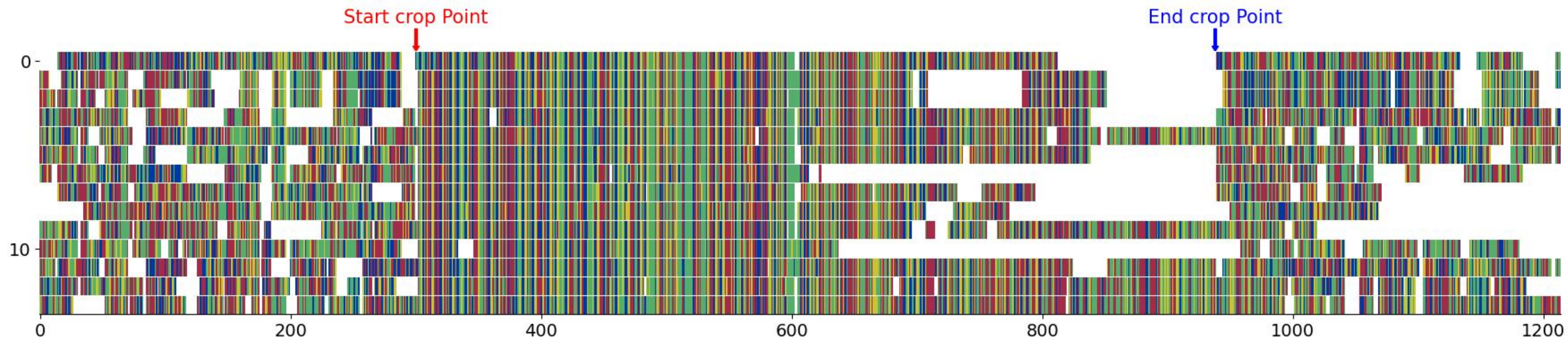
