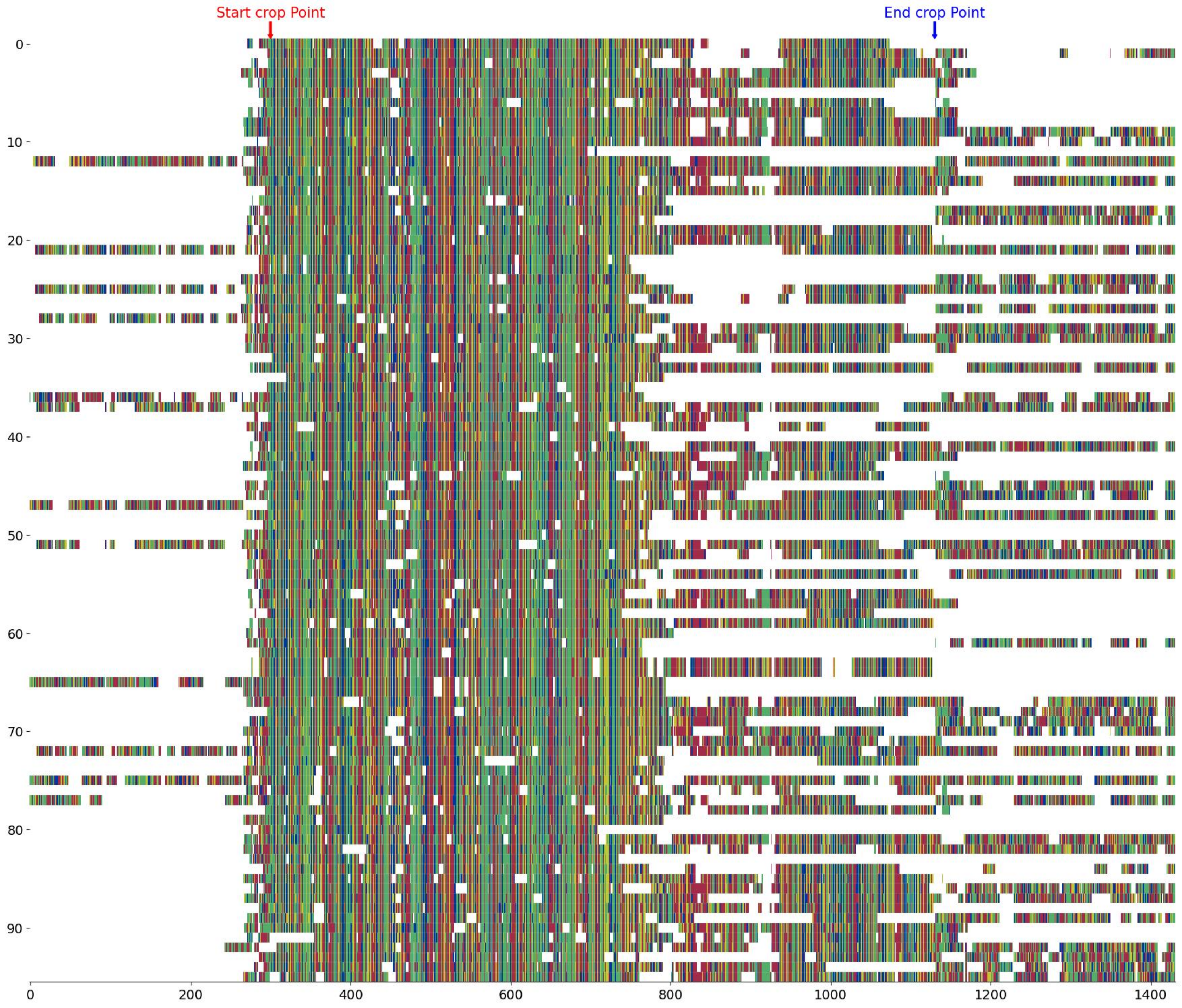




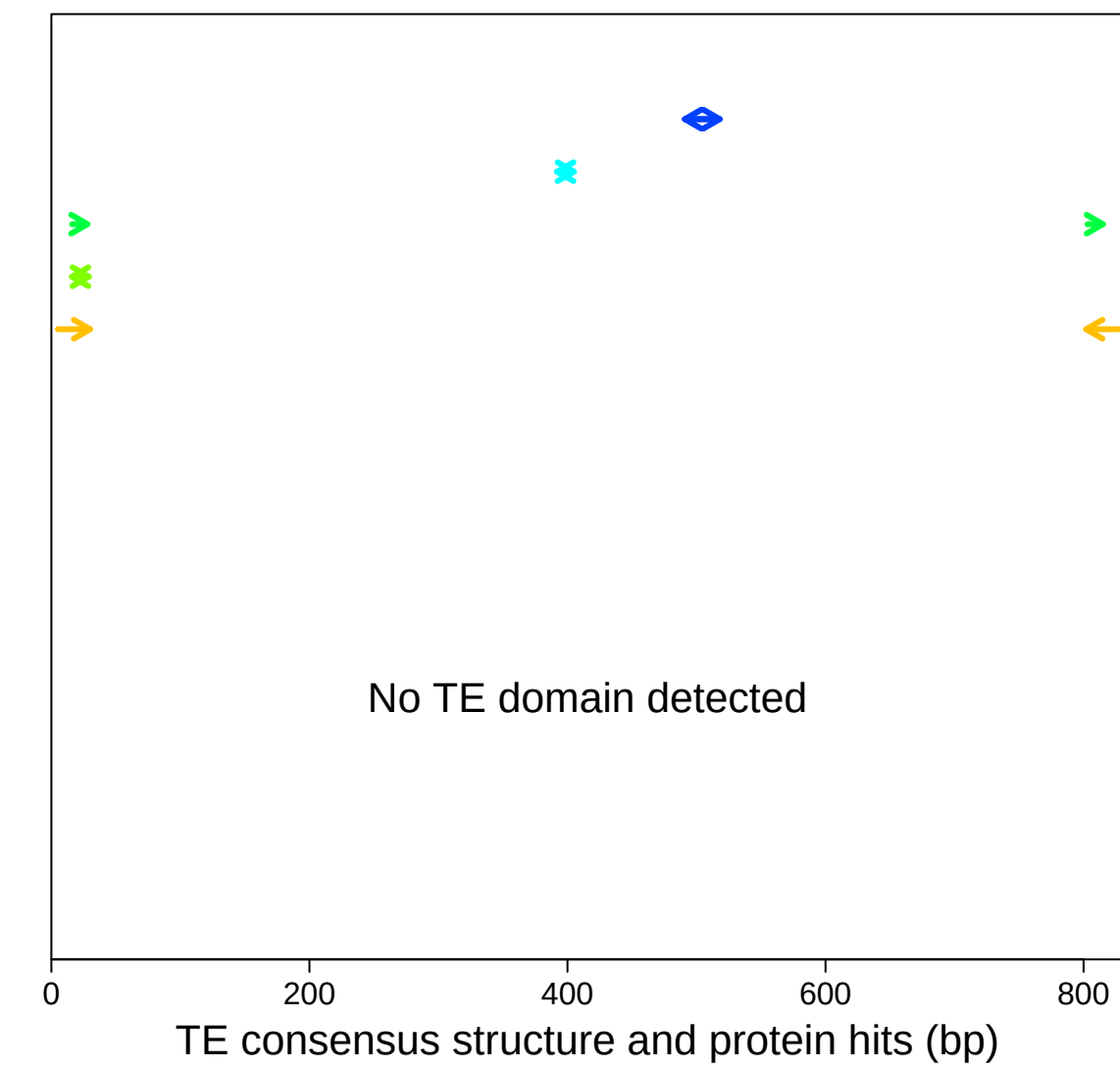
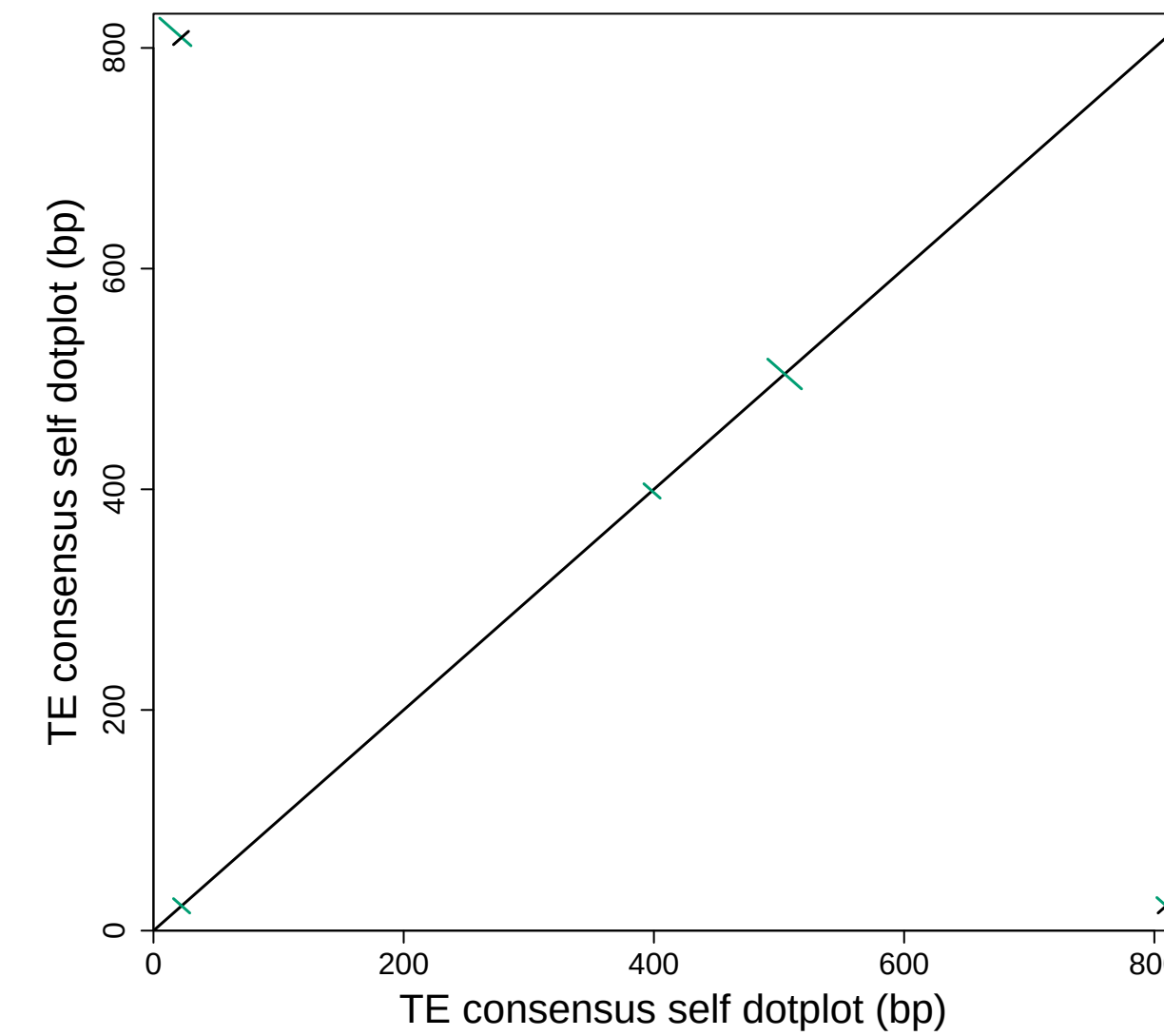
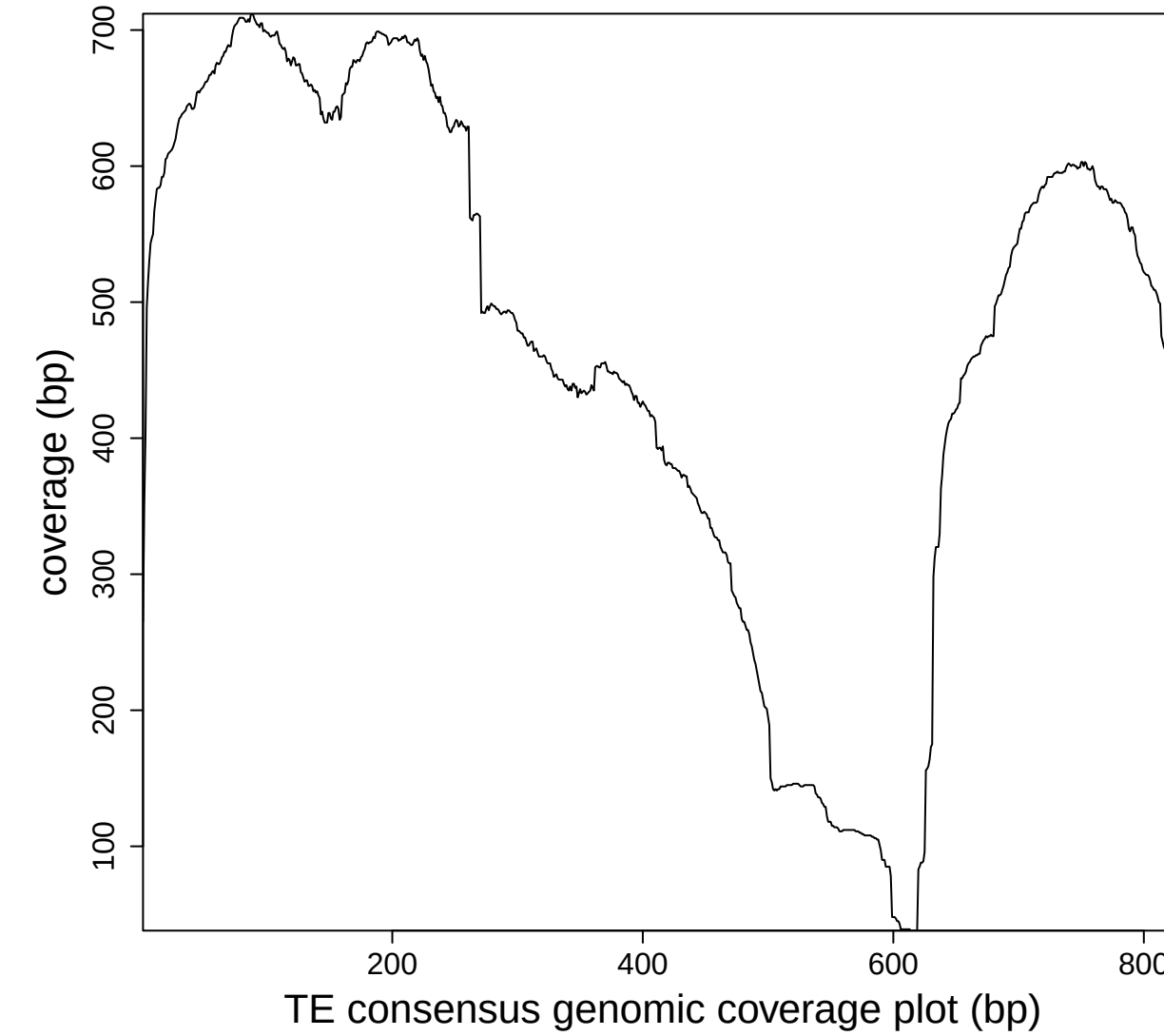
