

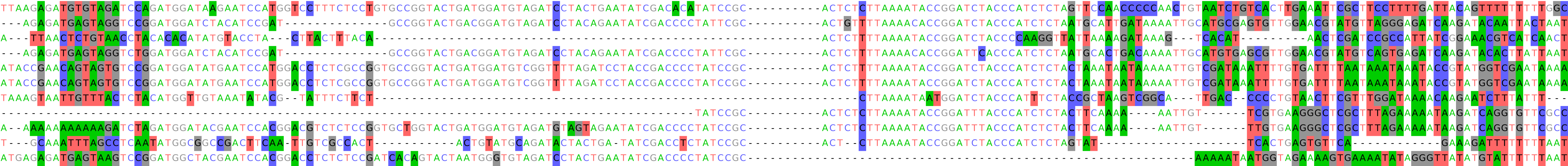
Start crop Point



End crop Point

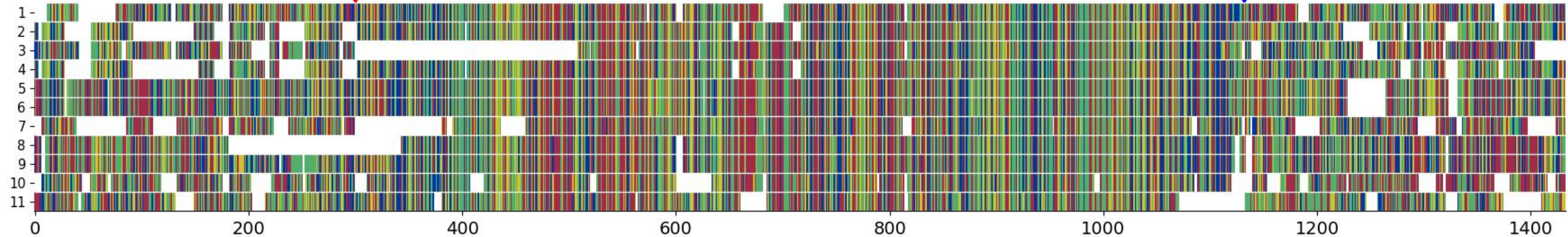