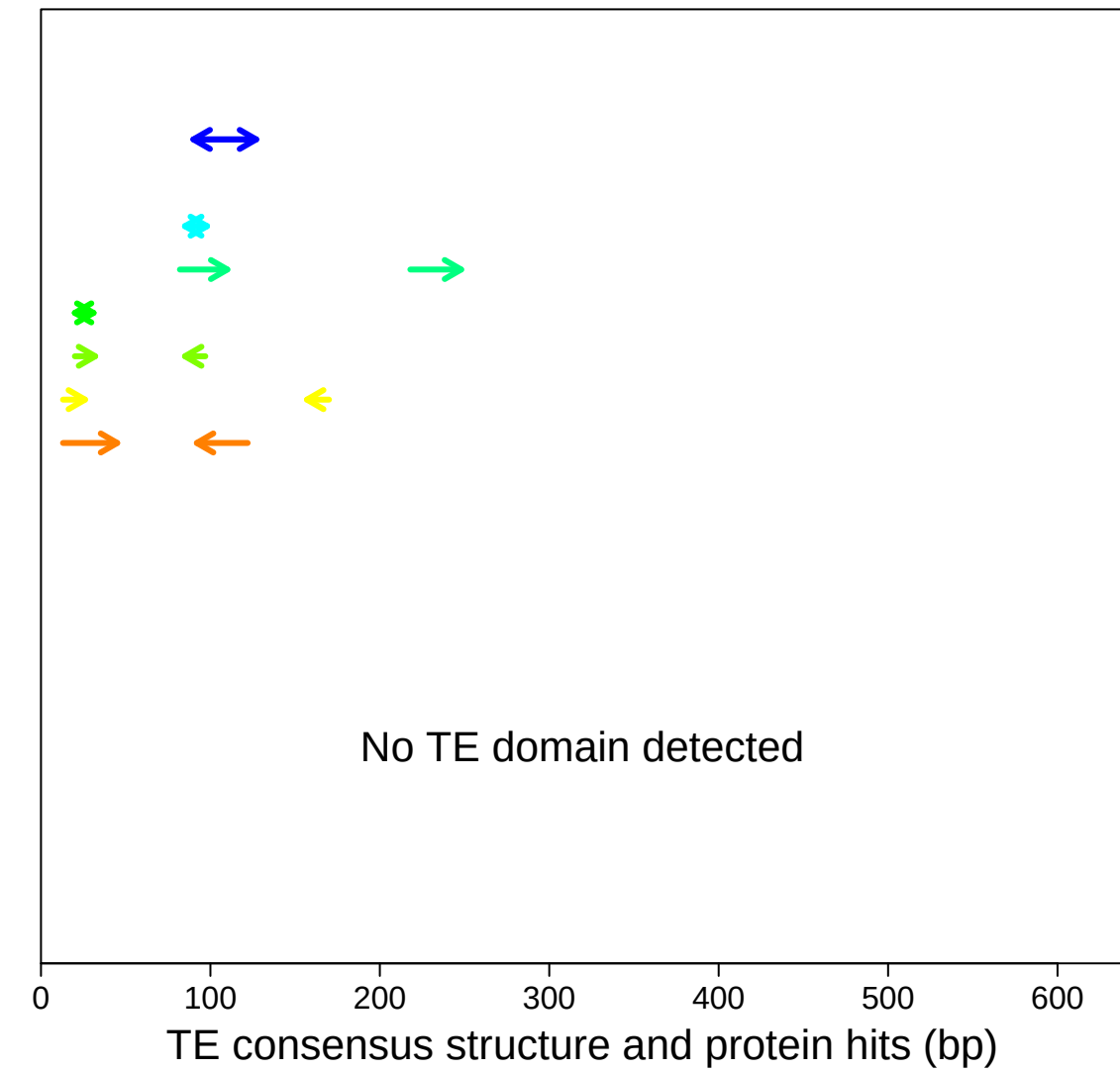
