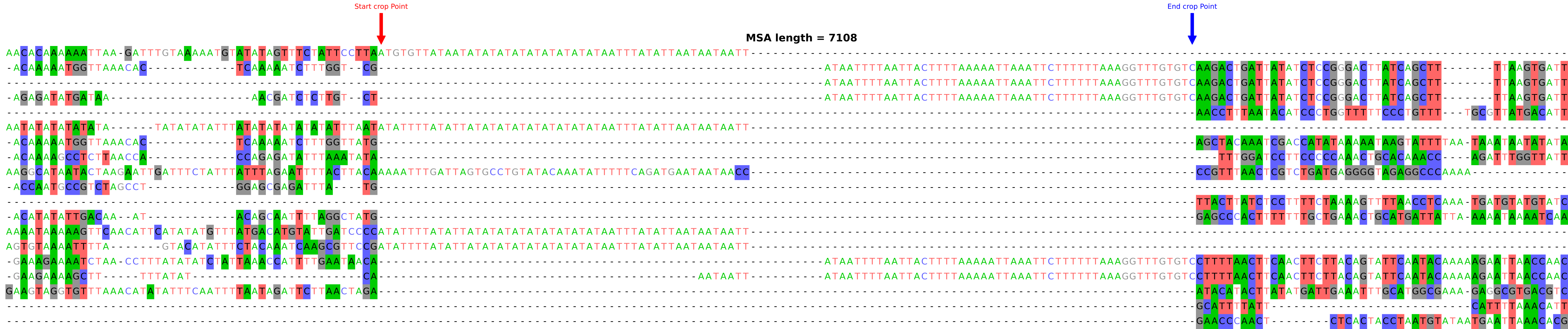
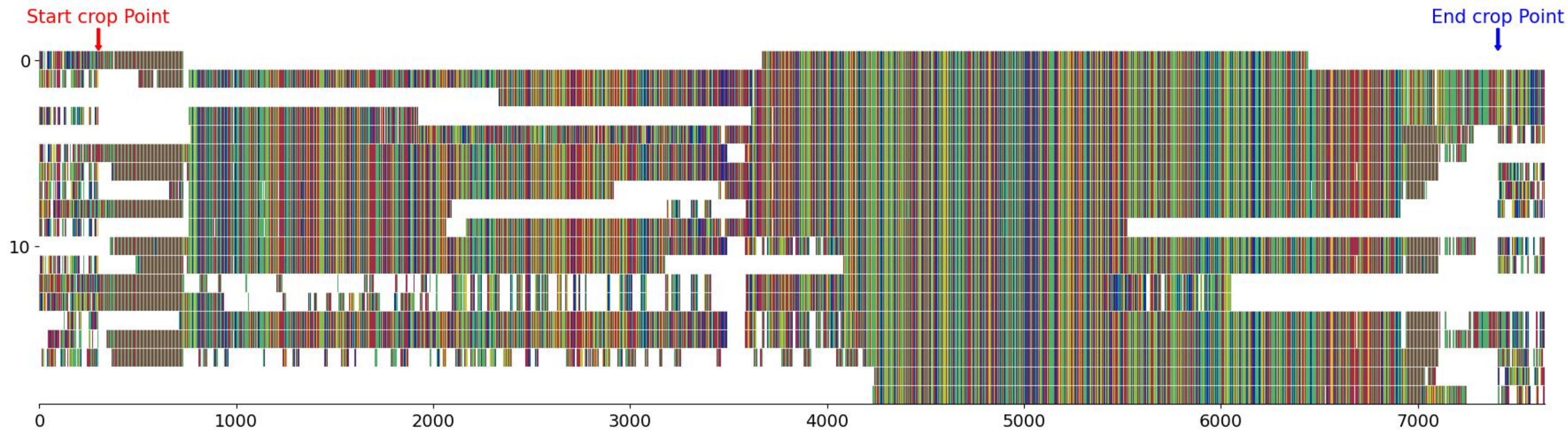




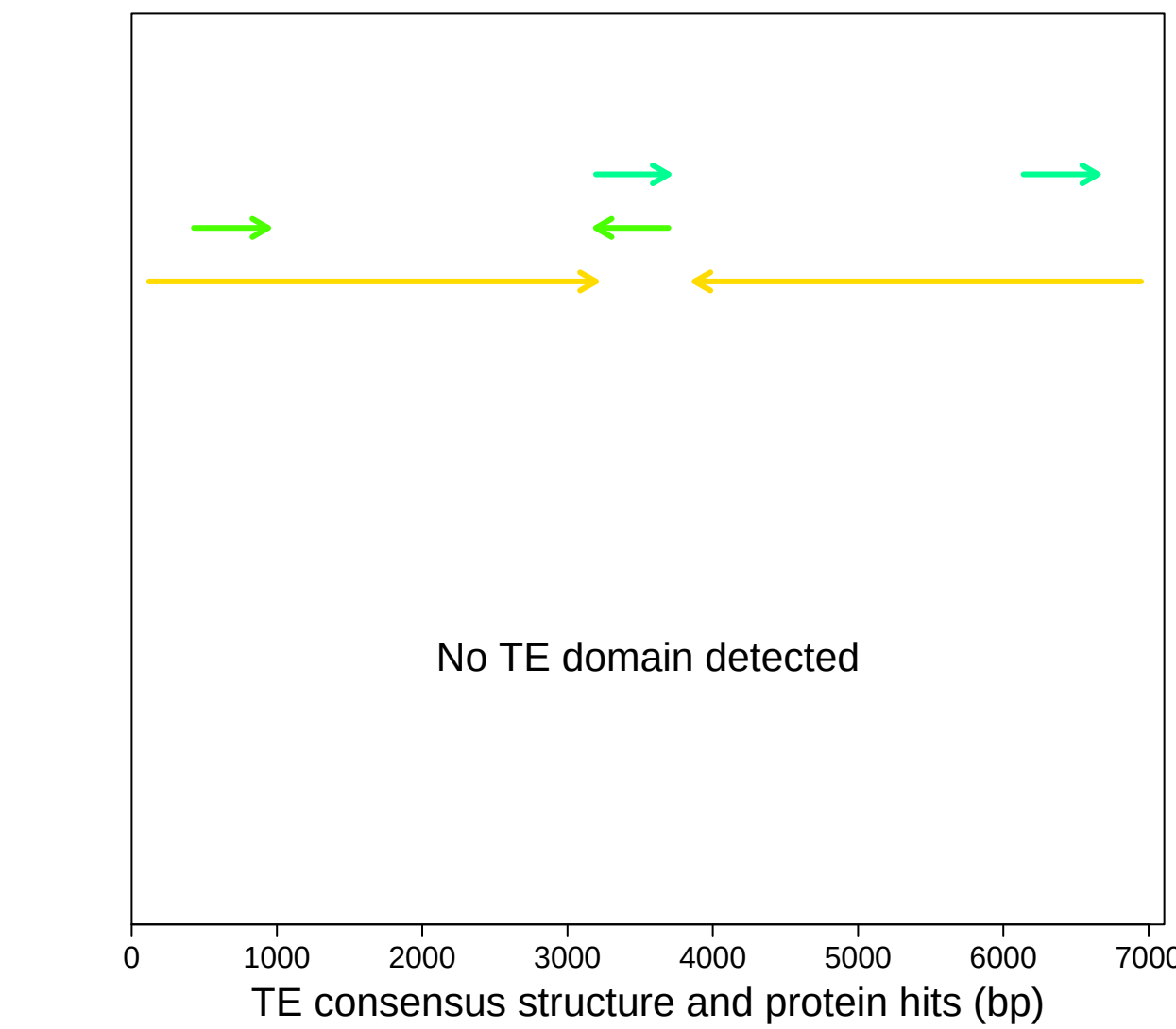
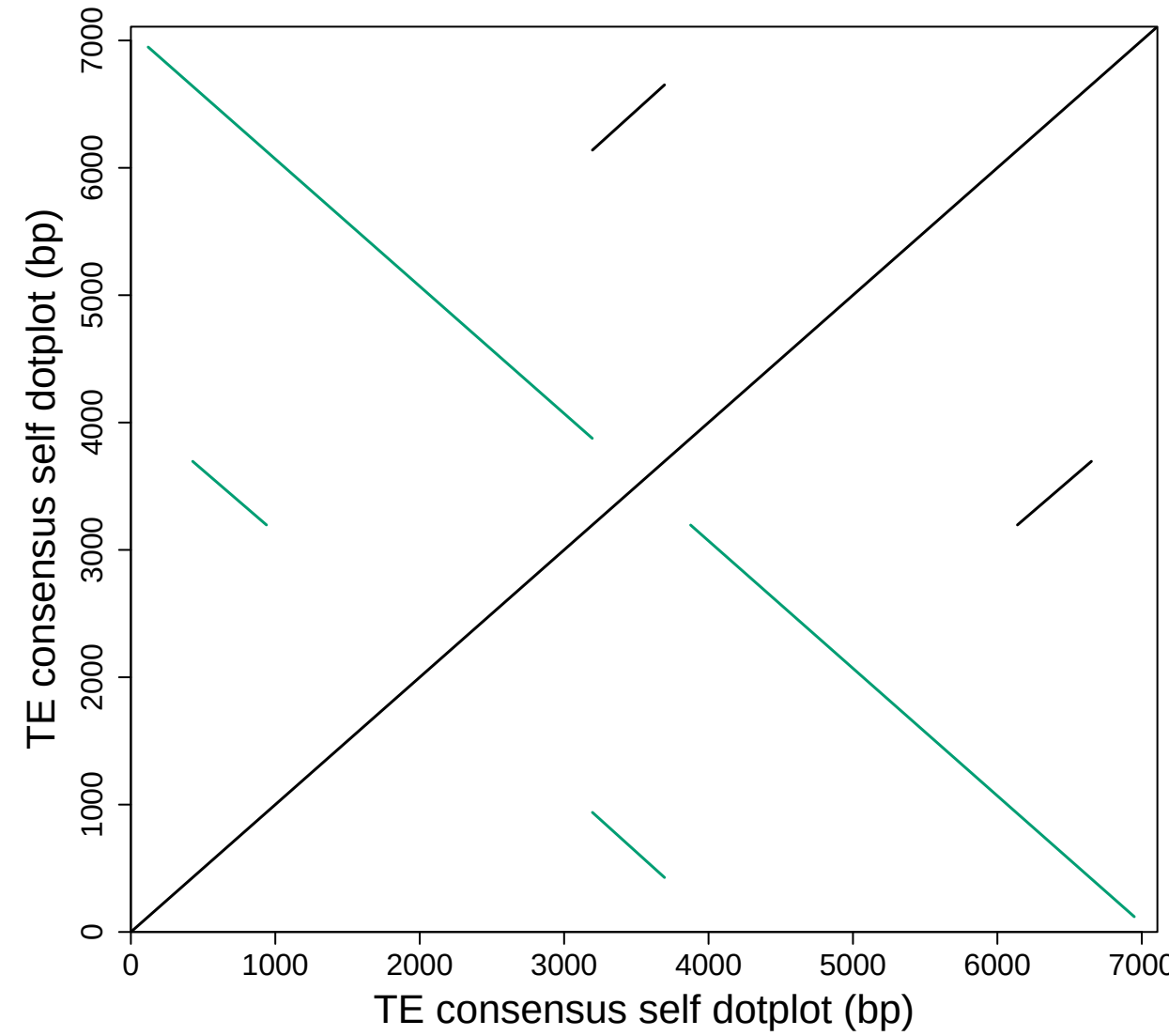