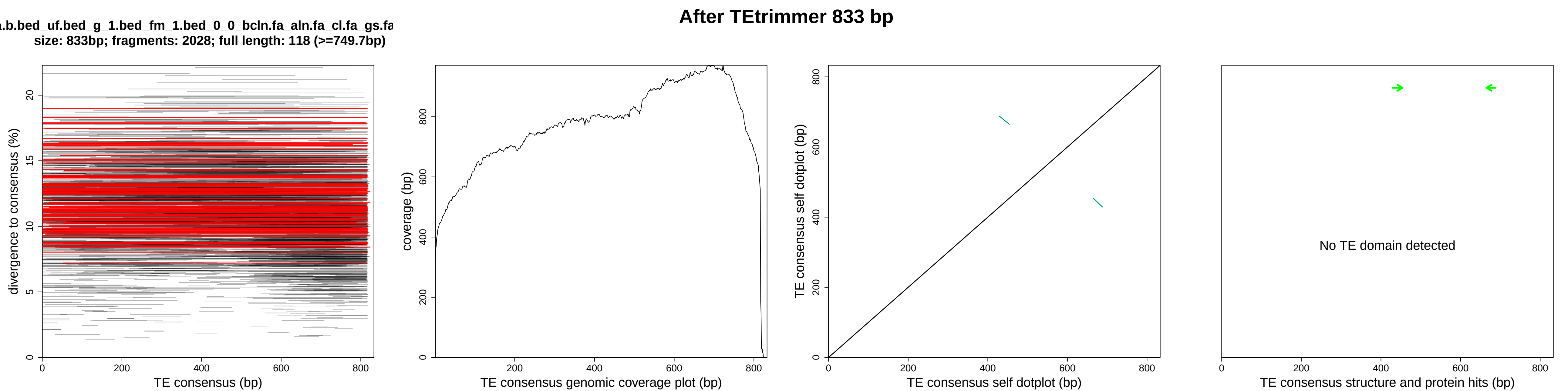
MSA length = 833



Start crop Point

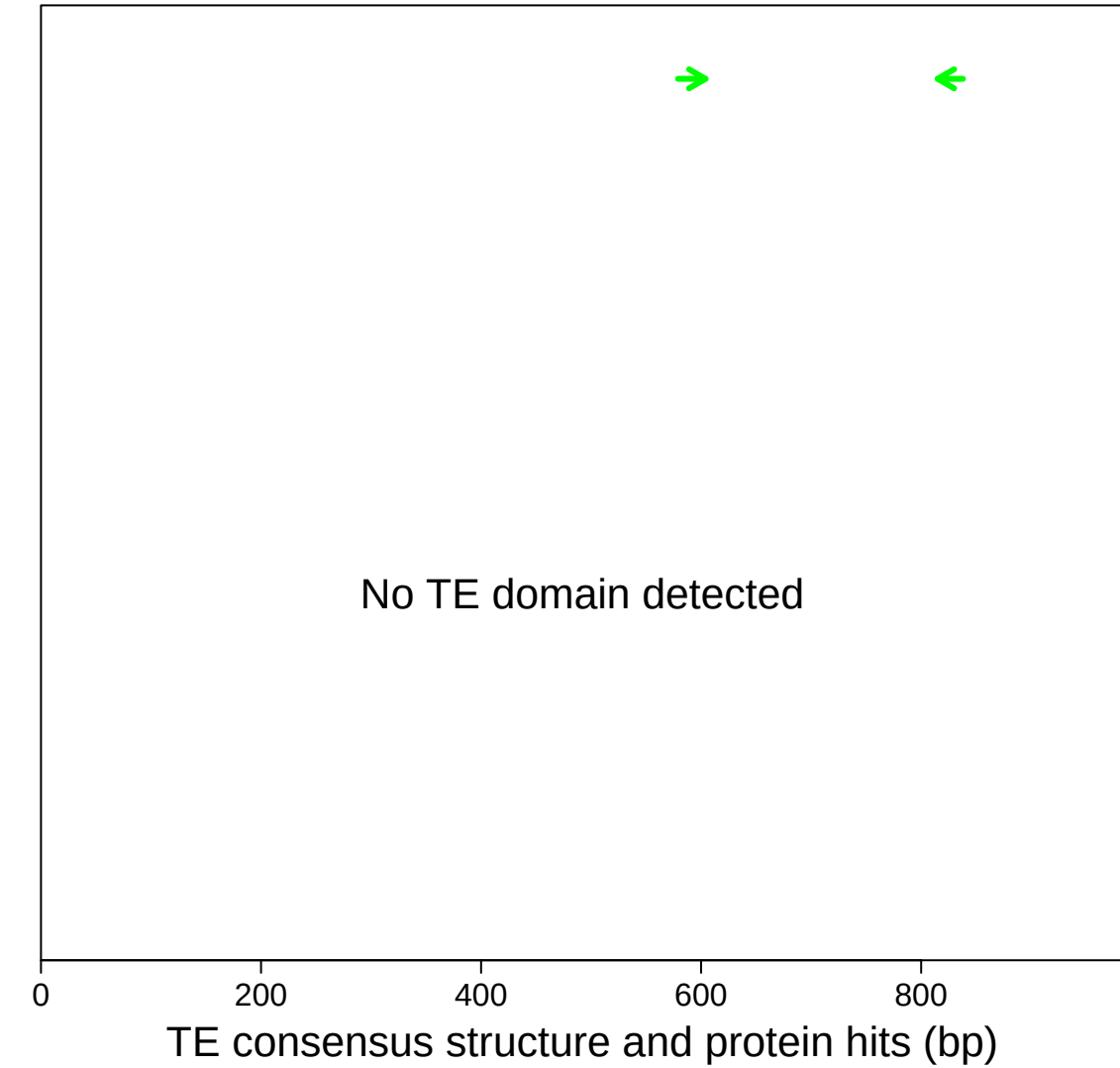