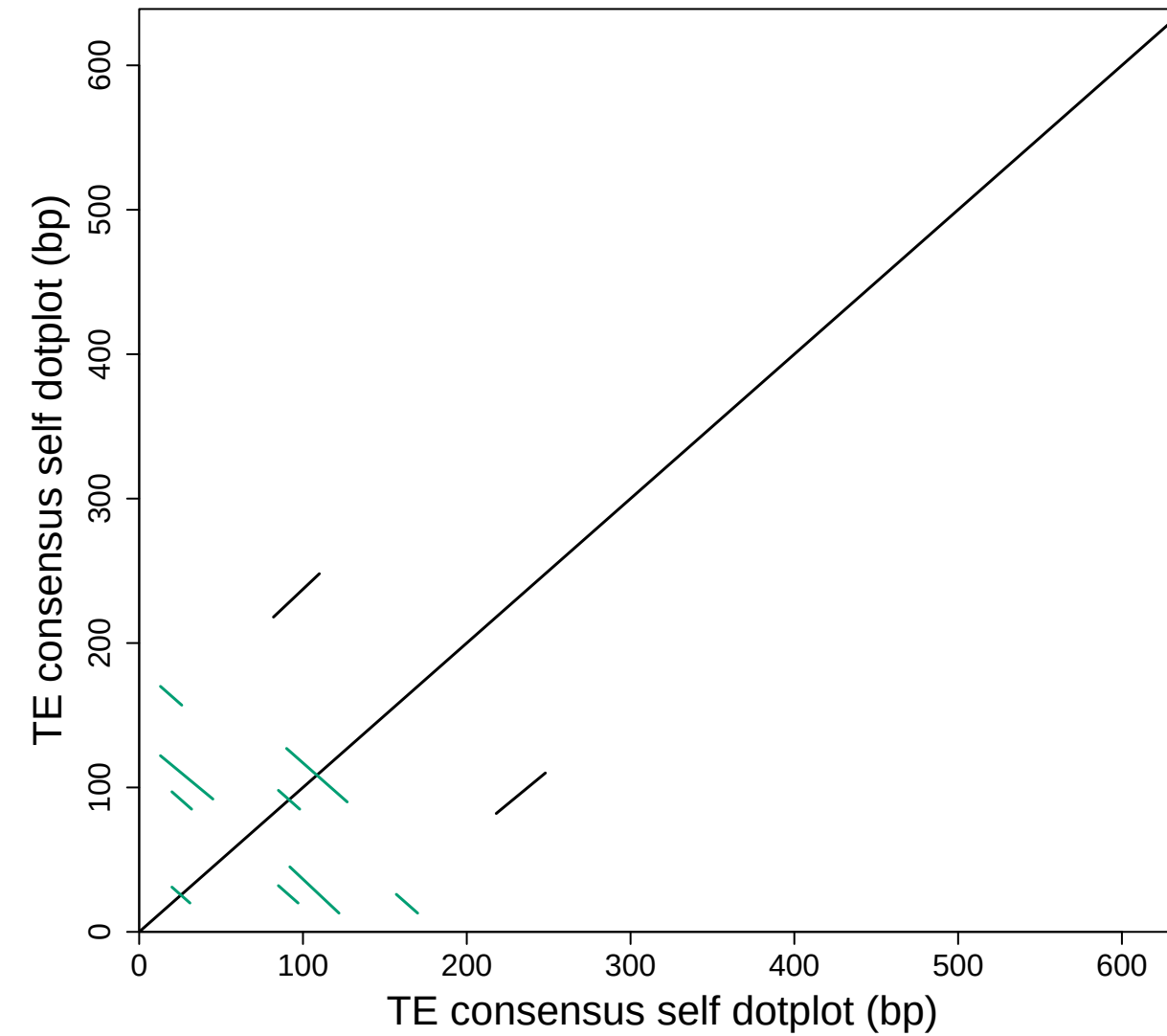