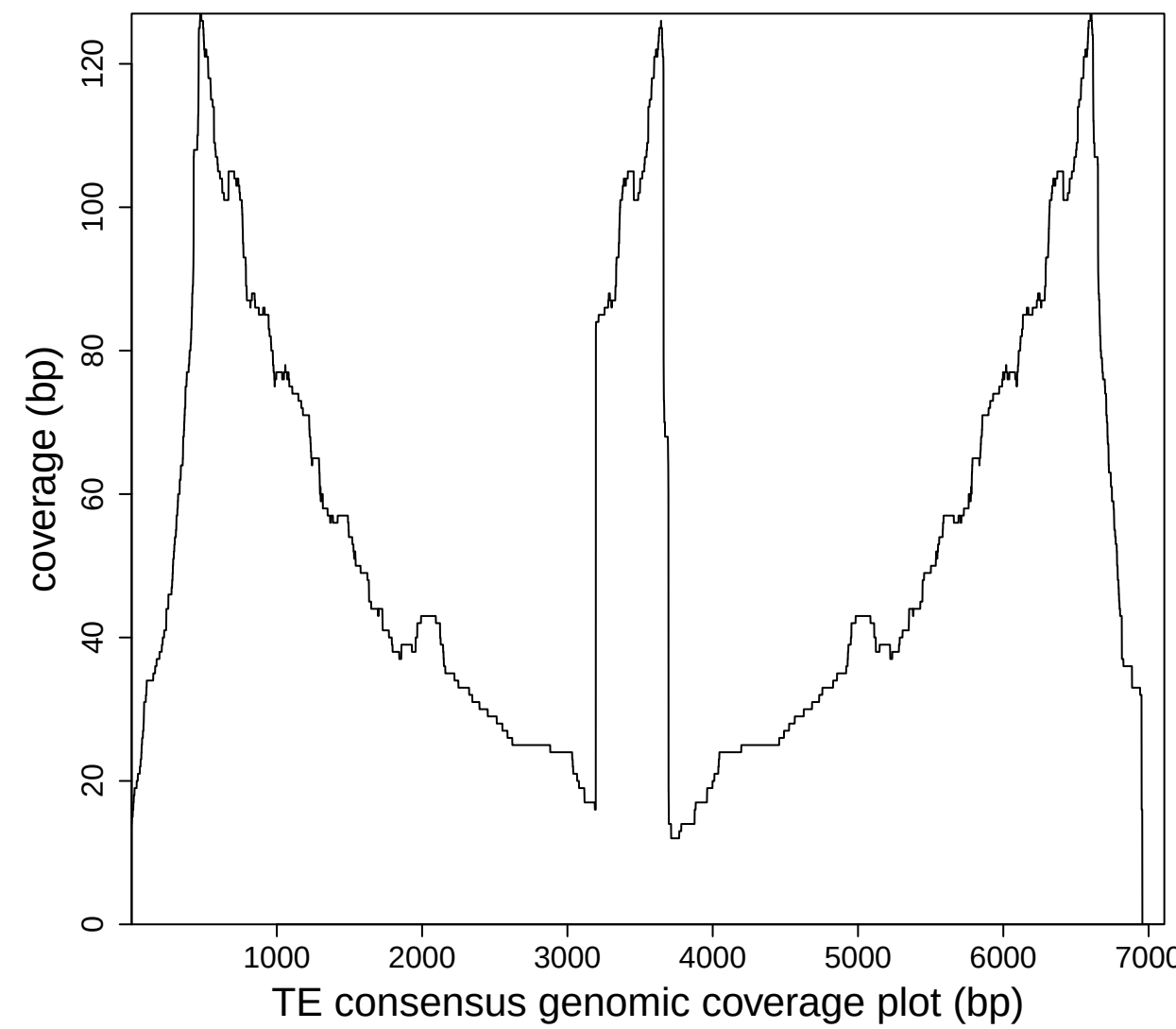
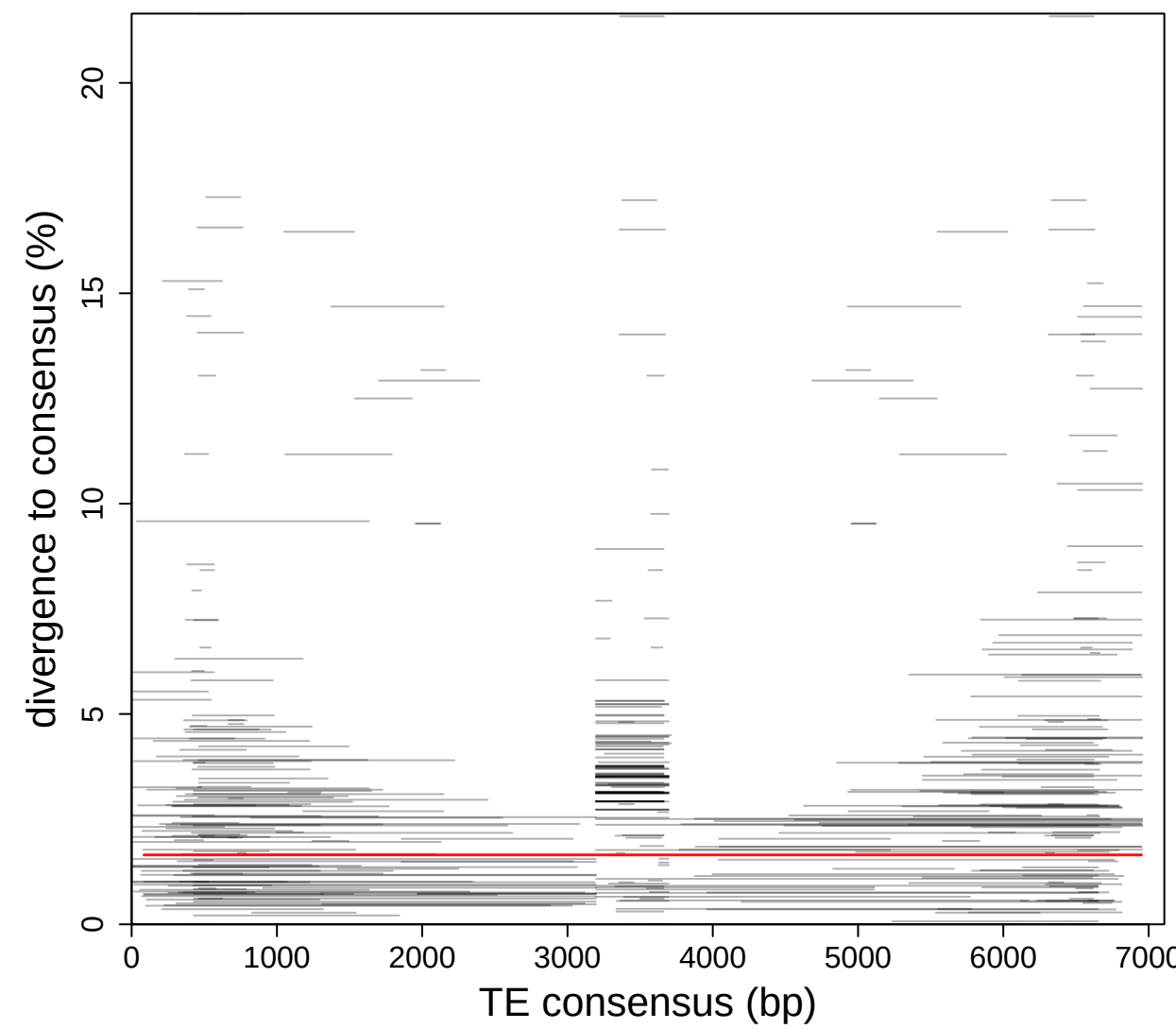
Start crop Point

End crop Point



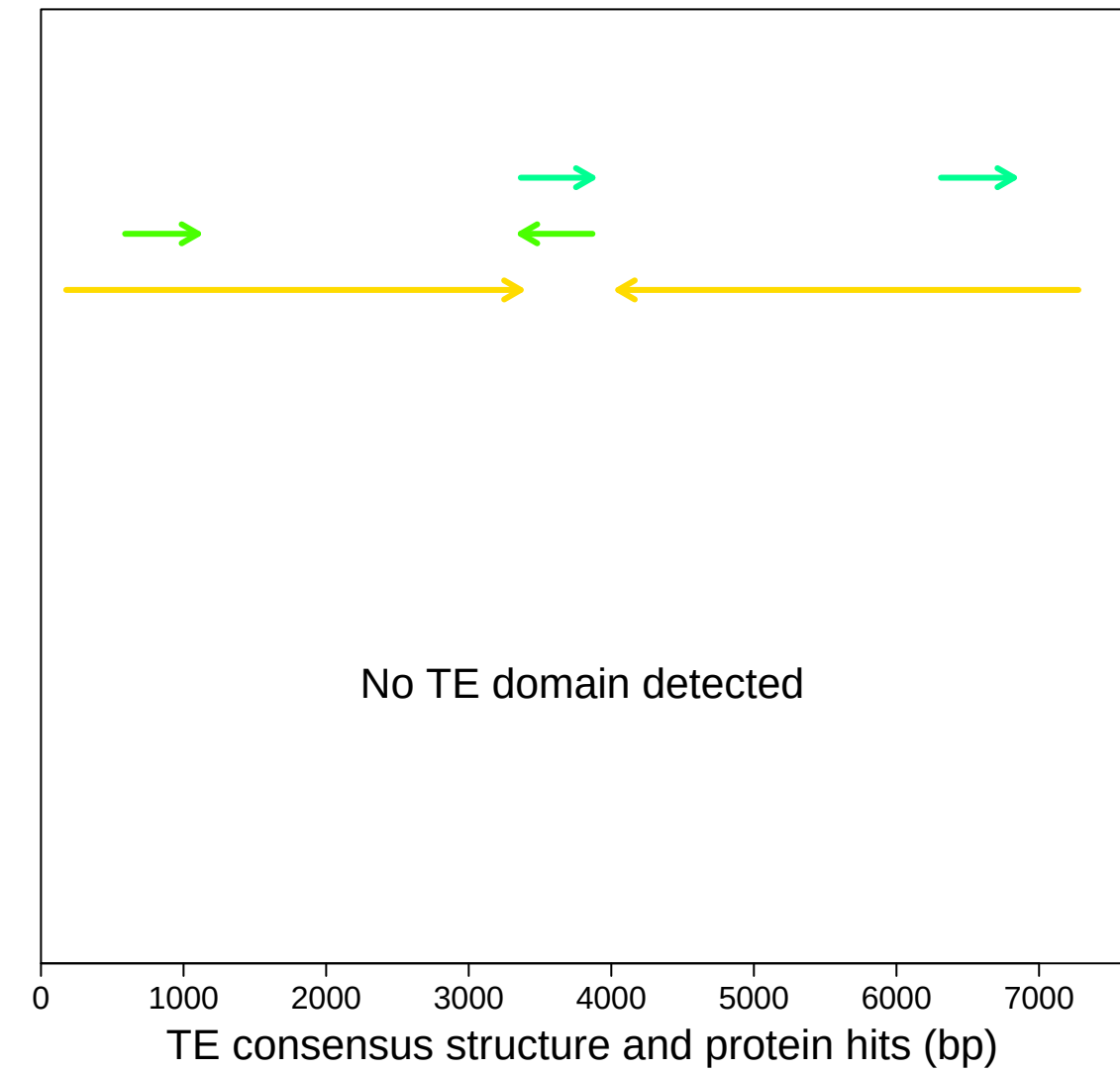