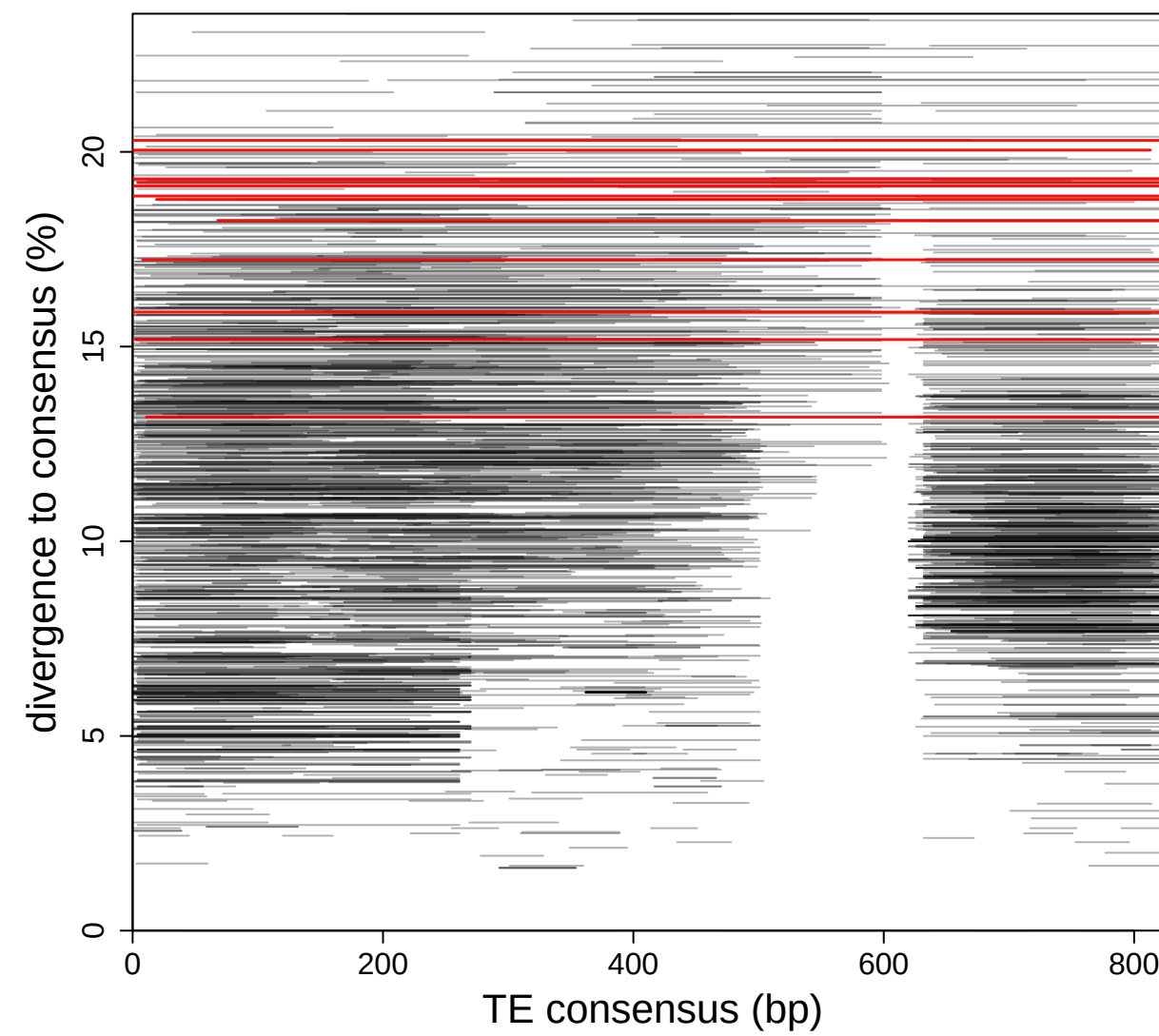
Start crop Point

End crop Point



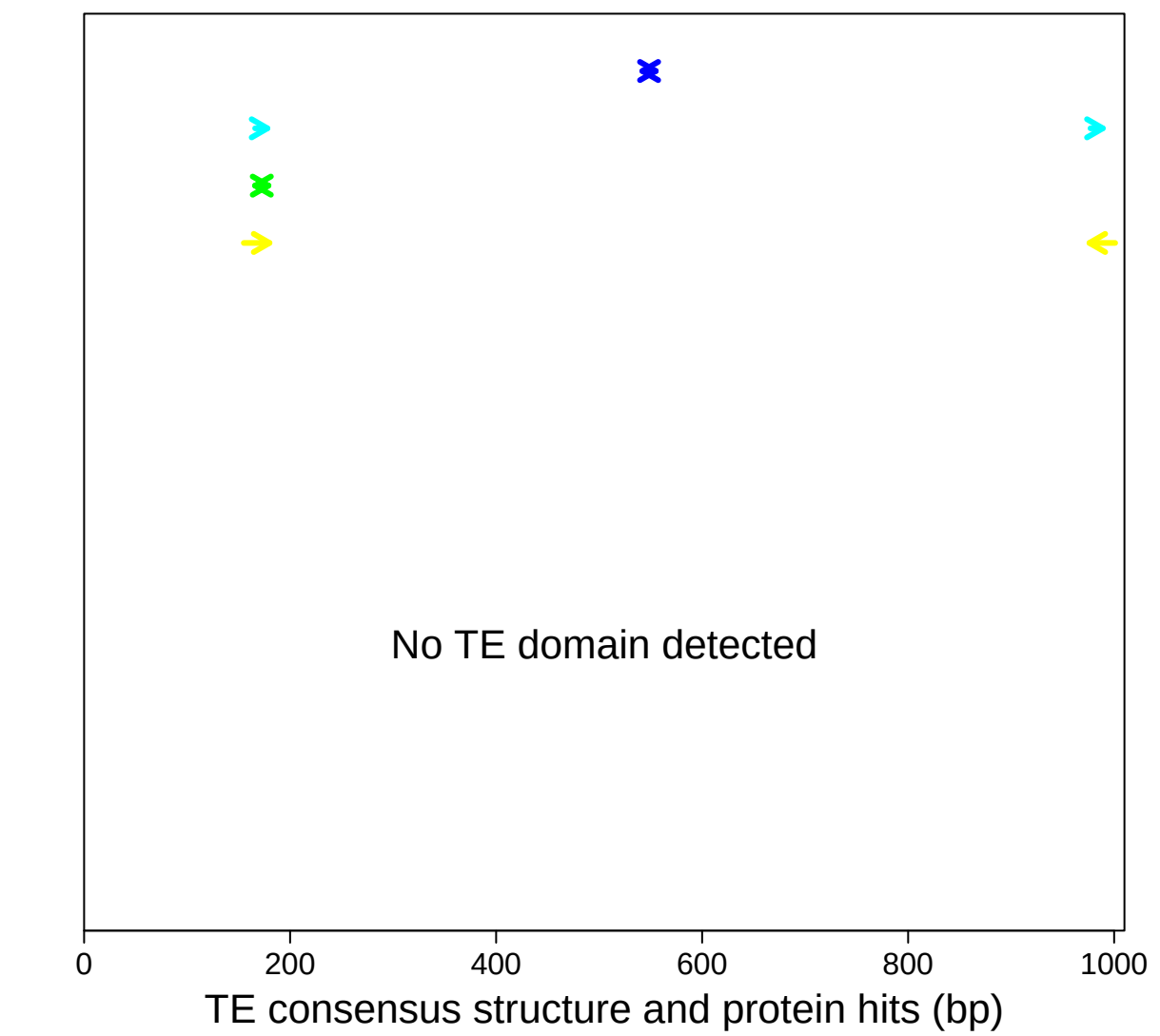
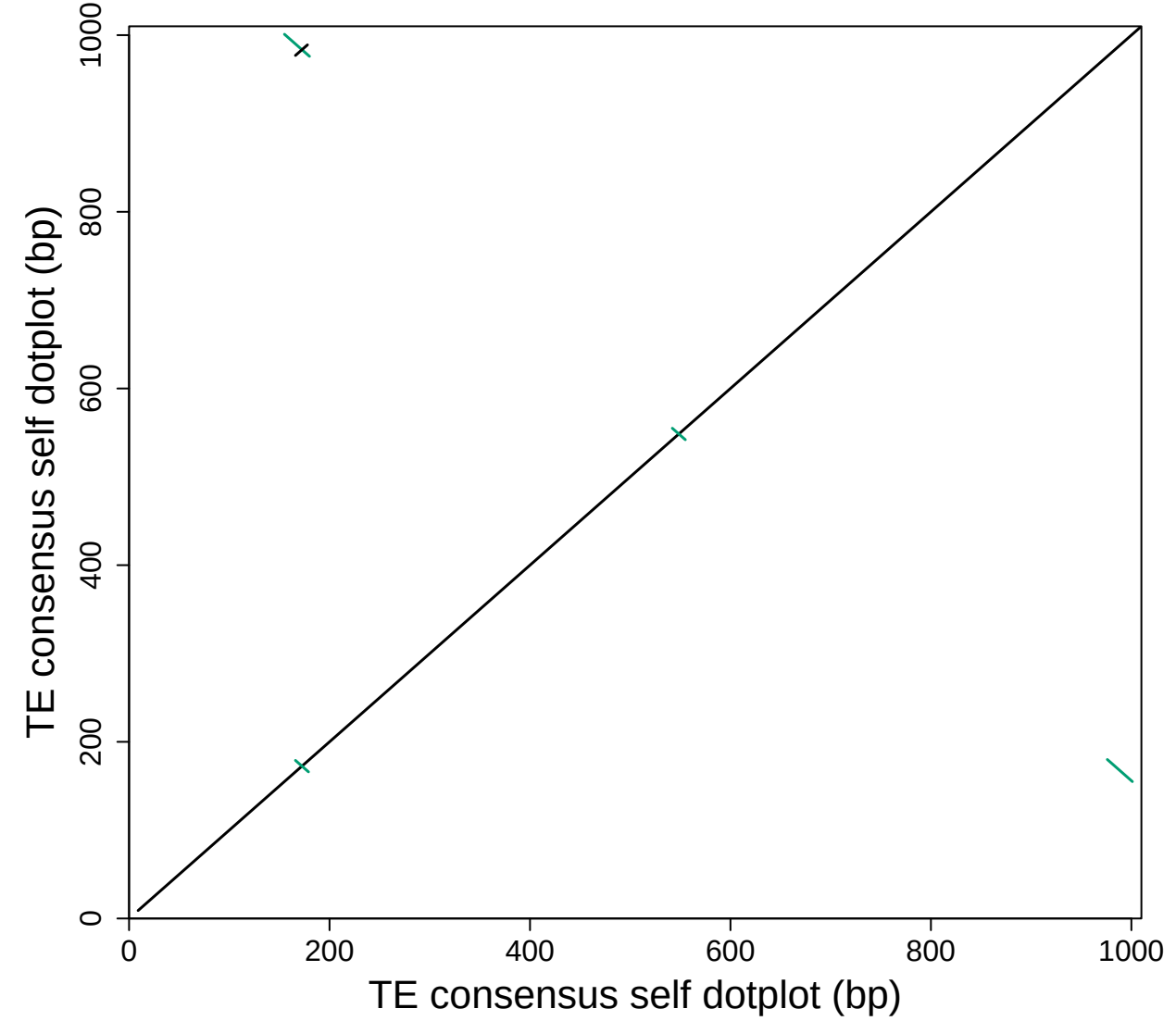
