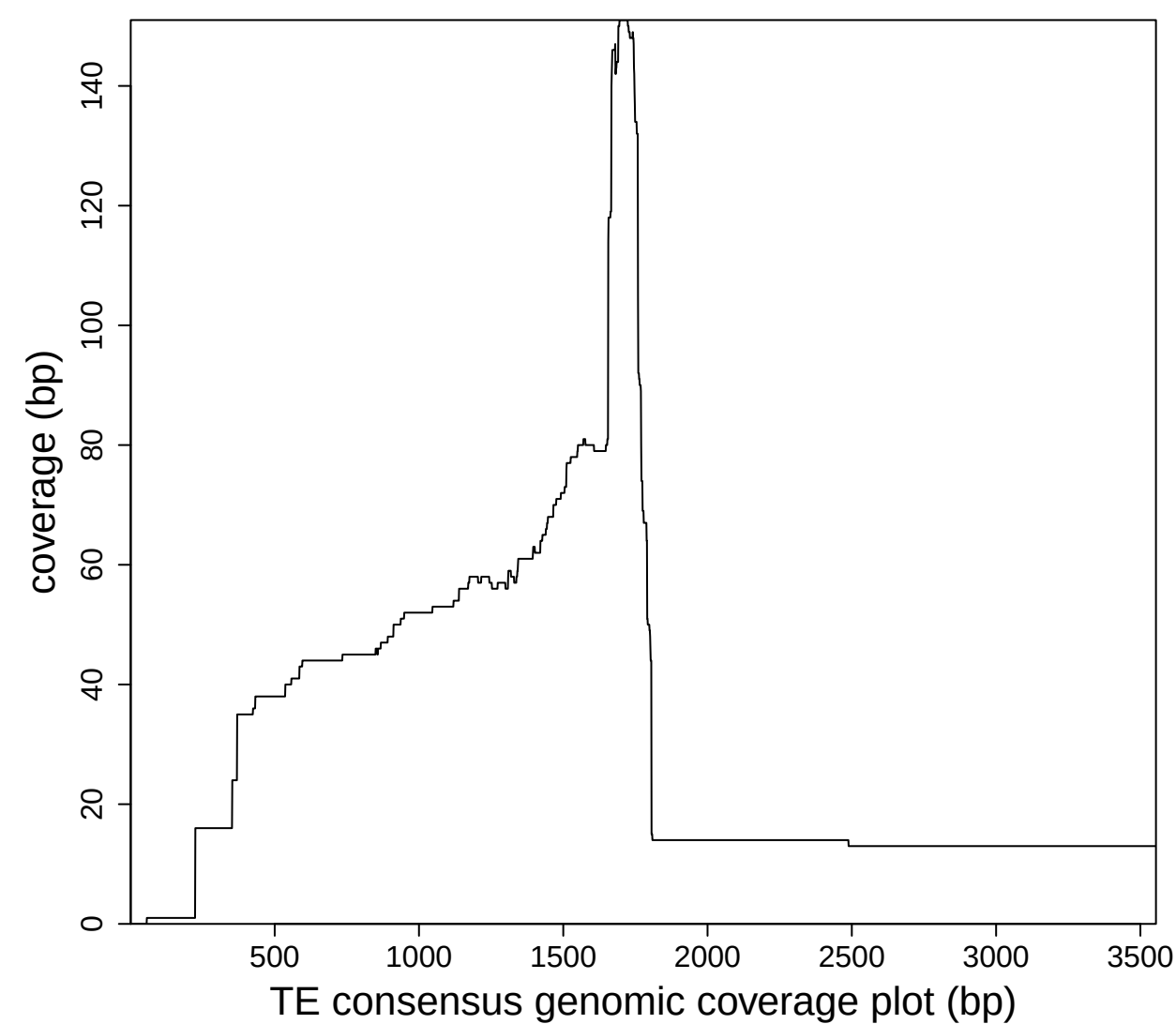