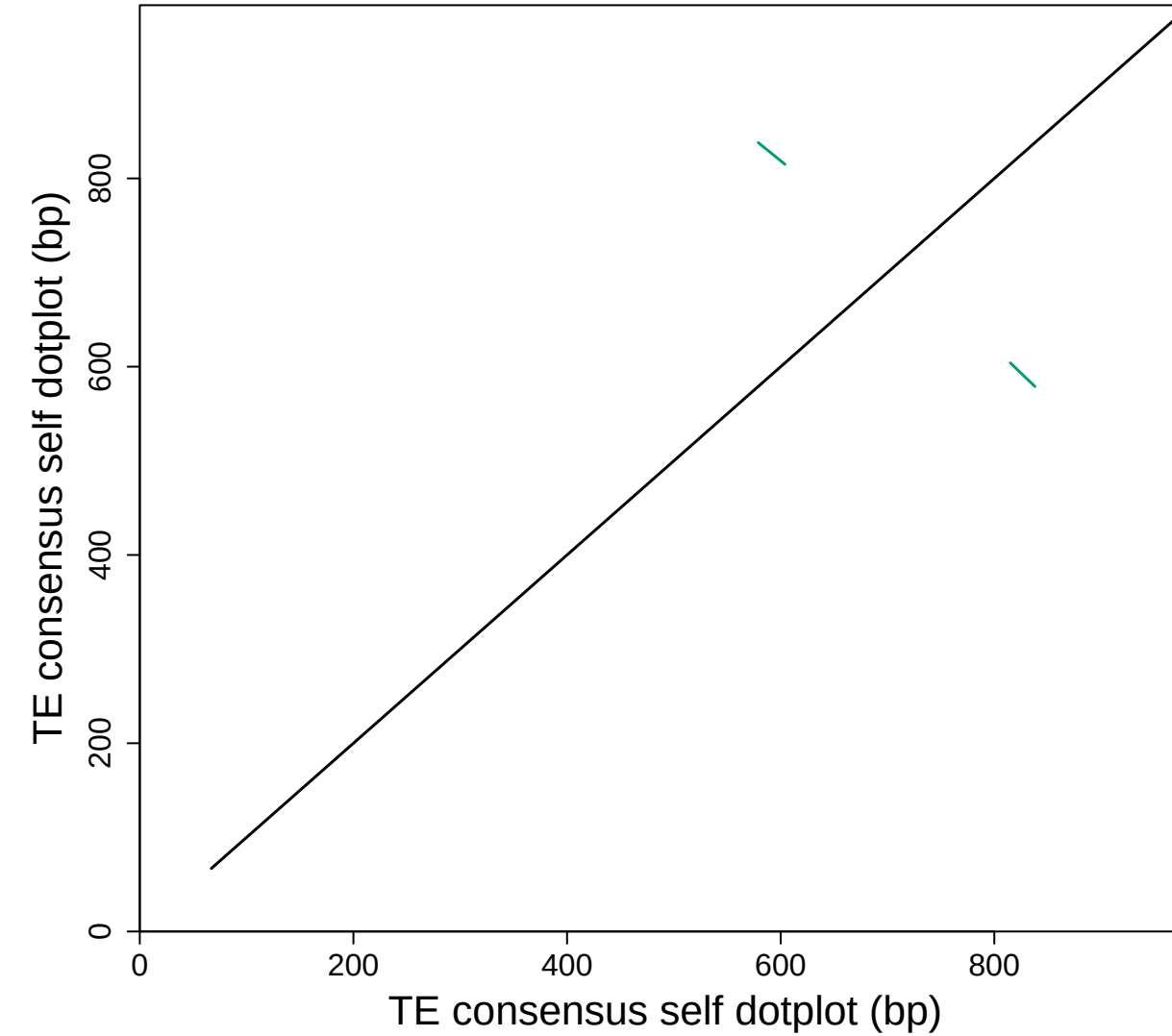
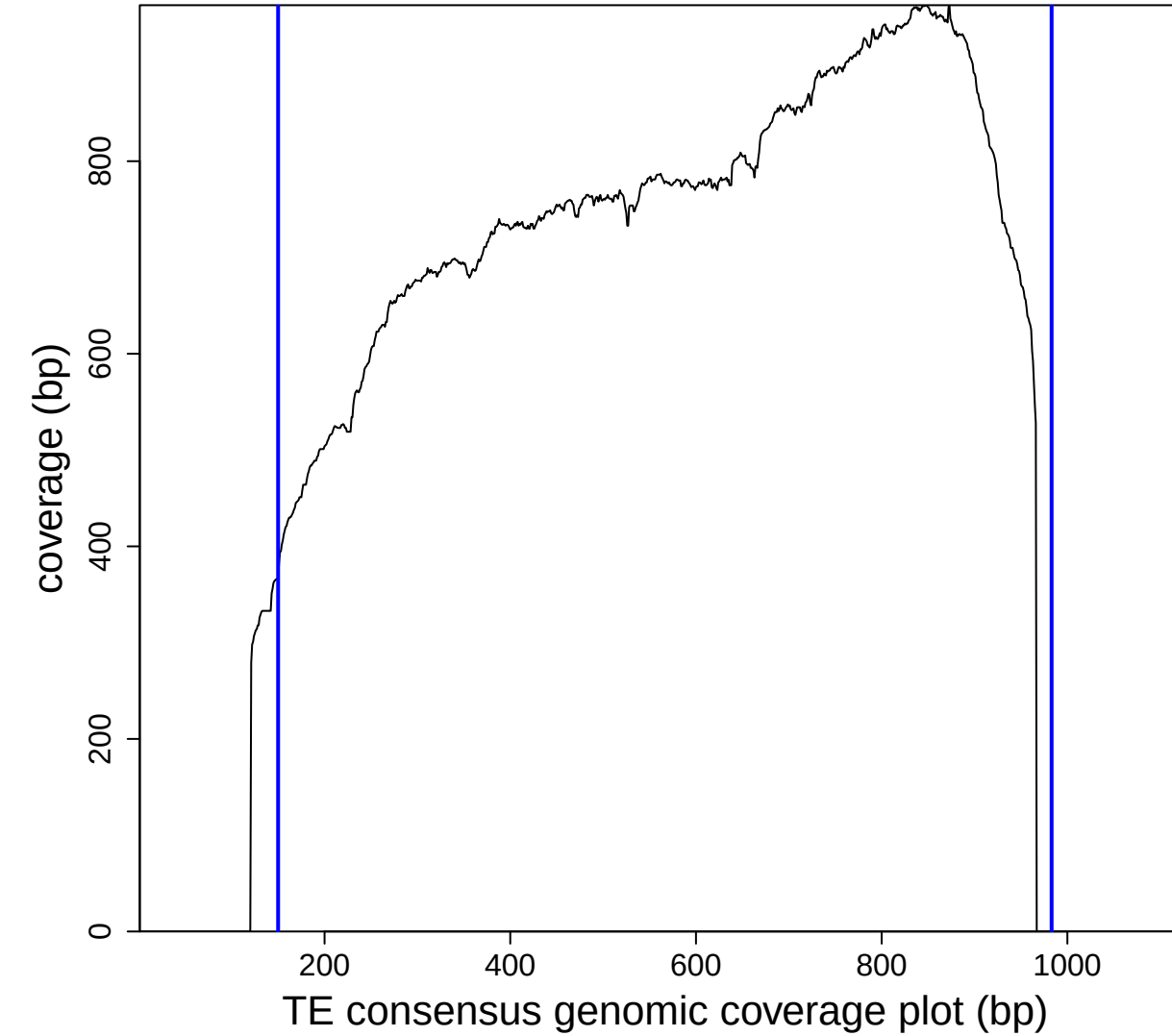
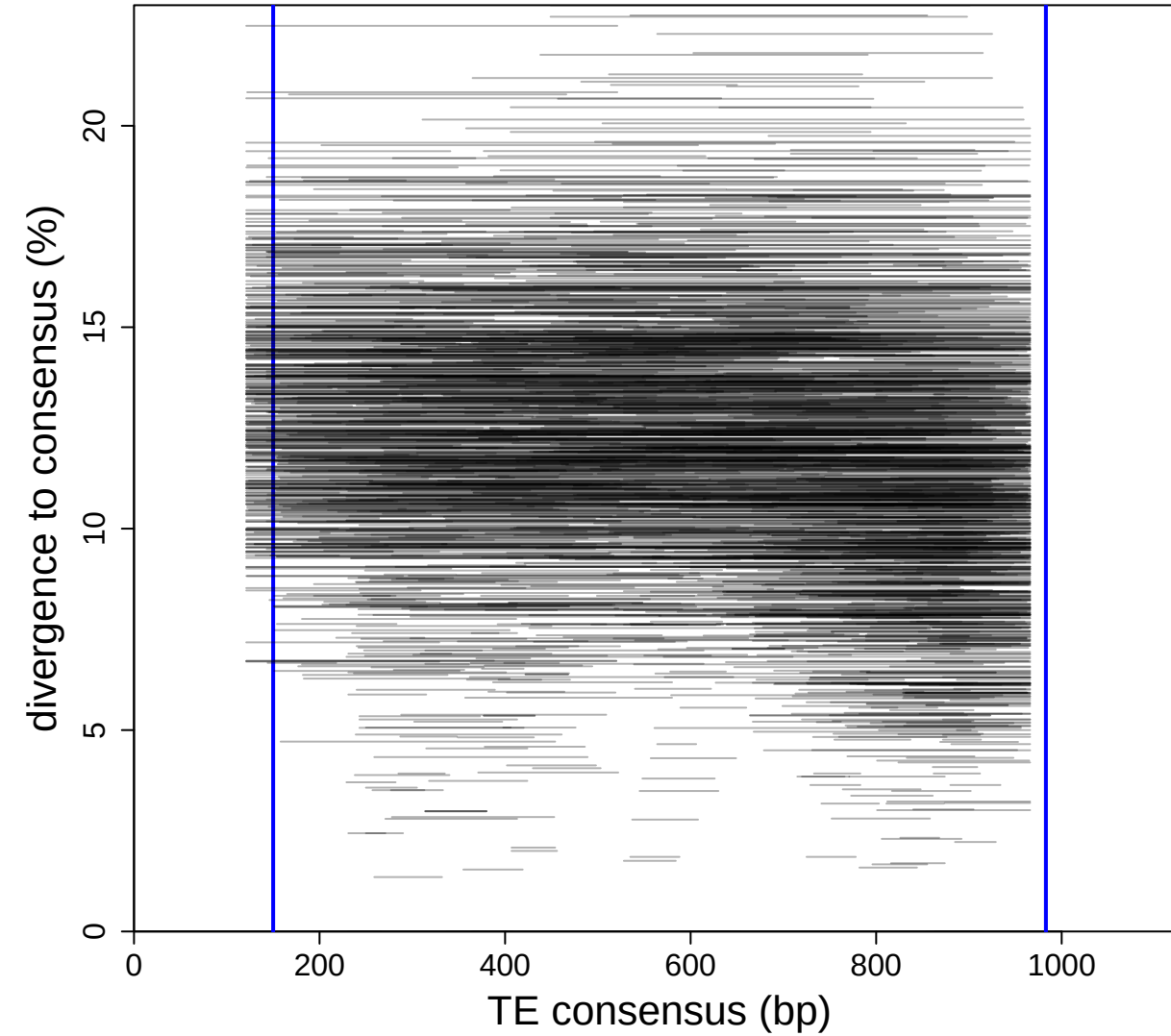