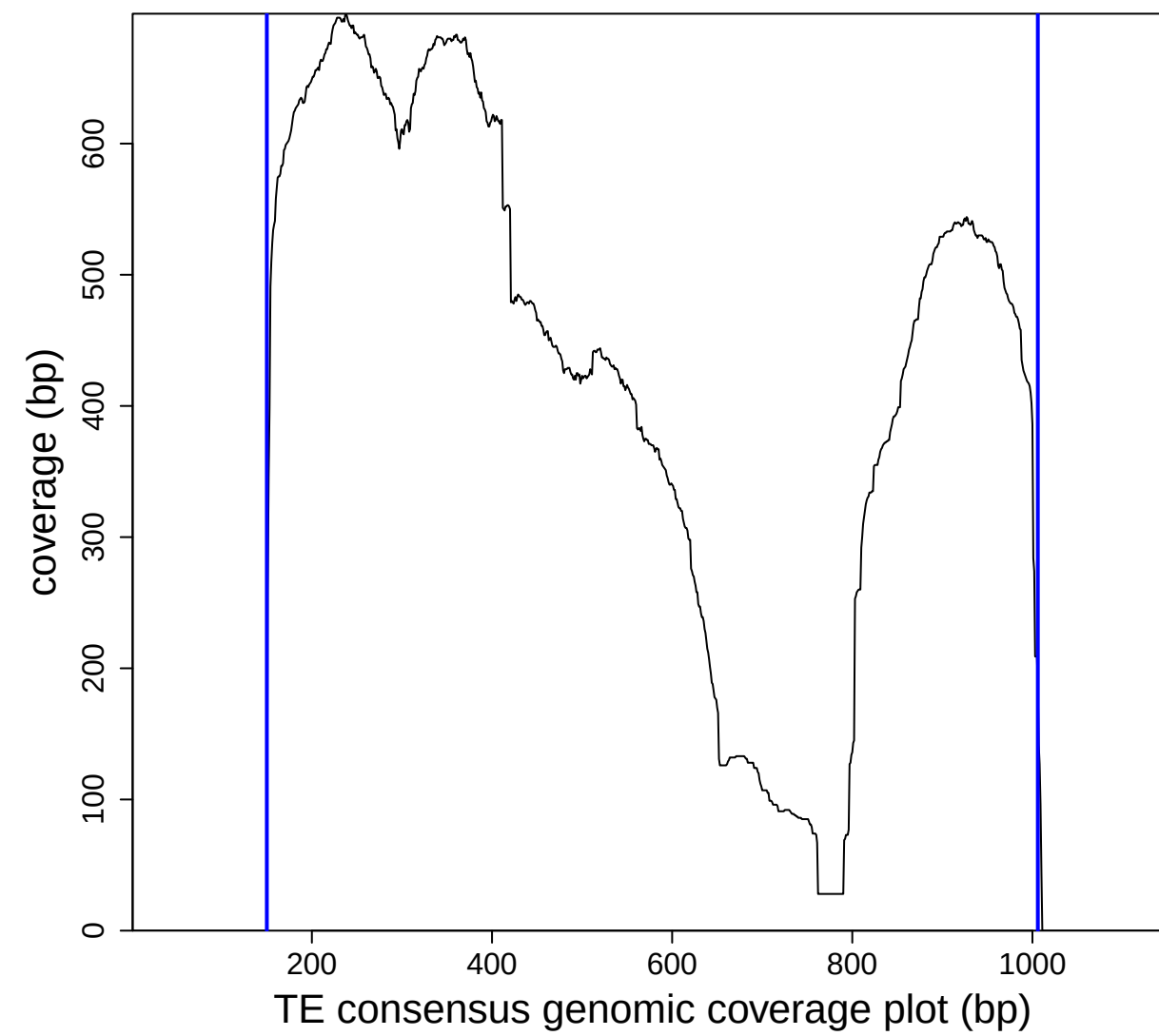
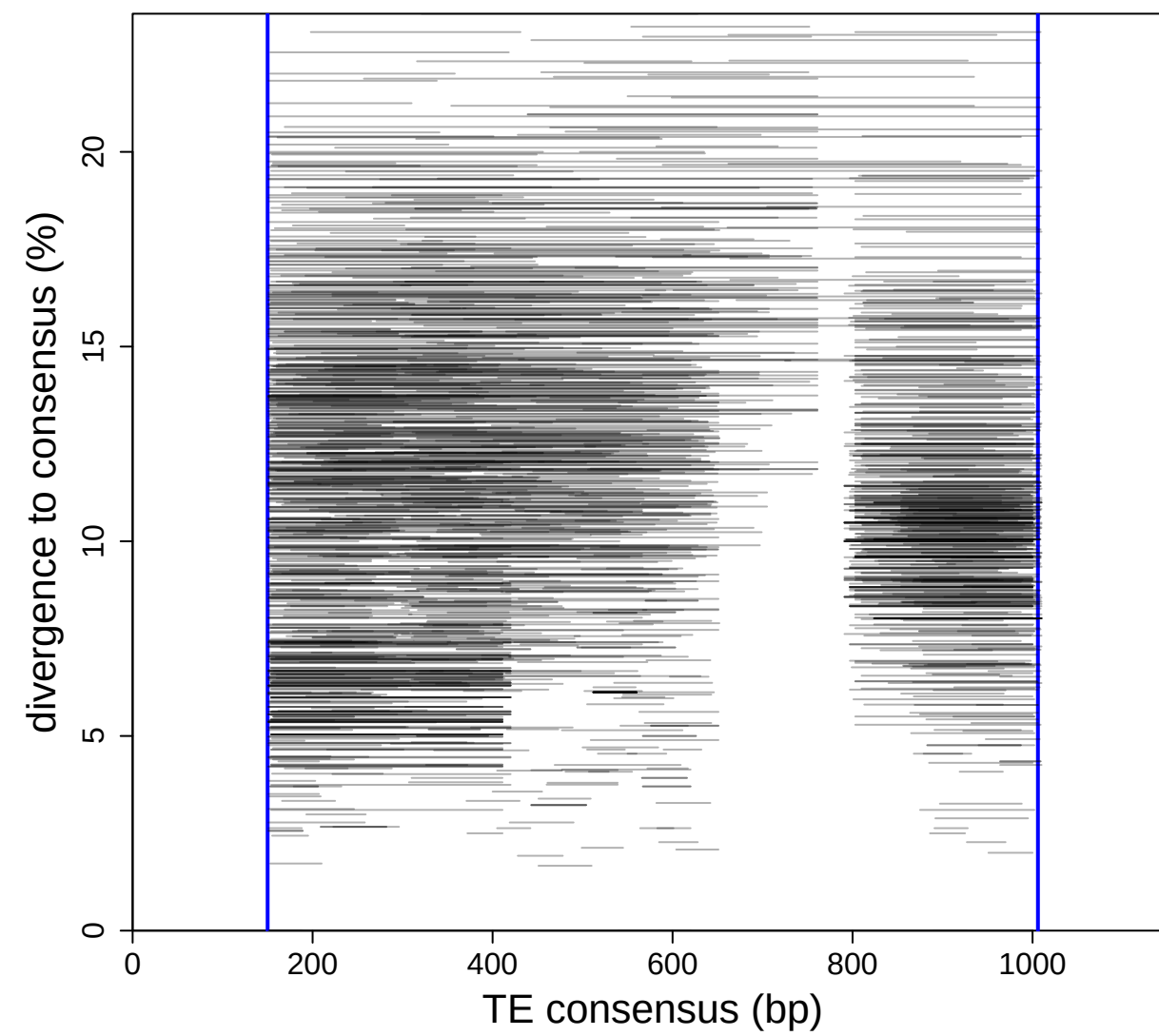
l.b.bed uf.bed g_1.bed fm_1.bed 0 0 bcln.fa aln.fa cl.fa gs.fa