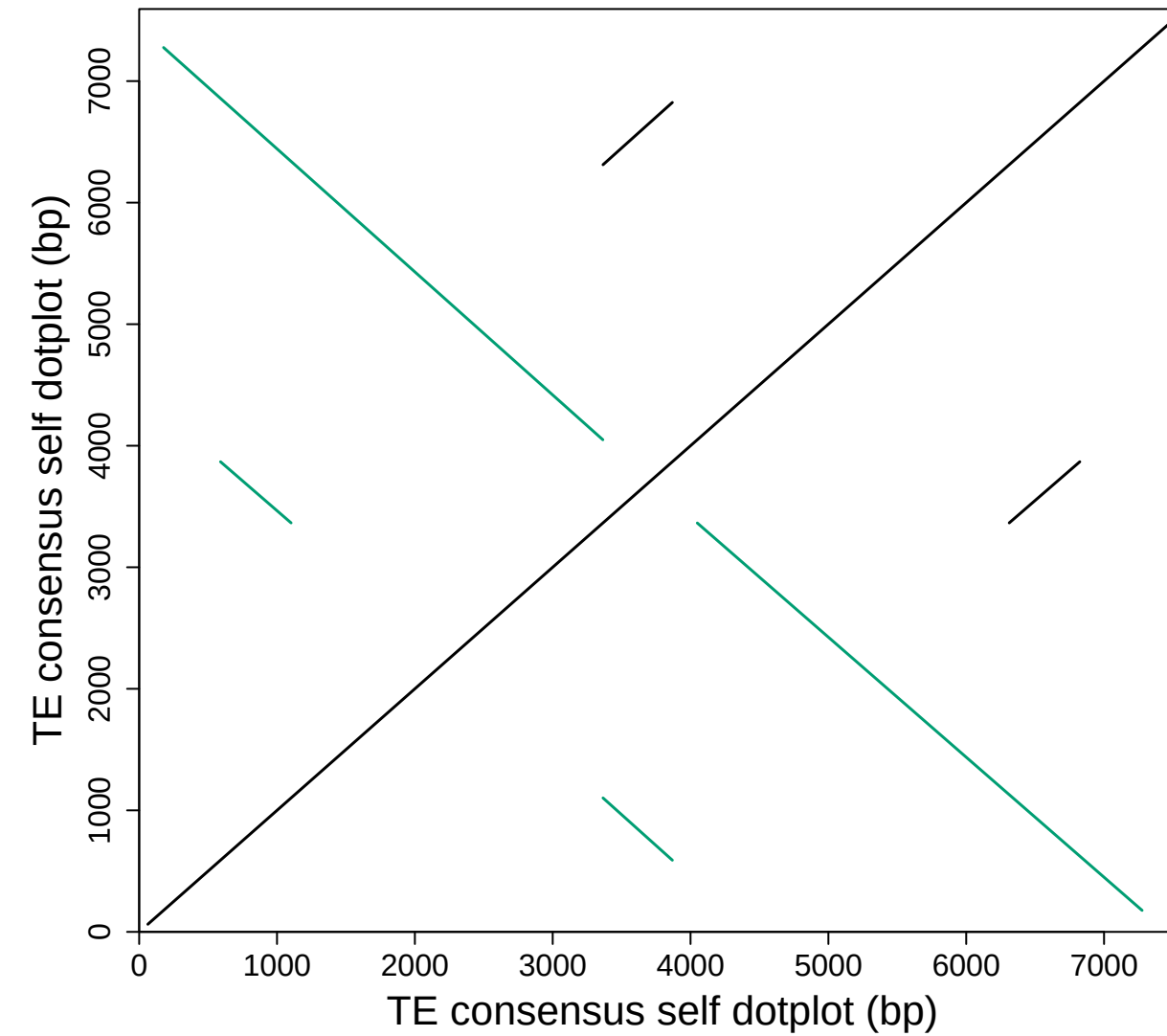
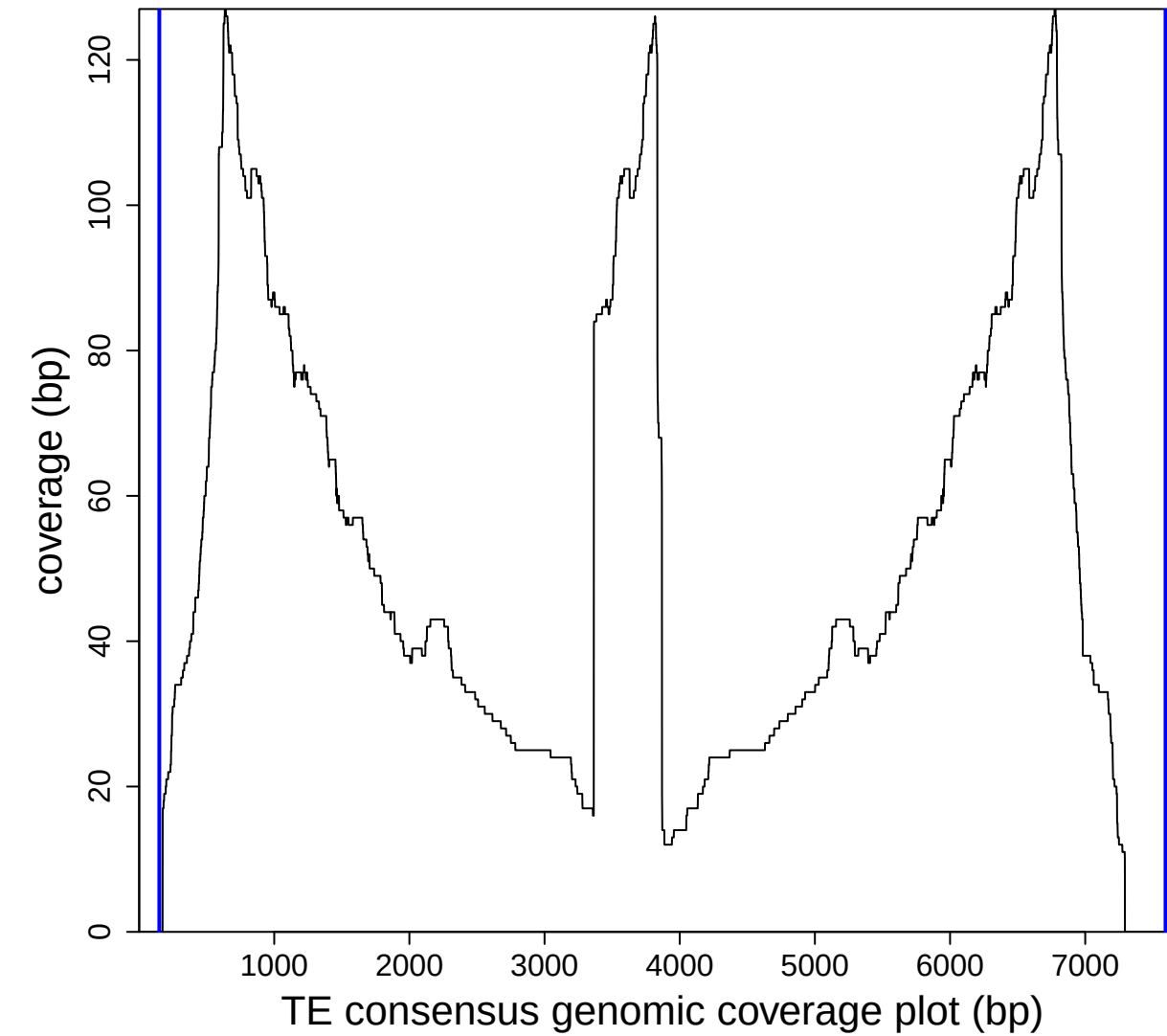
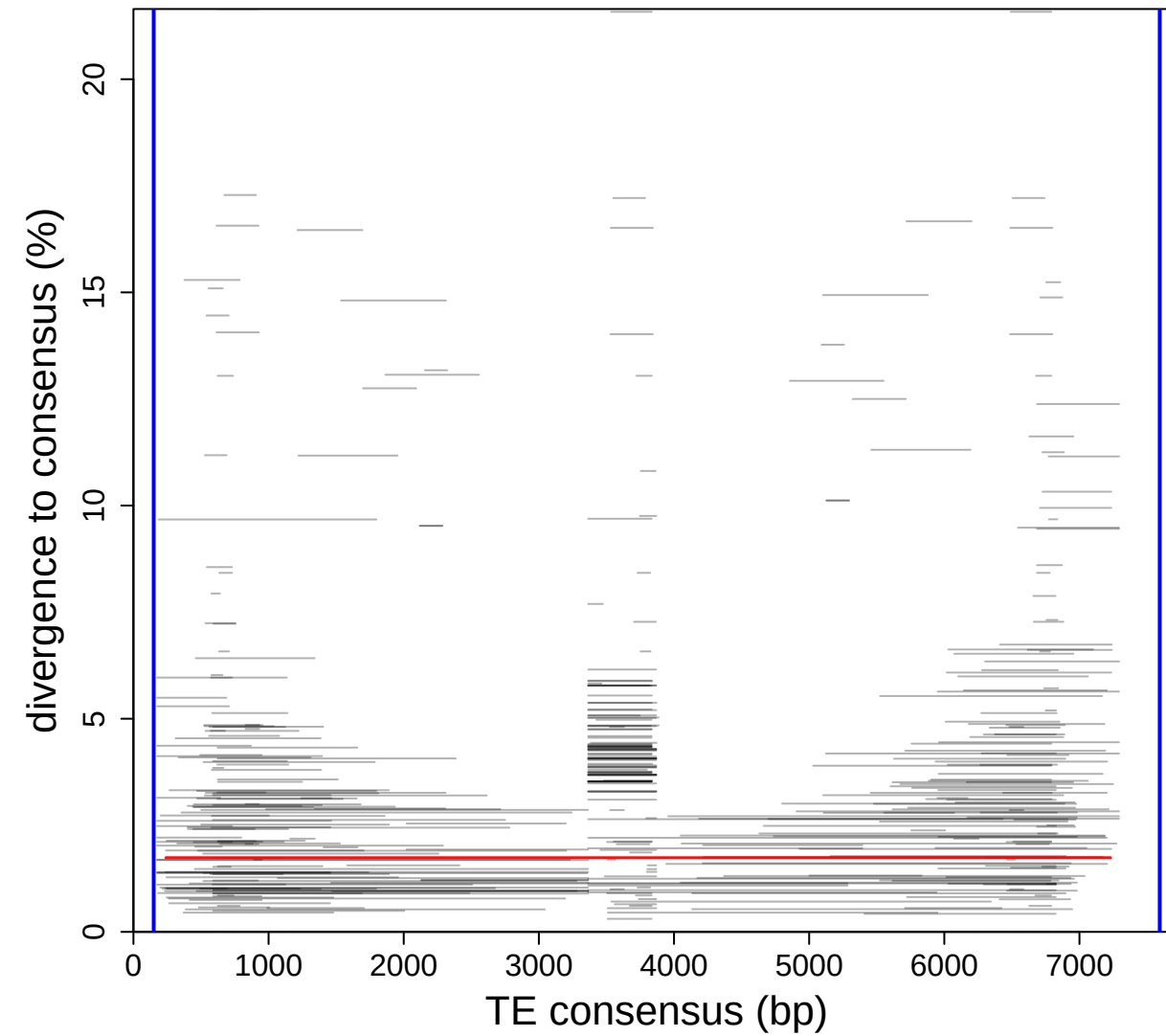
bed uf.bed g 2.bed fm 1.bed 