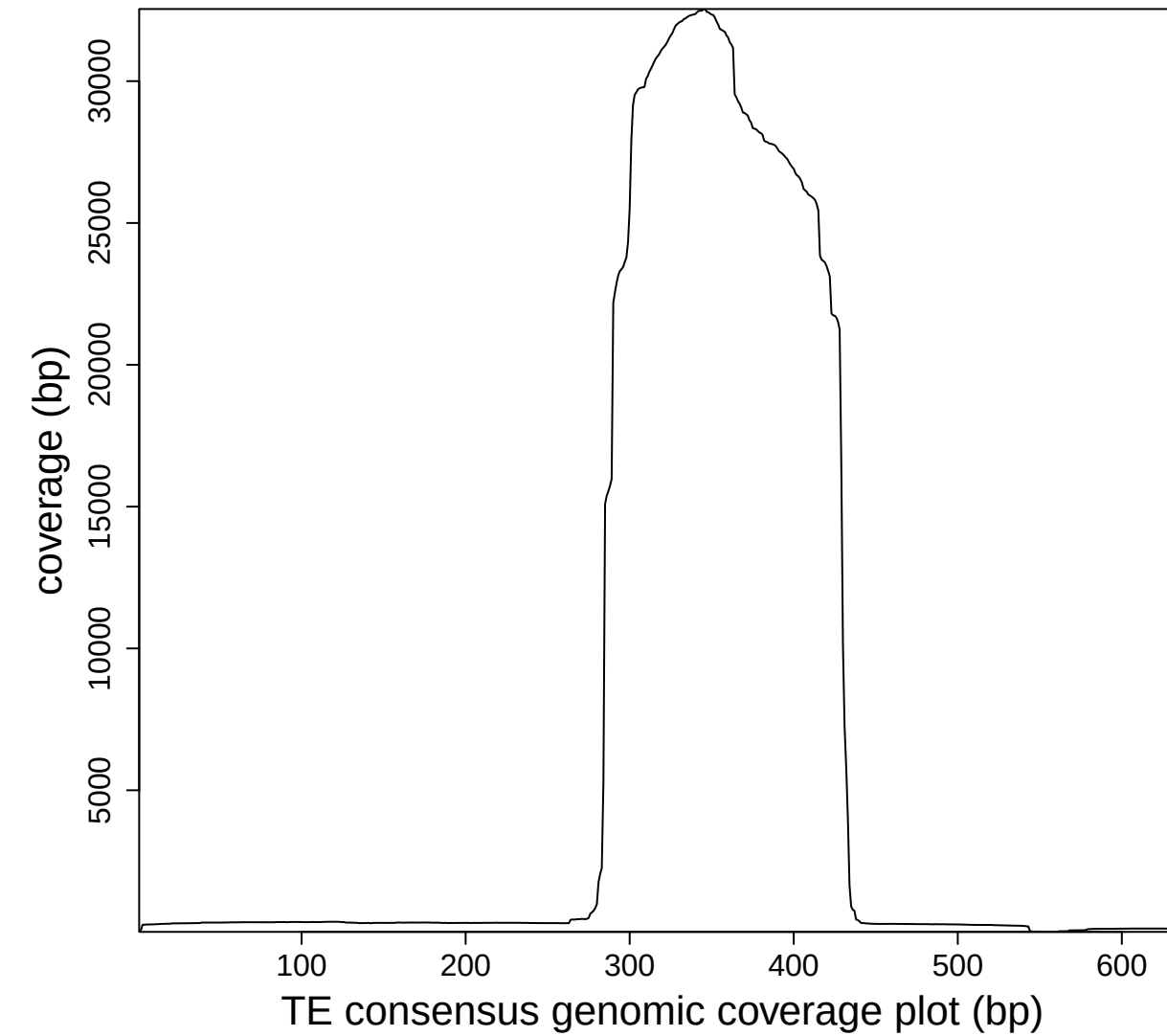
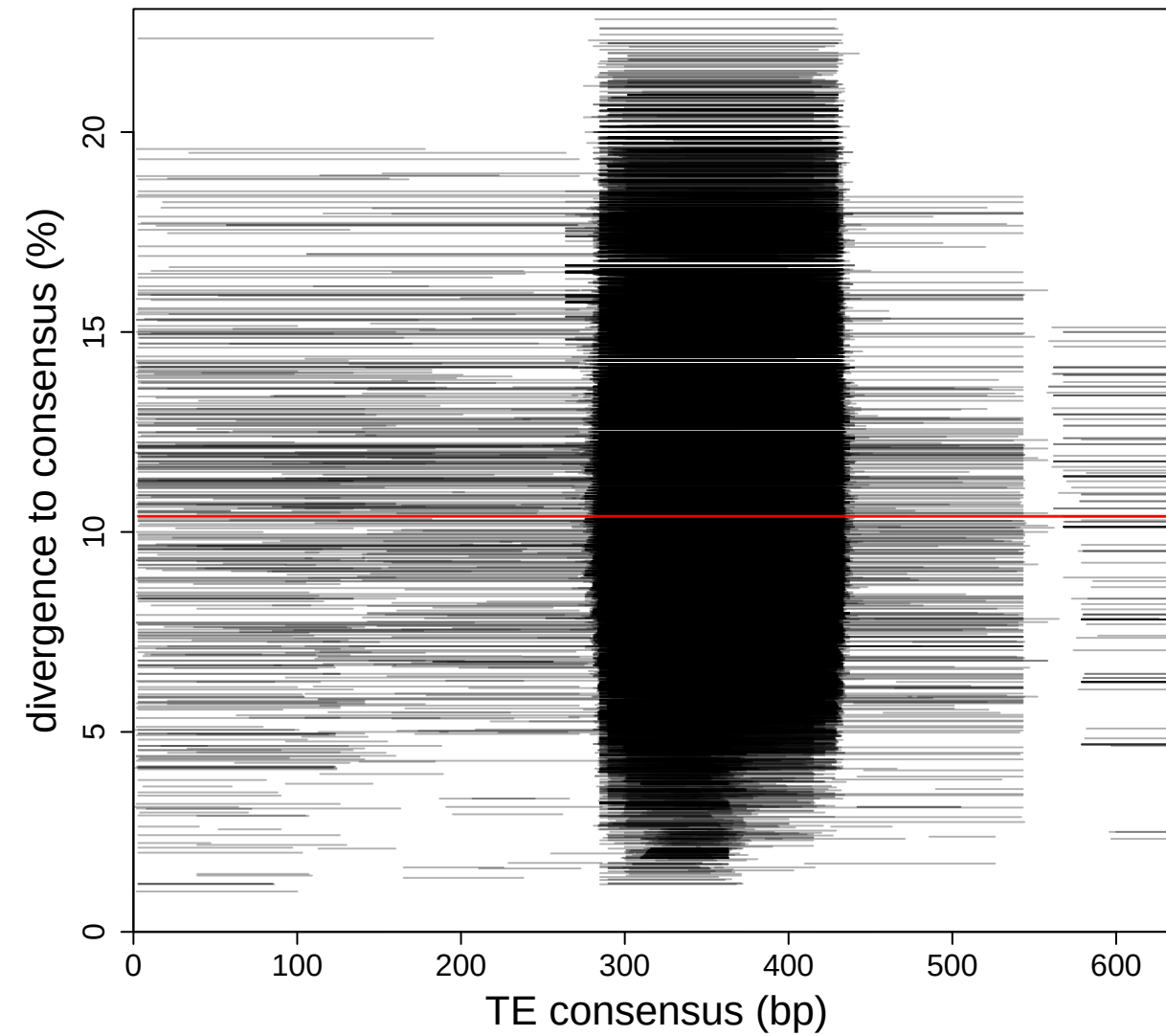