End crop Point



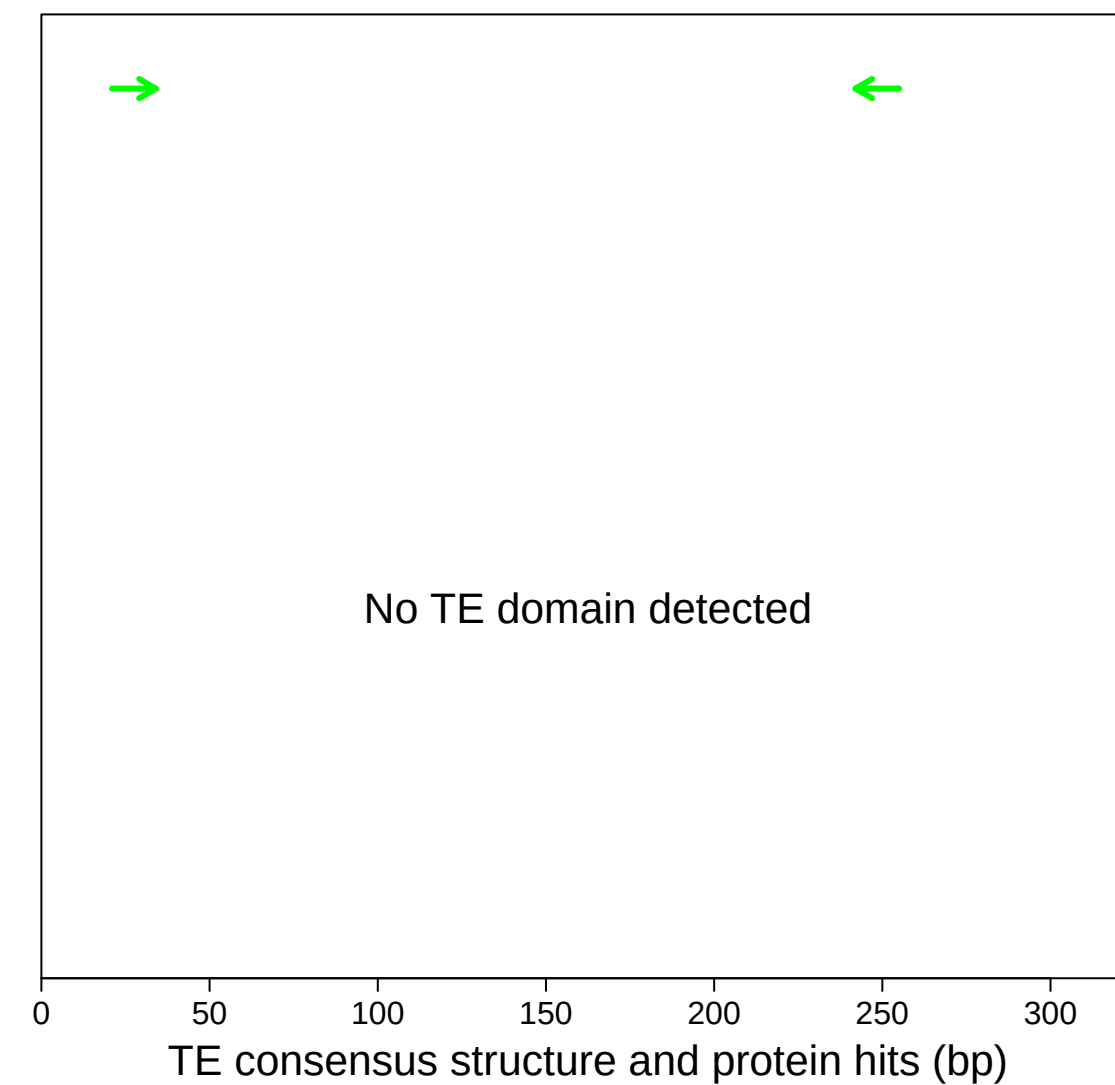
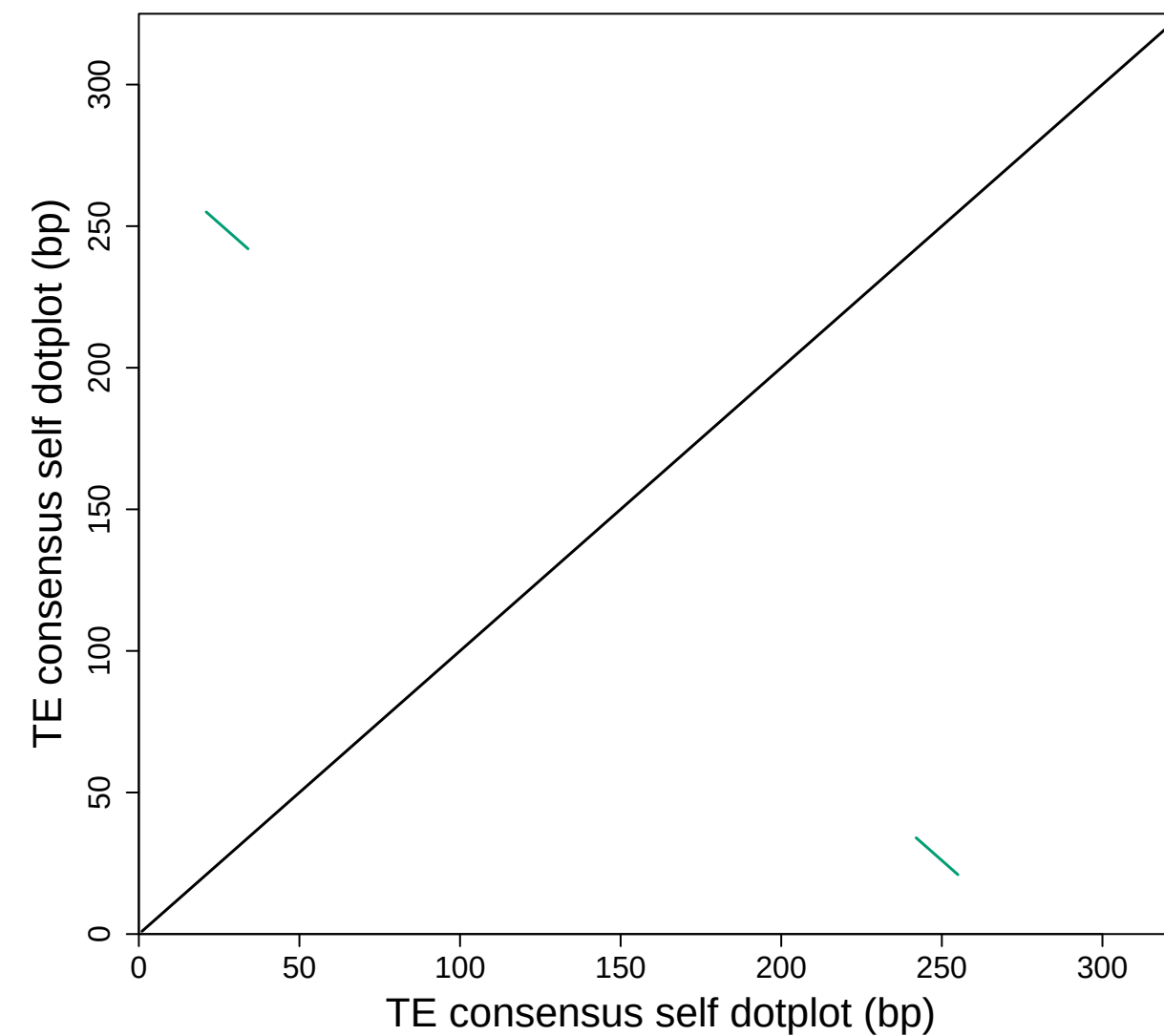
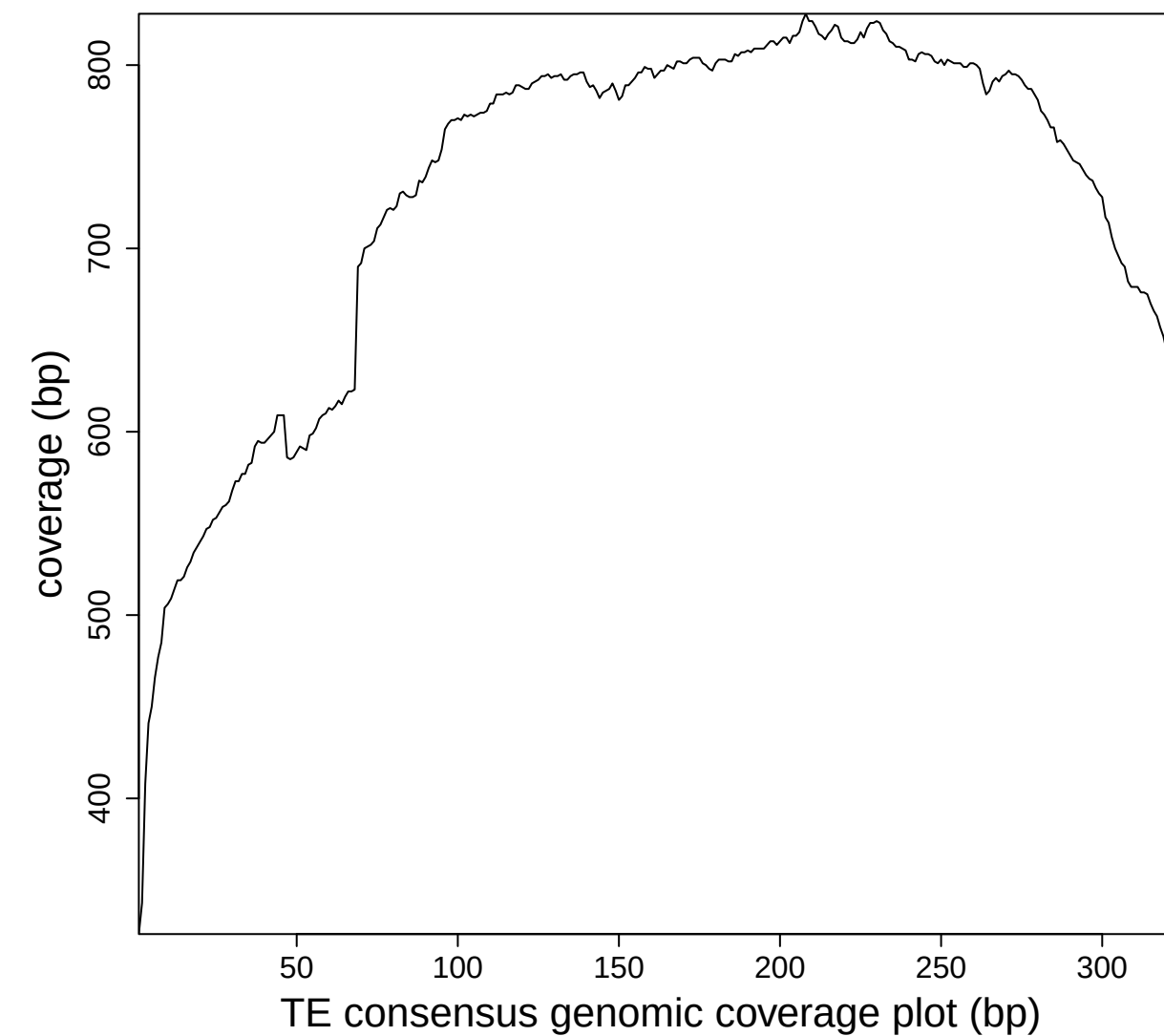