0 0 bcln.fa aln.fa cl.fa gs.fa ce
size: 7108bp; fragments: 437; full length: 1 (>=6397.2bp)

After TEtrimmer 7108 bp



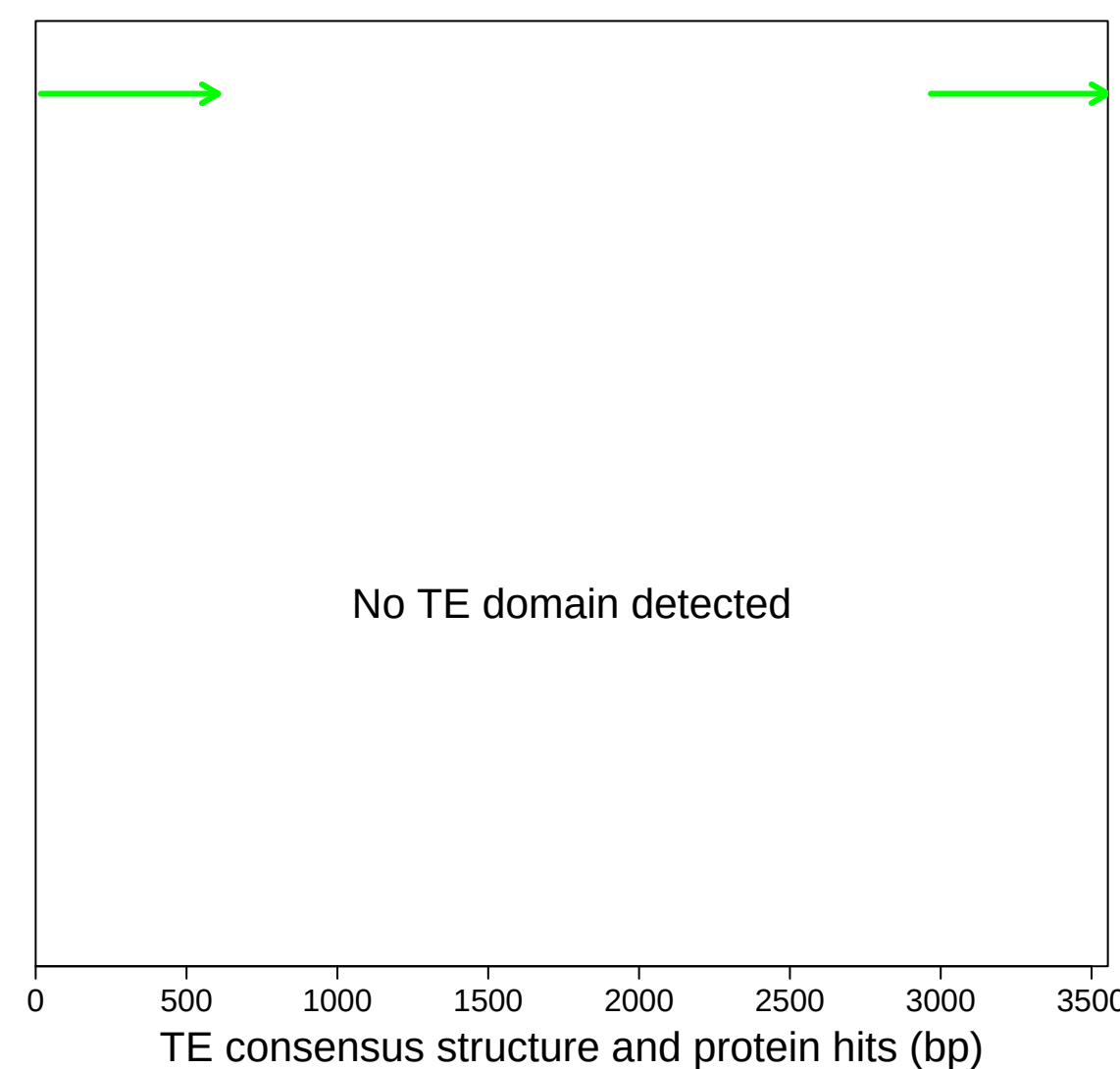