size: 831bp; fragments: 1859; full length: 12 (≥ 747.9 bp)

After TEtrimmer 831 bp



1.fasta.b.bed uf.bed g 1.bed fm 1.bed 0 0 bcIn.fa aln.fa cl.fa
size: 1156bp; fragments: 1753; full length: 0 (>=1040.4bp)

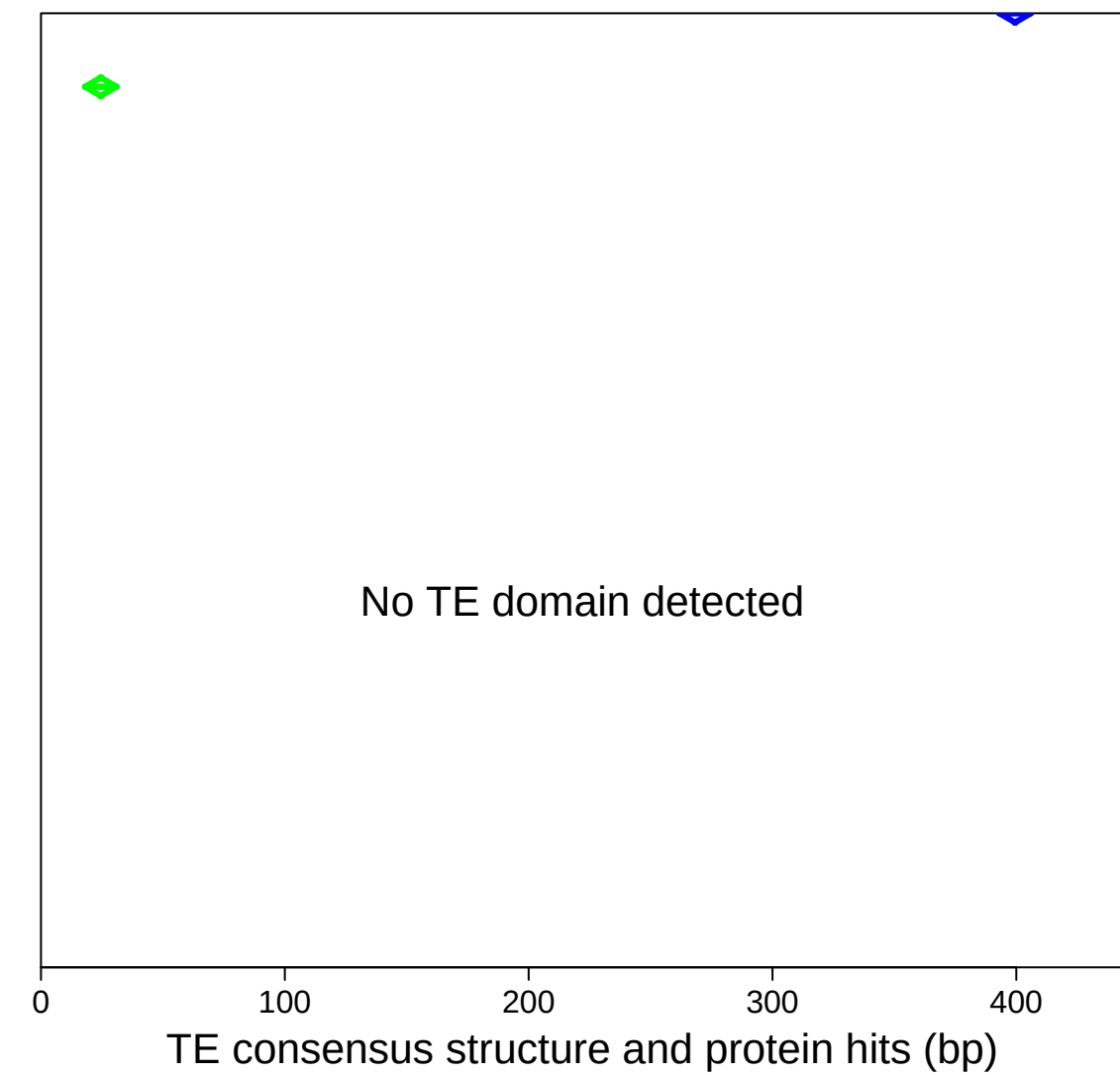
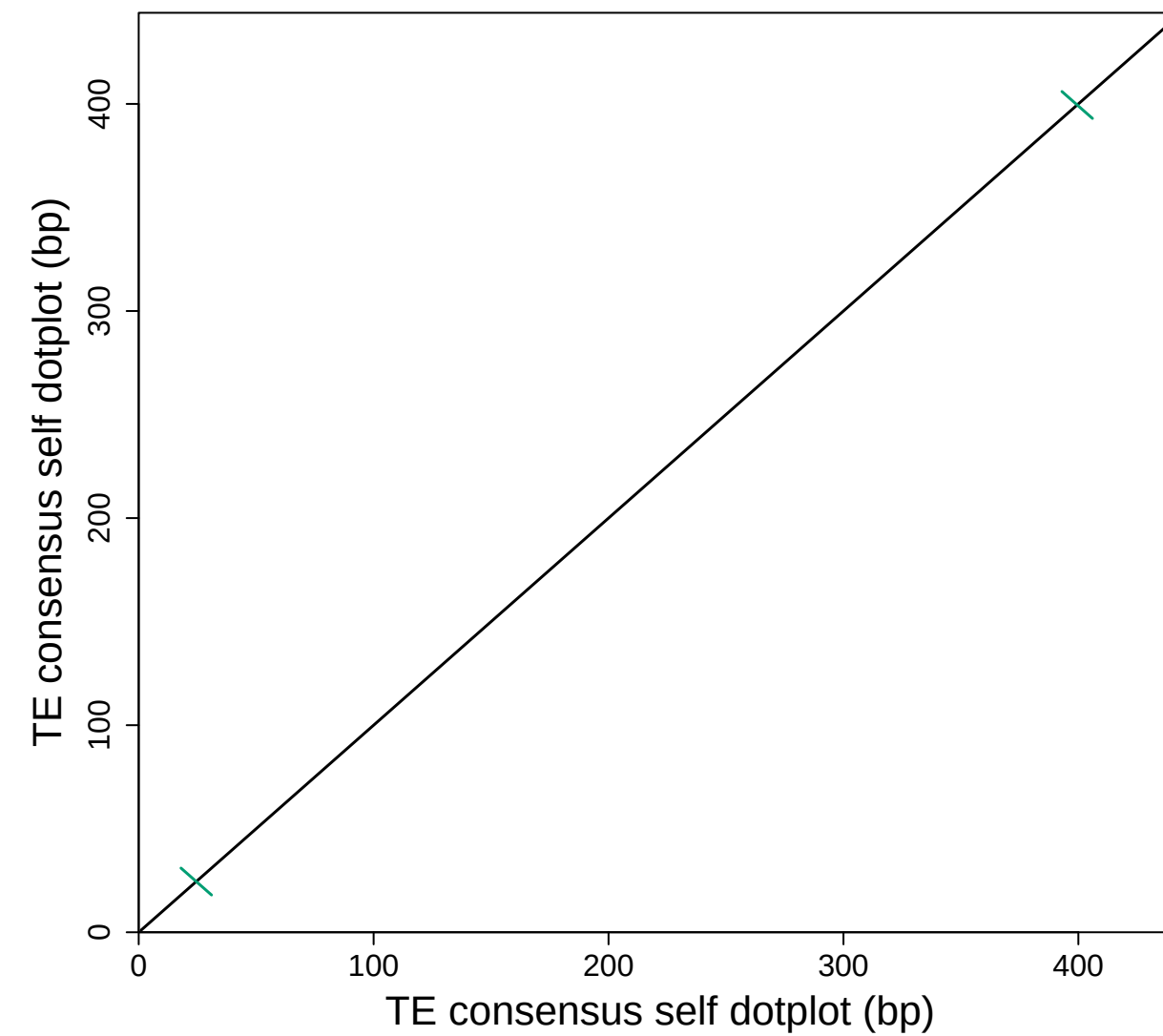