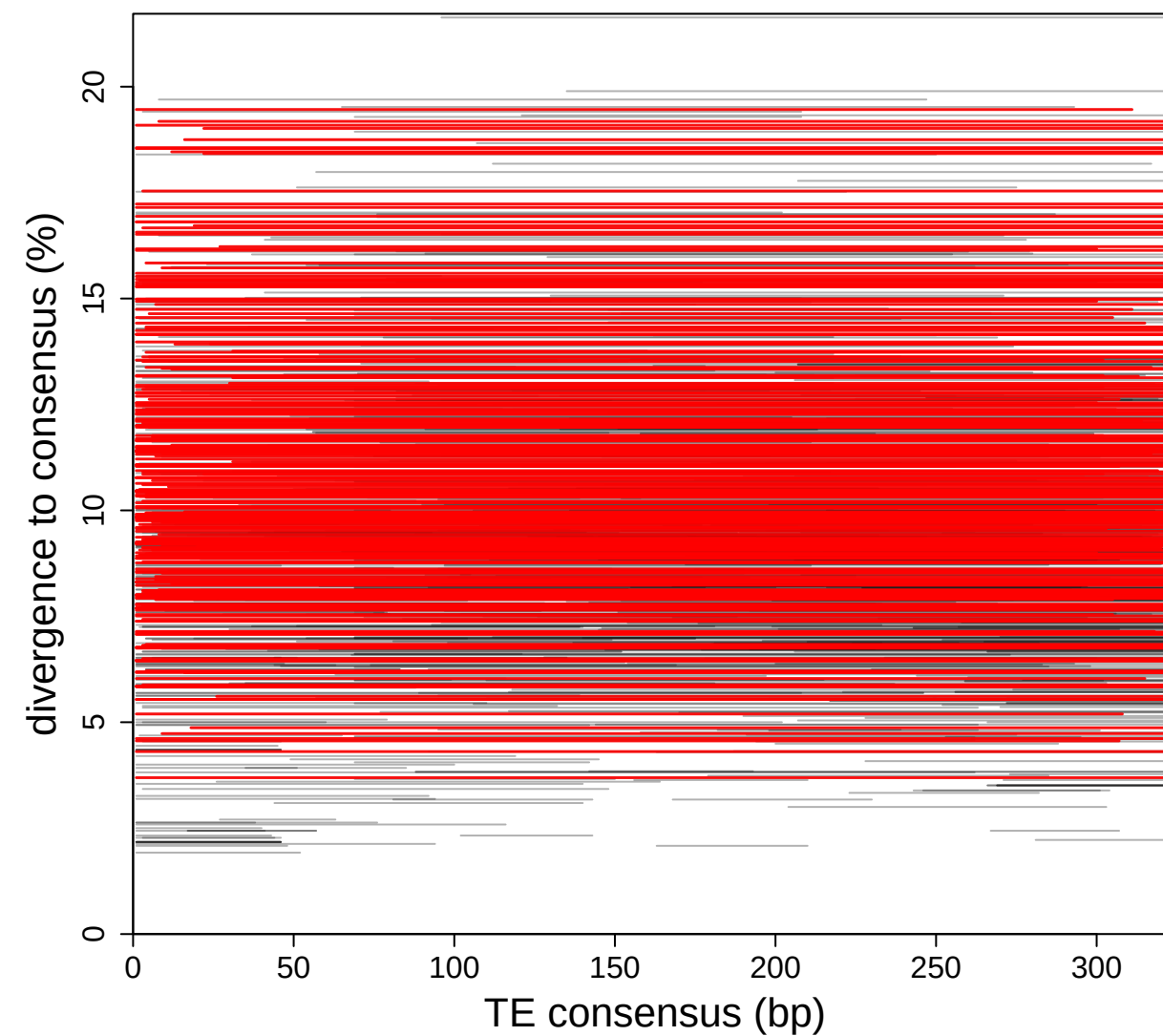


4.fasta.b.bed uf.bed g 1.bed fm 1.bed 0 0 bcln.fa aln.fa cl.fa