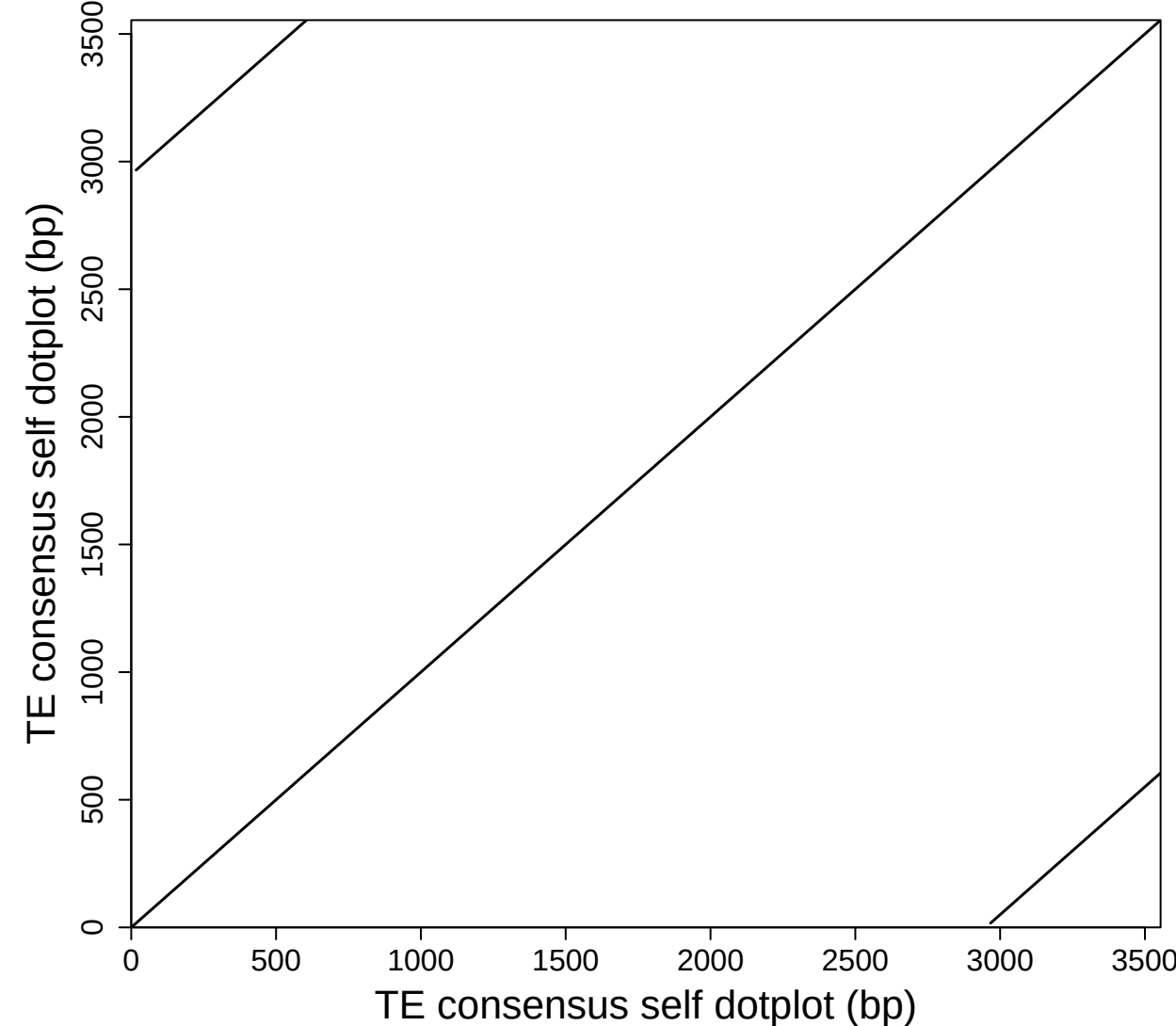
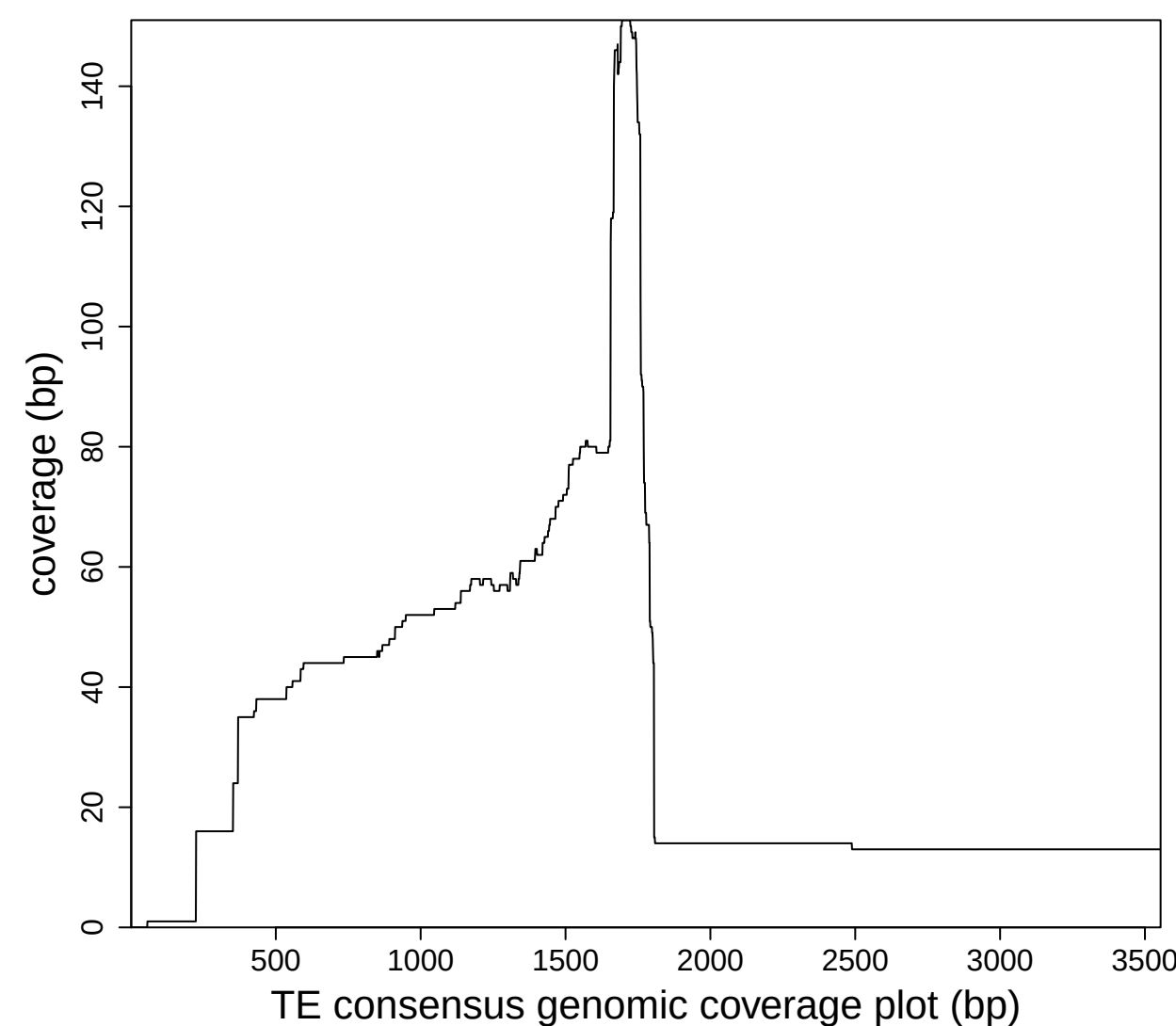
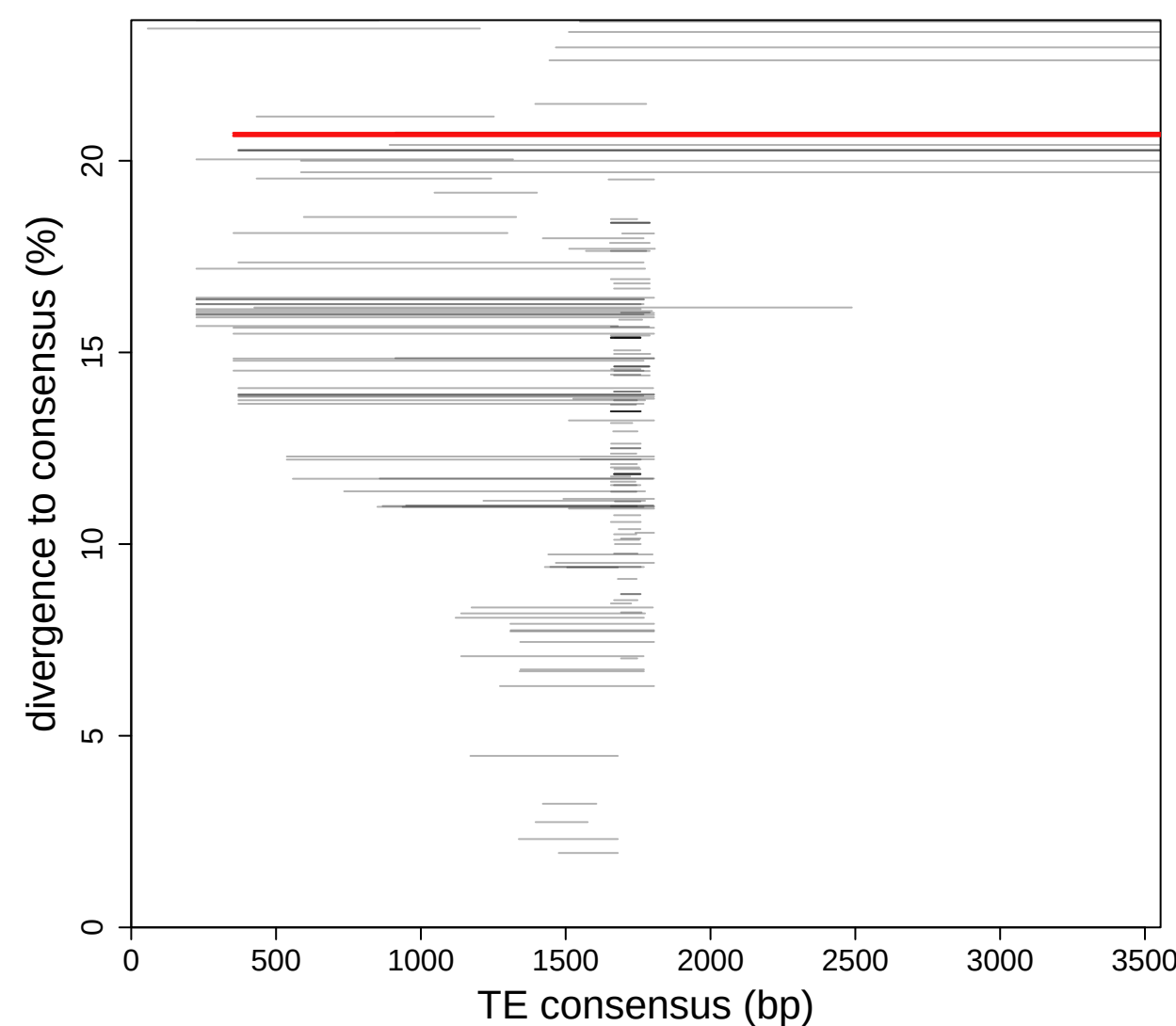
6.fasta.b.bed uf.bed g 2.bed fm 