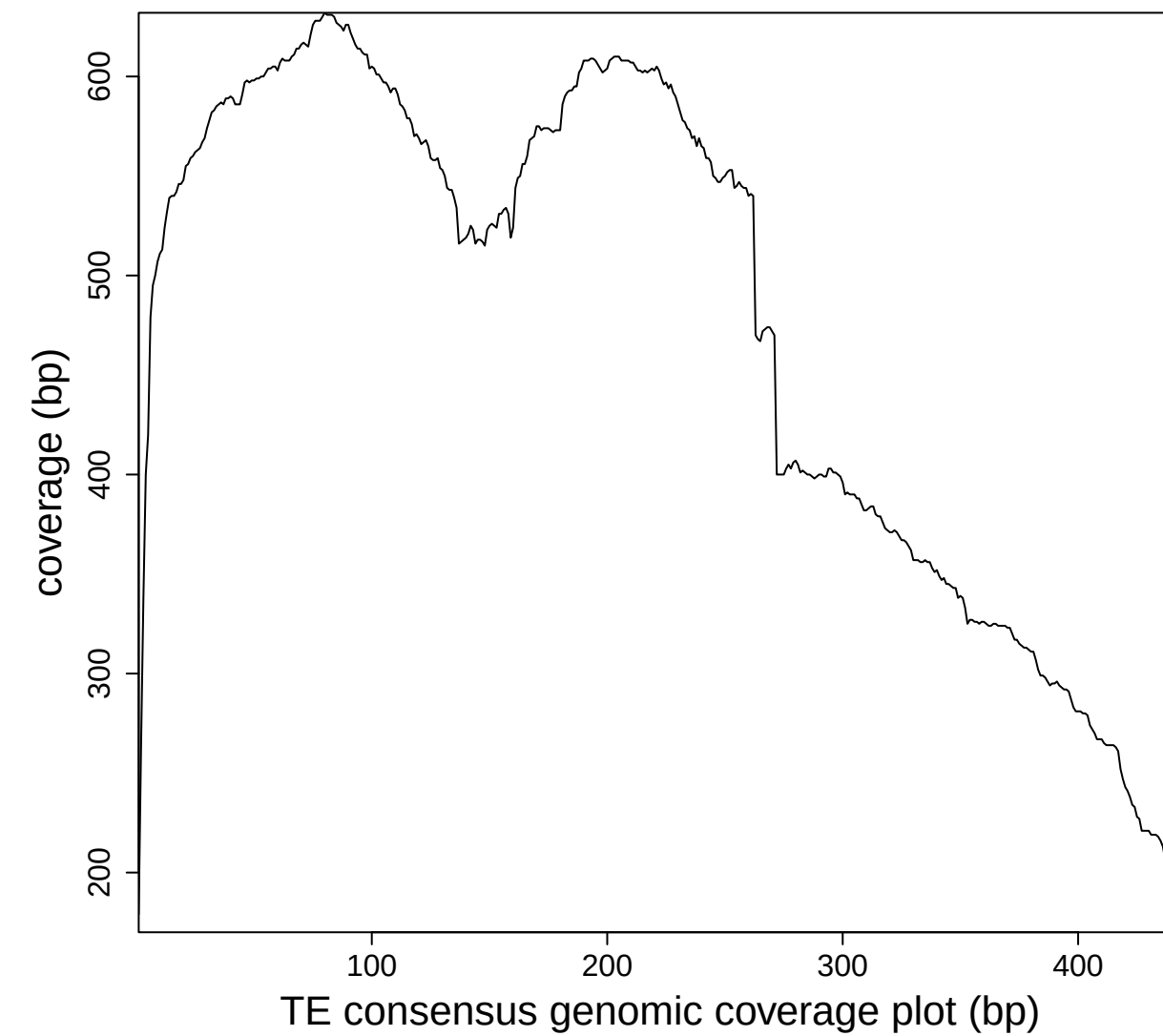
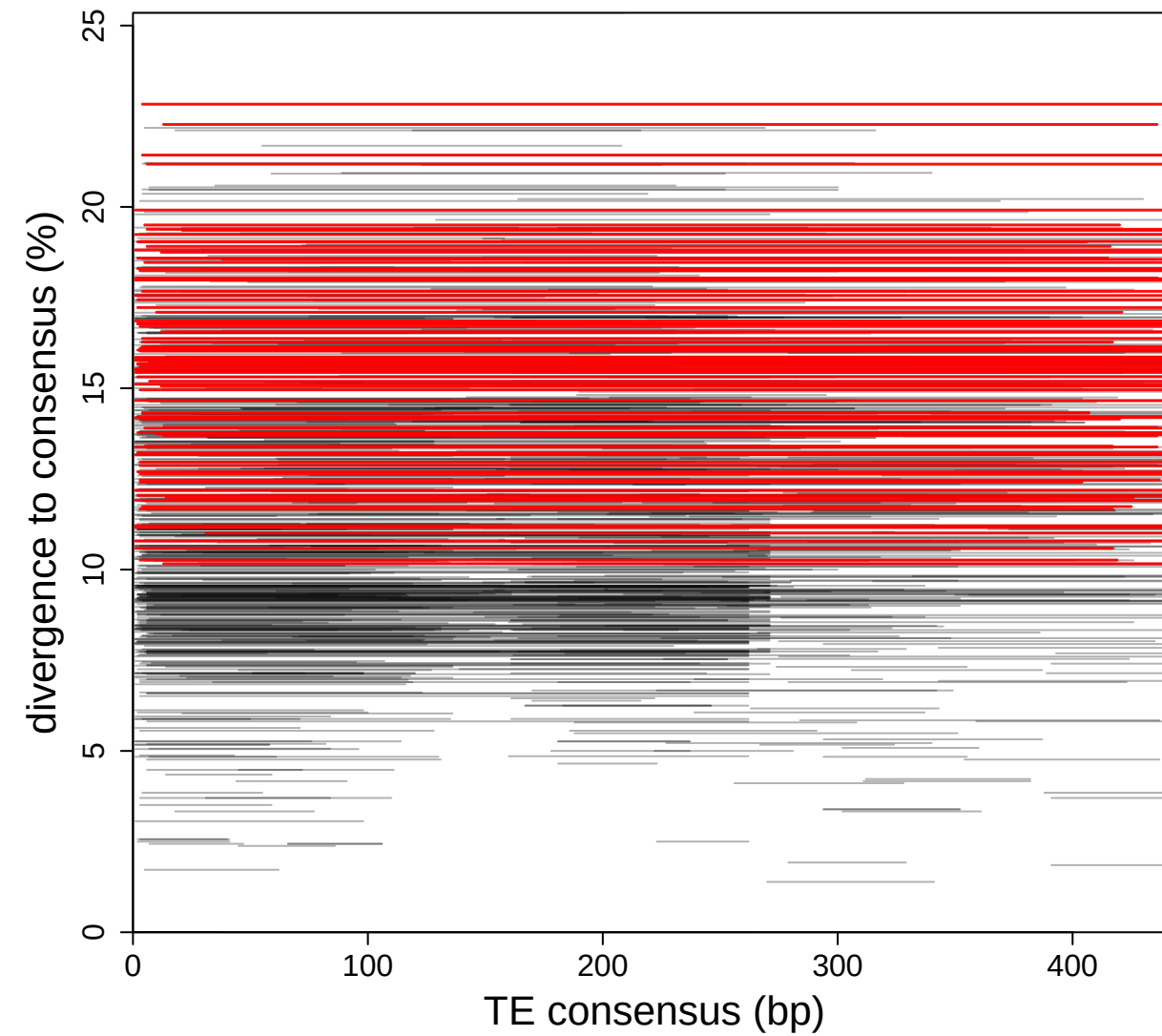
After TETrimmer Extended plot Blue lines are boundaries



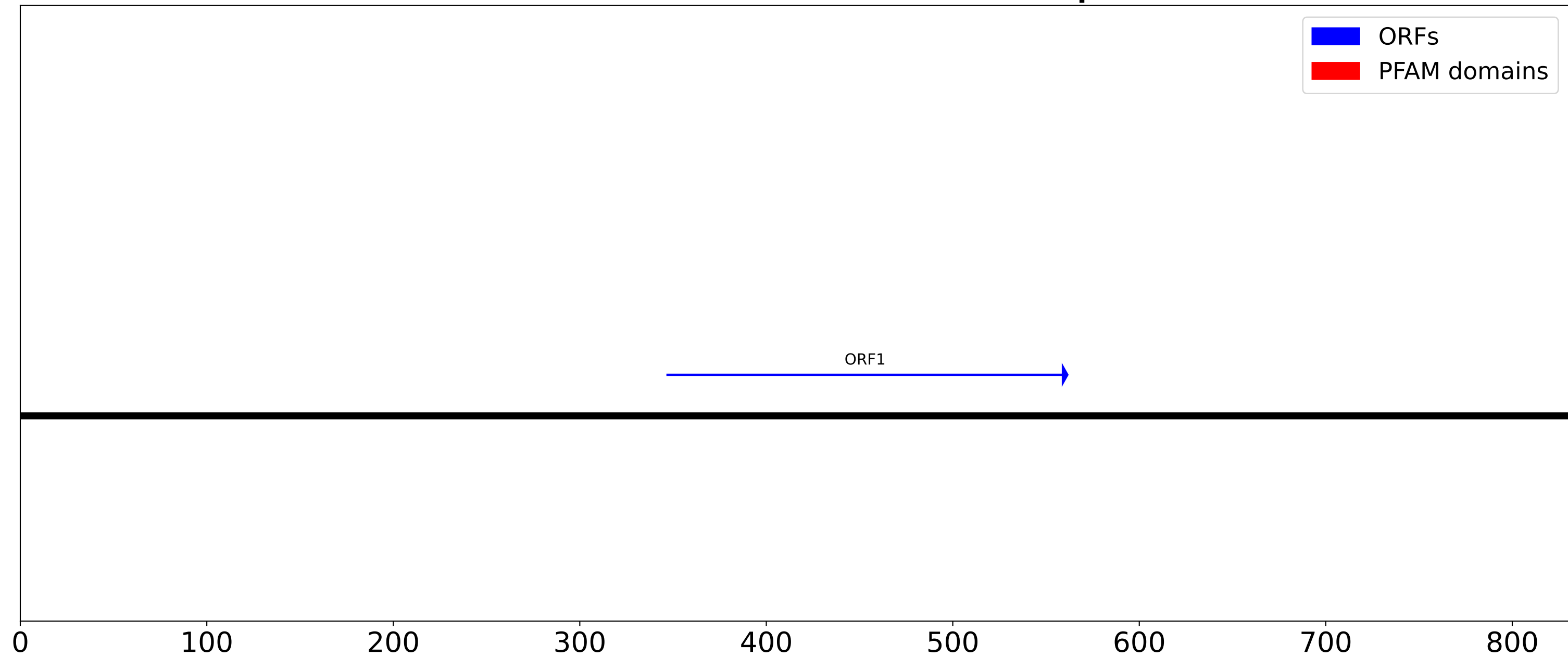