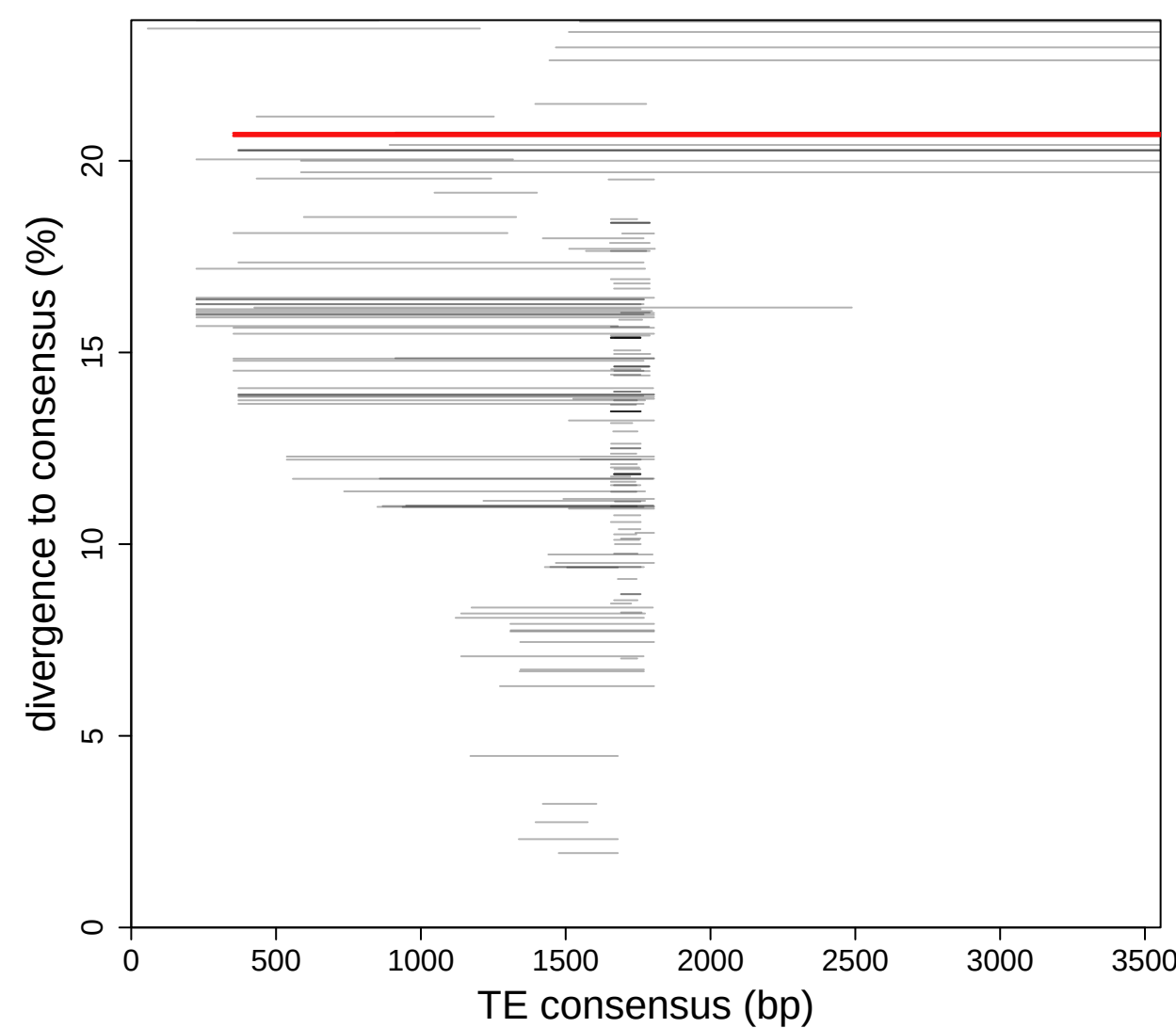