1.bed 0 0 bcln.fa aln.fa cl.fa
size: 7744bp; fragments: 437; full length: 1 (>=6969.6bp)

After TEtrimmer Extended plot Blue lines are boundaries

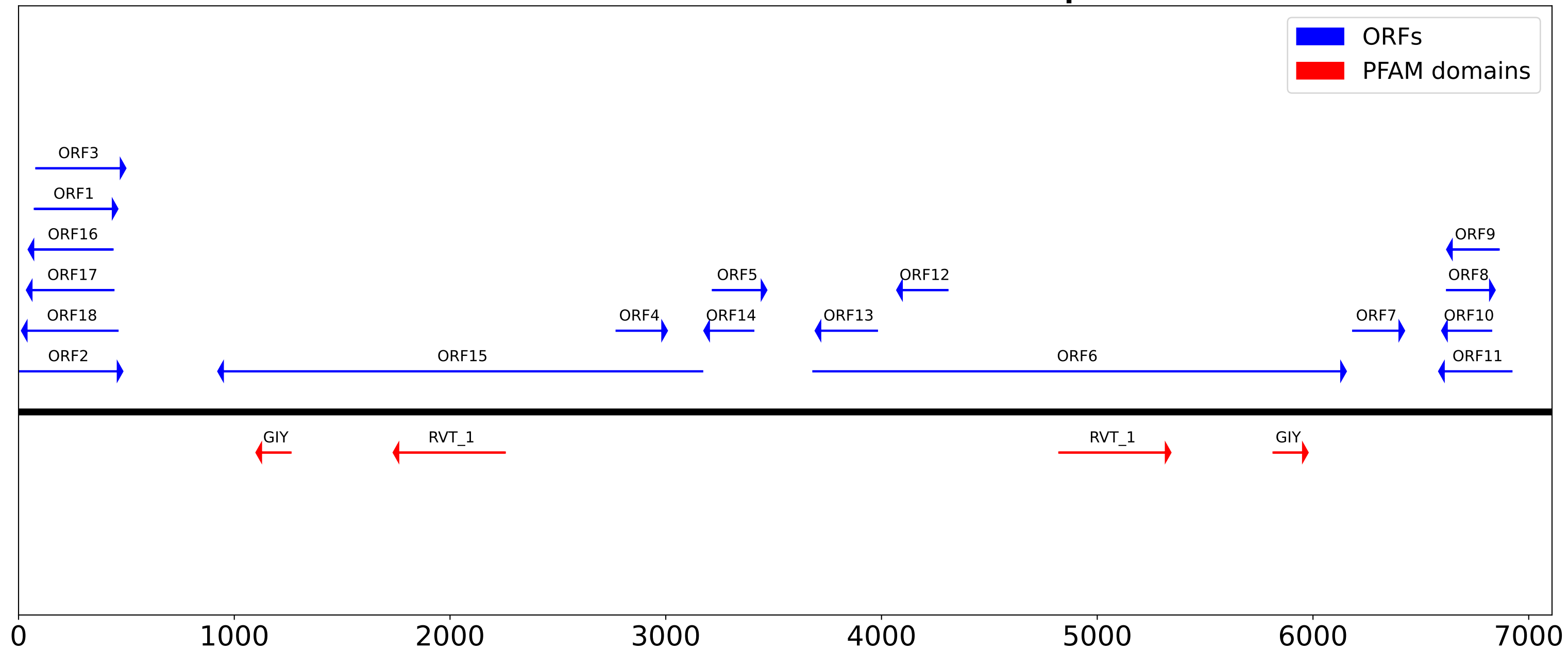


TE: rnd_5_family_946
size: 3554bp; fragments: 167; full length: 2 (≥ 3198.6 bp)

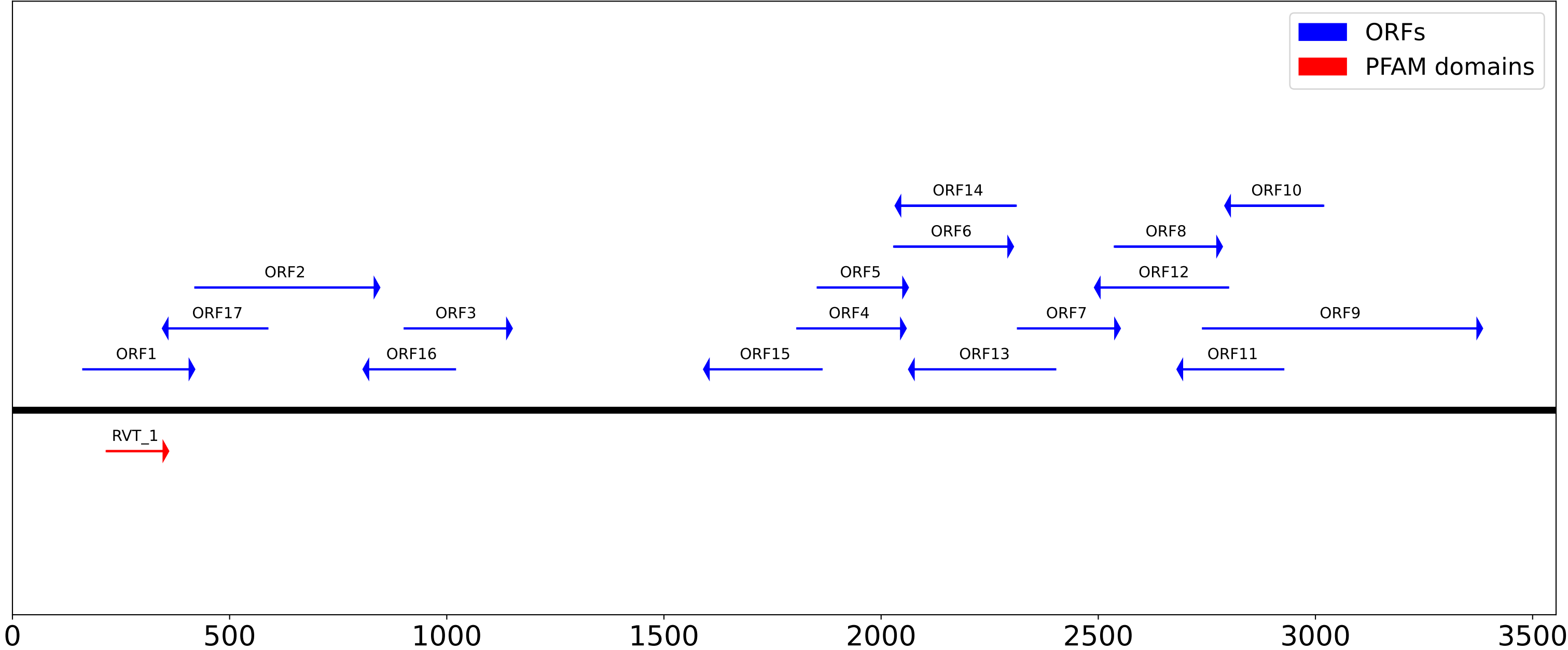