size: 1133bp; fragments: 1931; full length: 0 (≥ 1019.7 bp)

After TEtrimmer Extended plot Blue lines are boundaries

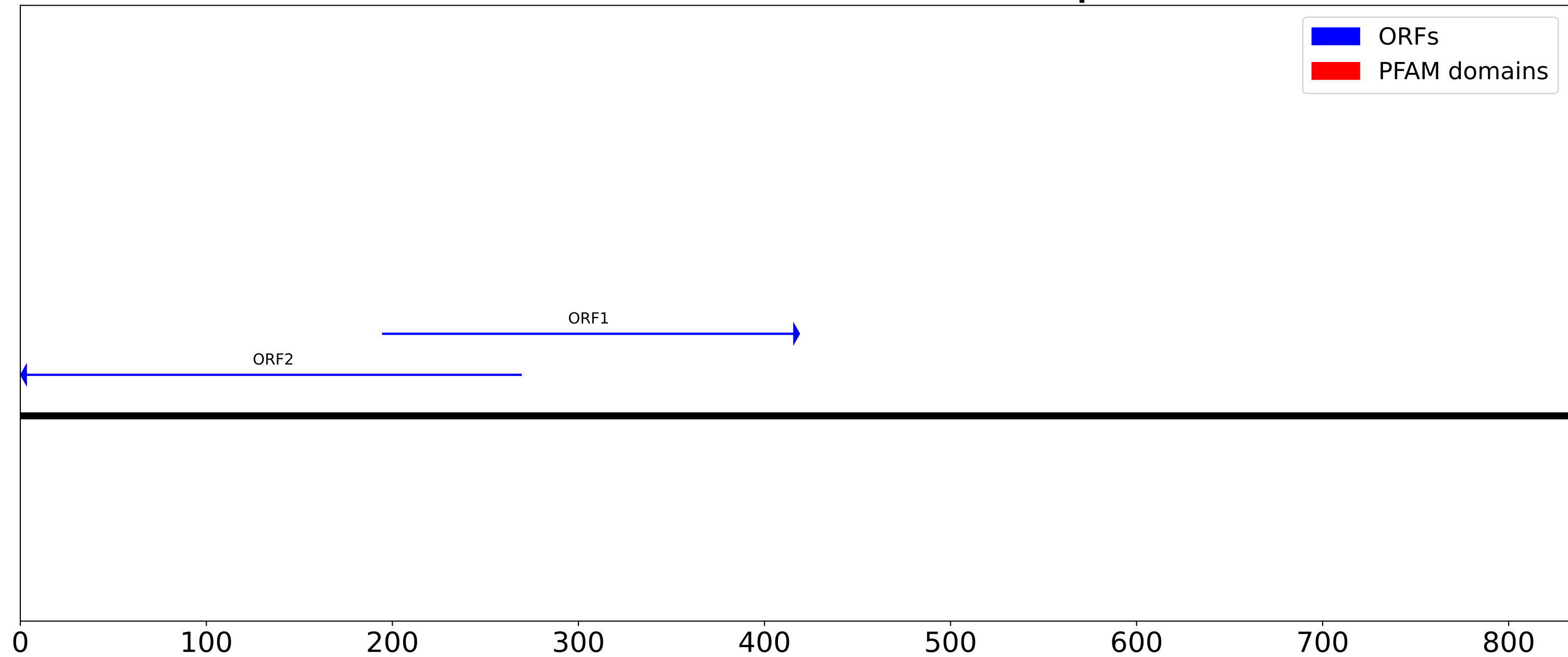


TE: rnd_1_family_944
size: 325bp; fragments: 1125; full length: 299 (≥ 292.5 bp)