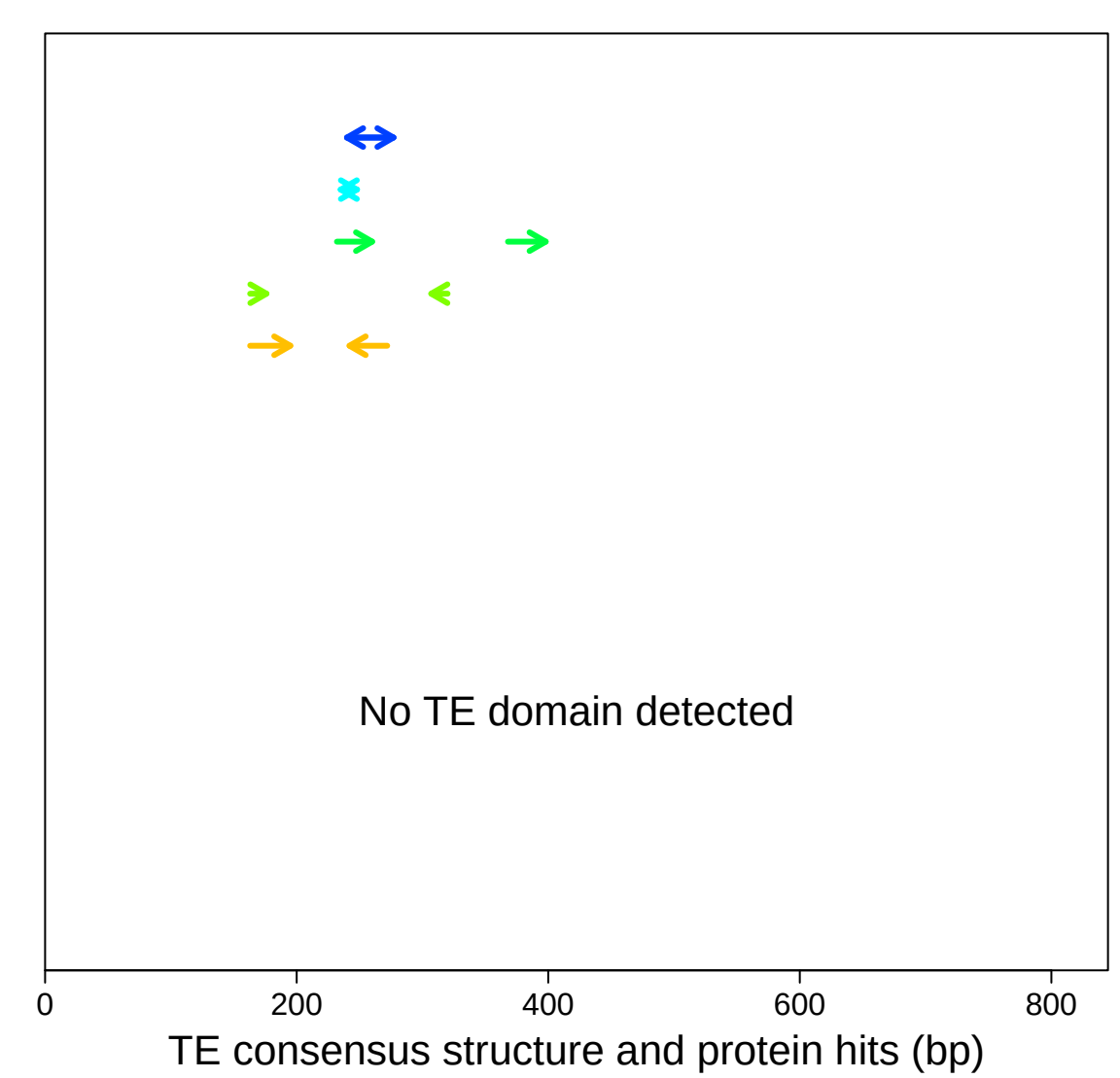