6.fasta.b.bed uf.bed g 3.bed fm 1.bed 0 0 bcln.fa aln.fa cl.fa
size: 10423bp; fragments: 525; full length: 0 (>=9380.7bp)

After TEtrimmer Extended plot Blue lines are boundaries

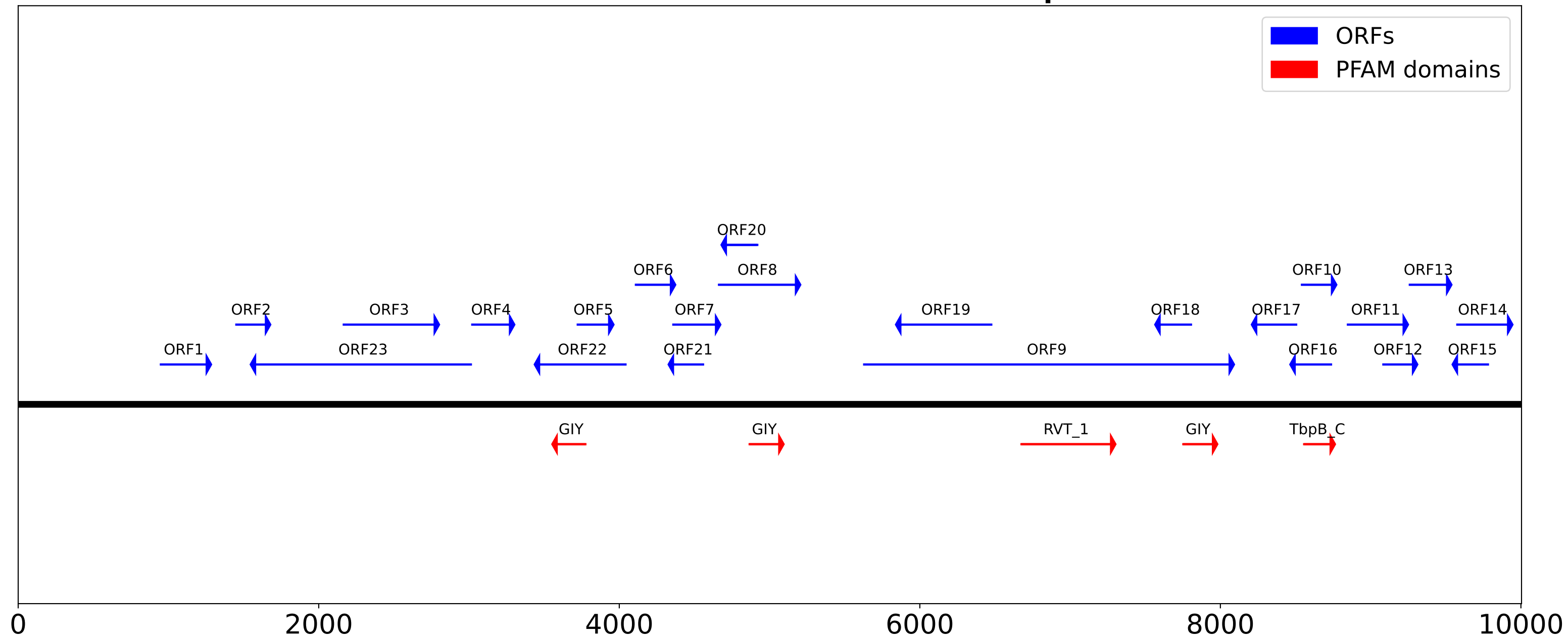


TE: rnd_5_family_946
size: 3554bp; fragments: 167; full length: 2 (≥ 3198.6 bp)