Before TEtrimmer 3554 bp



After TEtrimmer ORF and PFAM domain plot



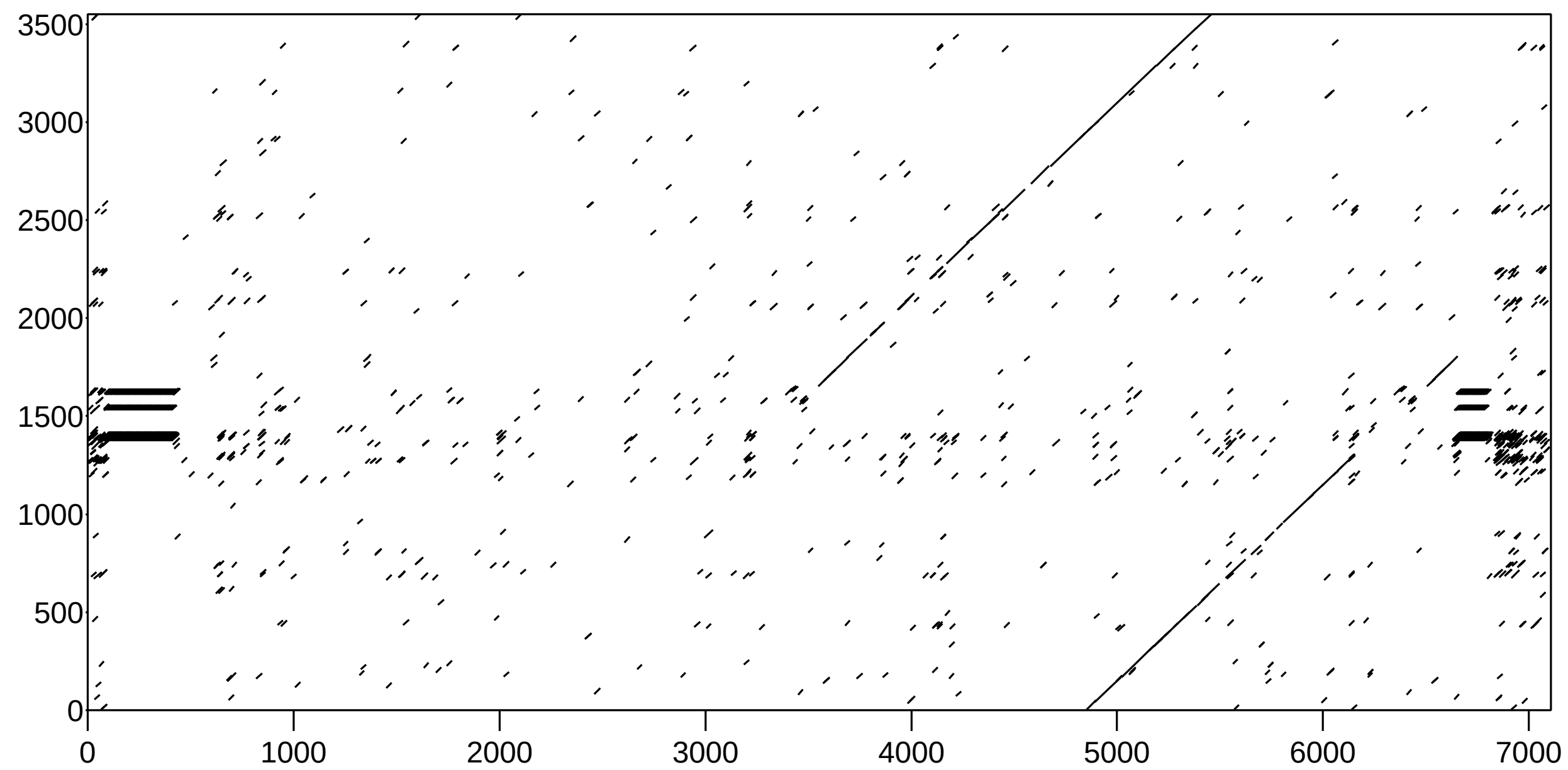
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)