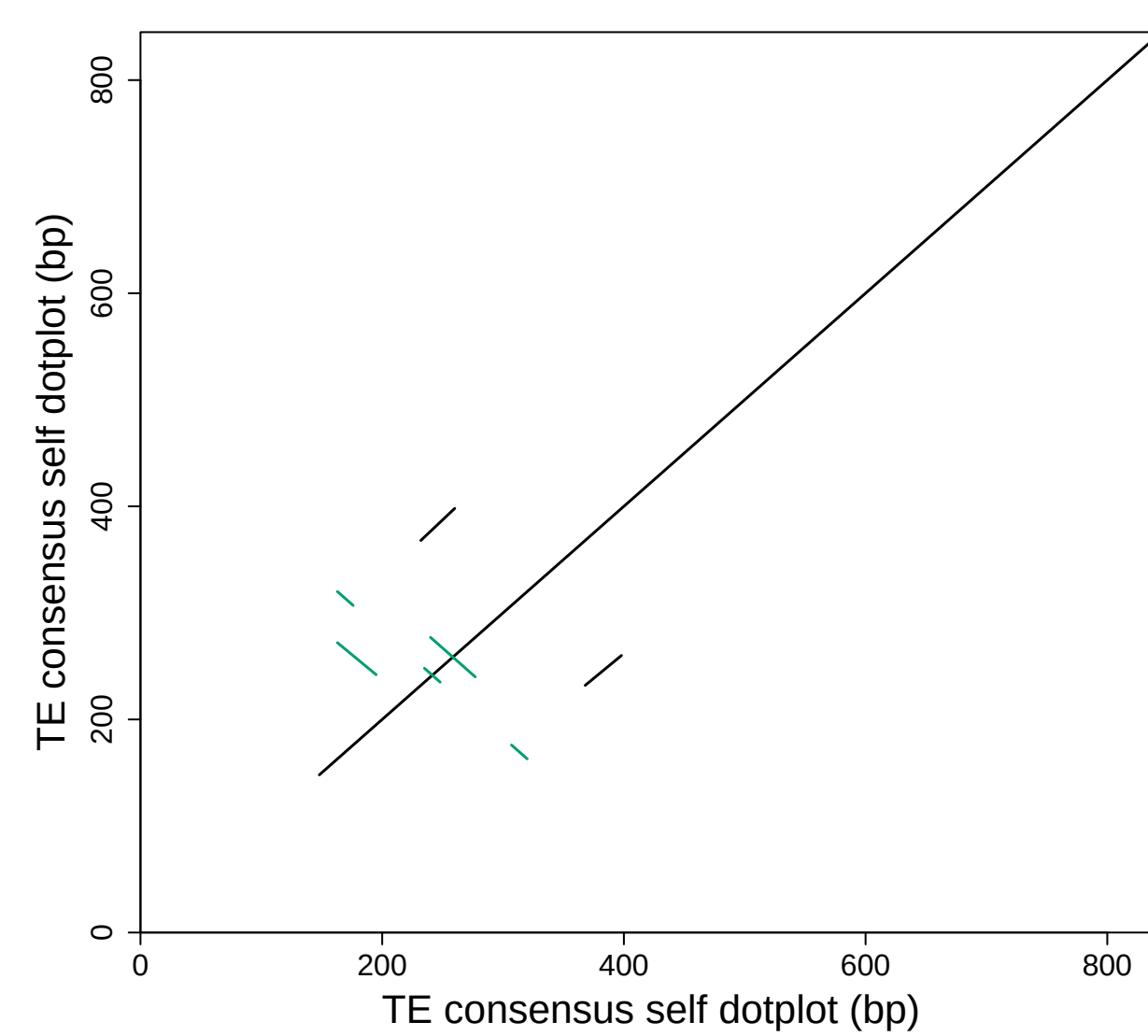