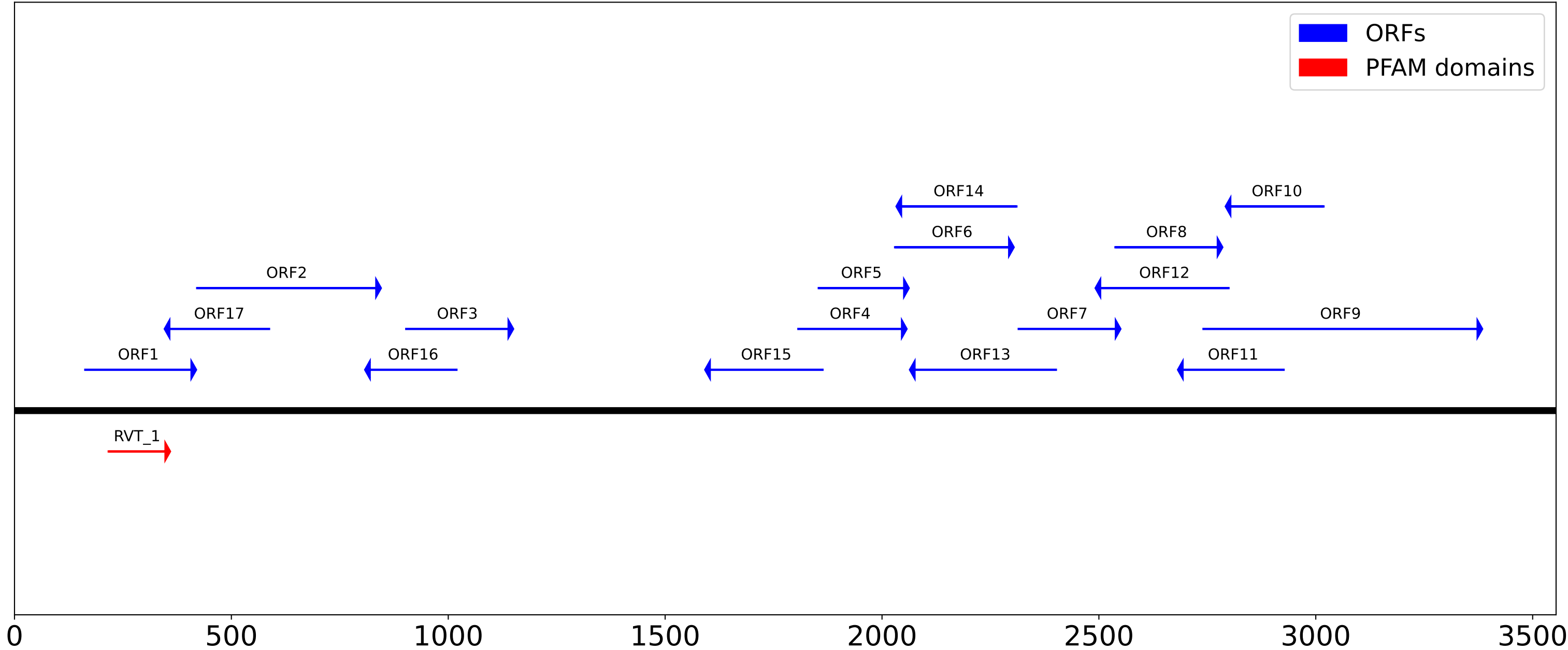


Before TETrimmer 3554 bp

After TEtrimmer ORF and PFAM domain plot



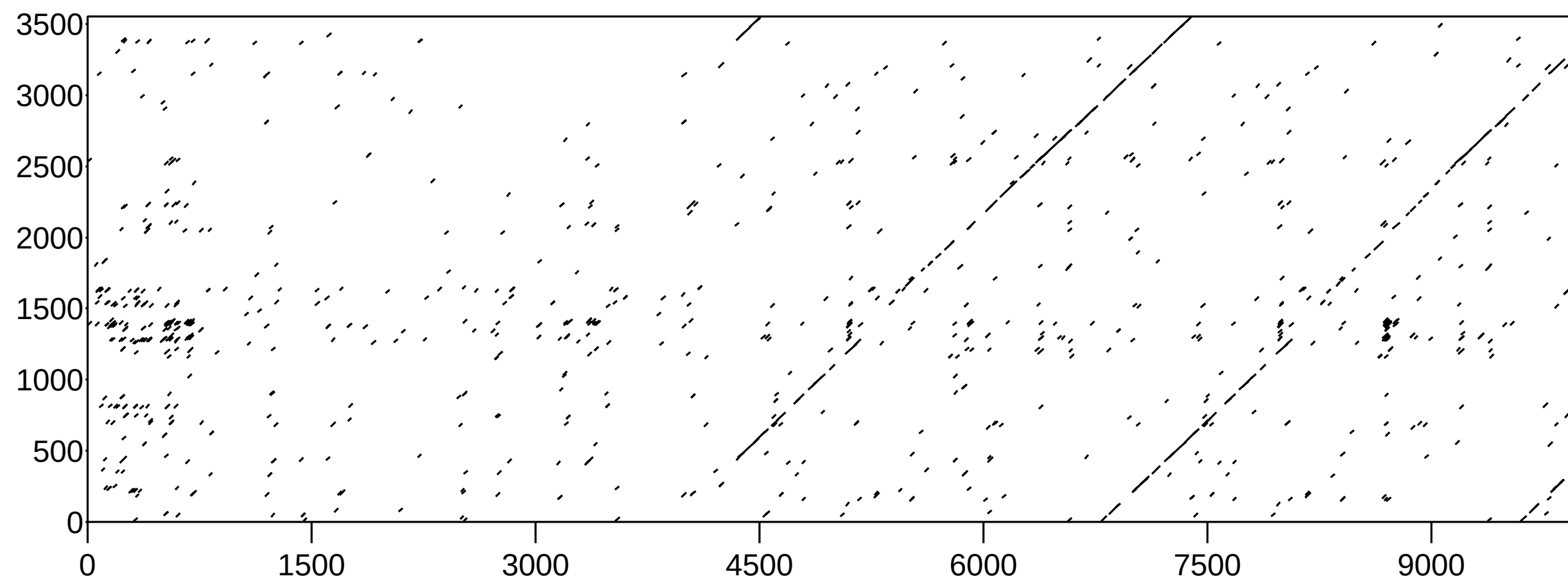
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)