Before TEtrimmer 325 bp

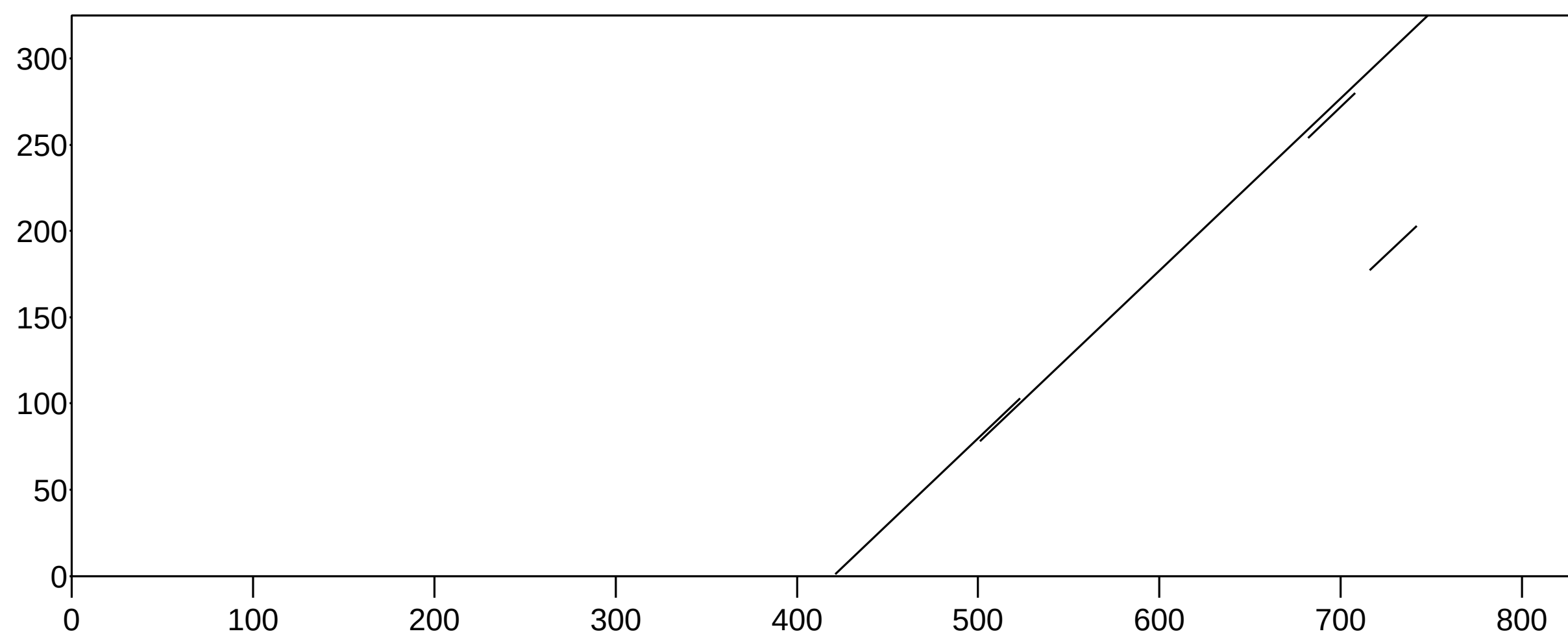
After TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)