Before TEtrimmer 444 bp

TE: rnd_3_family_351

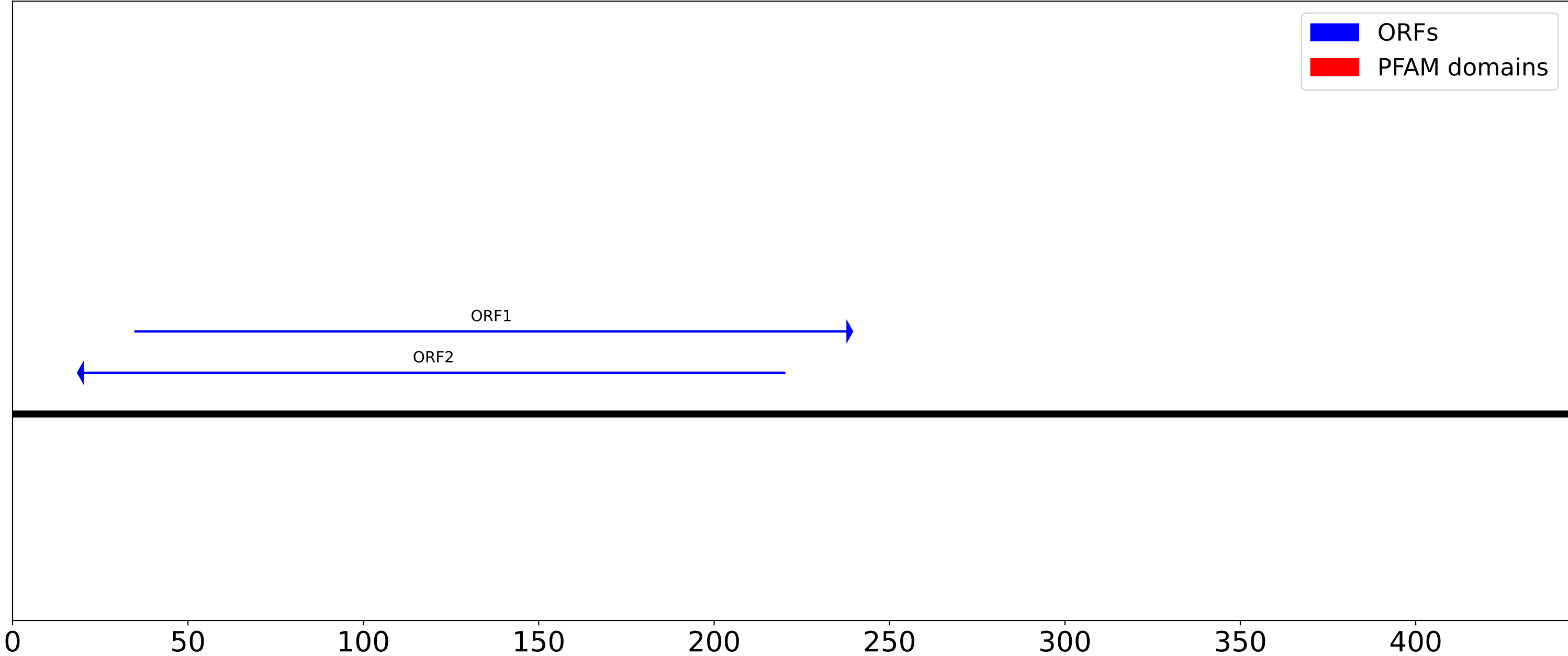
size: 444bp; fragments: 968; full length: 101 (≥ 399.6 bp)



After TEtrimmer ORF and PFAM domain plot



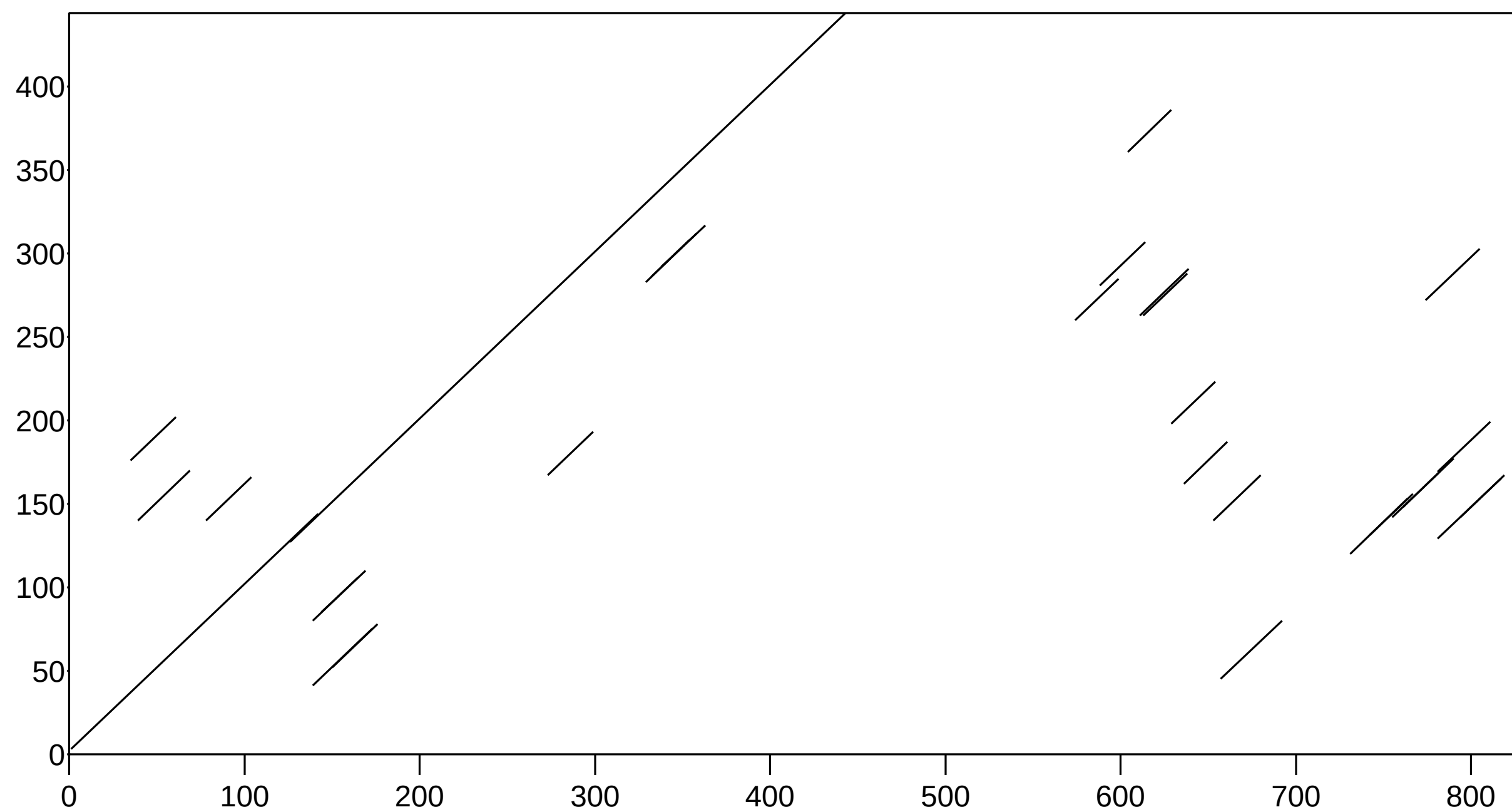
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)