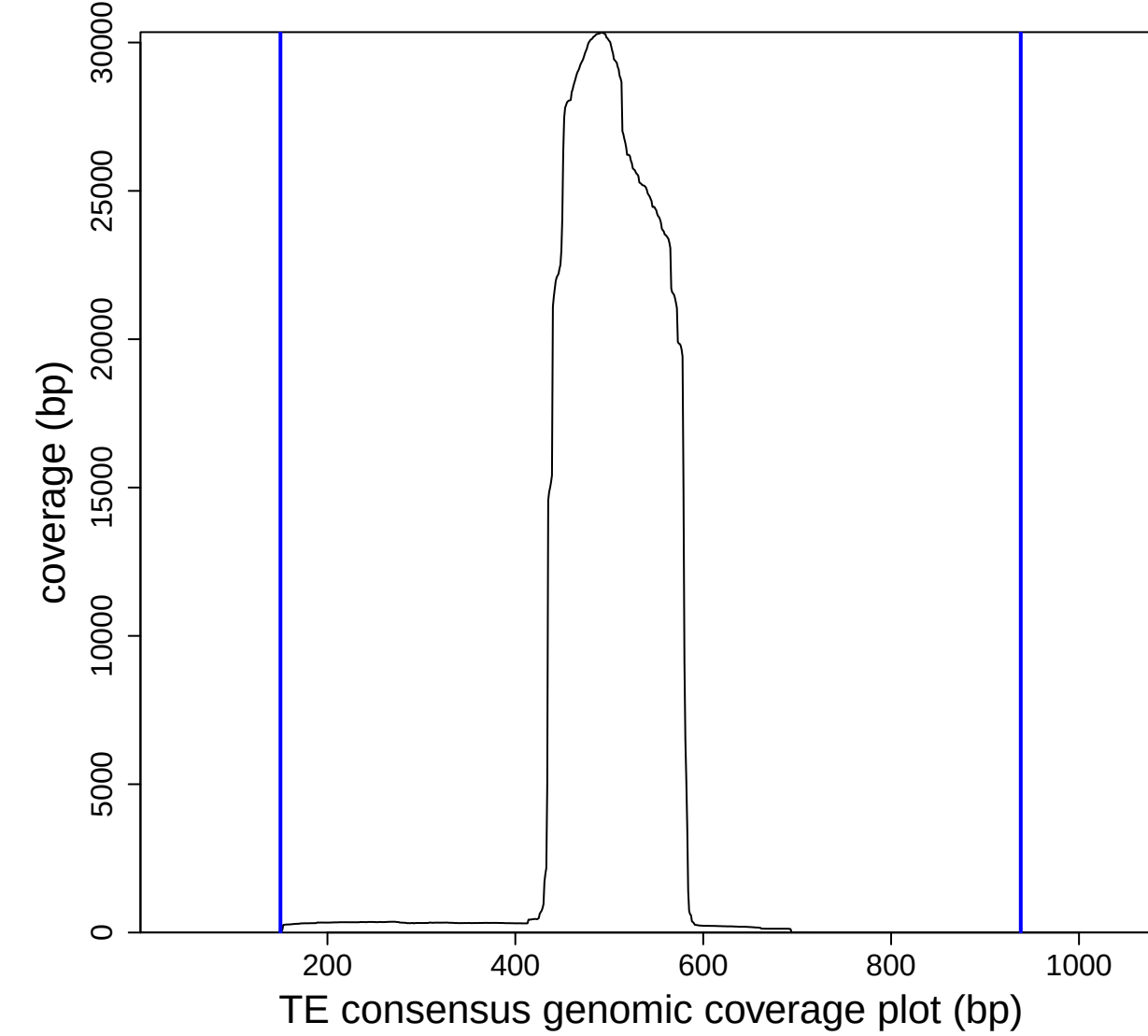
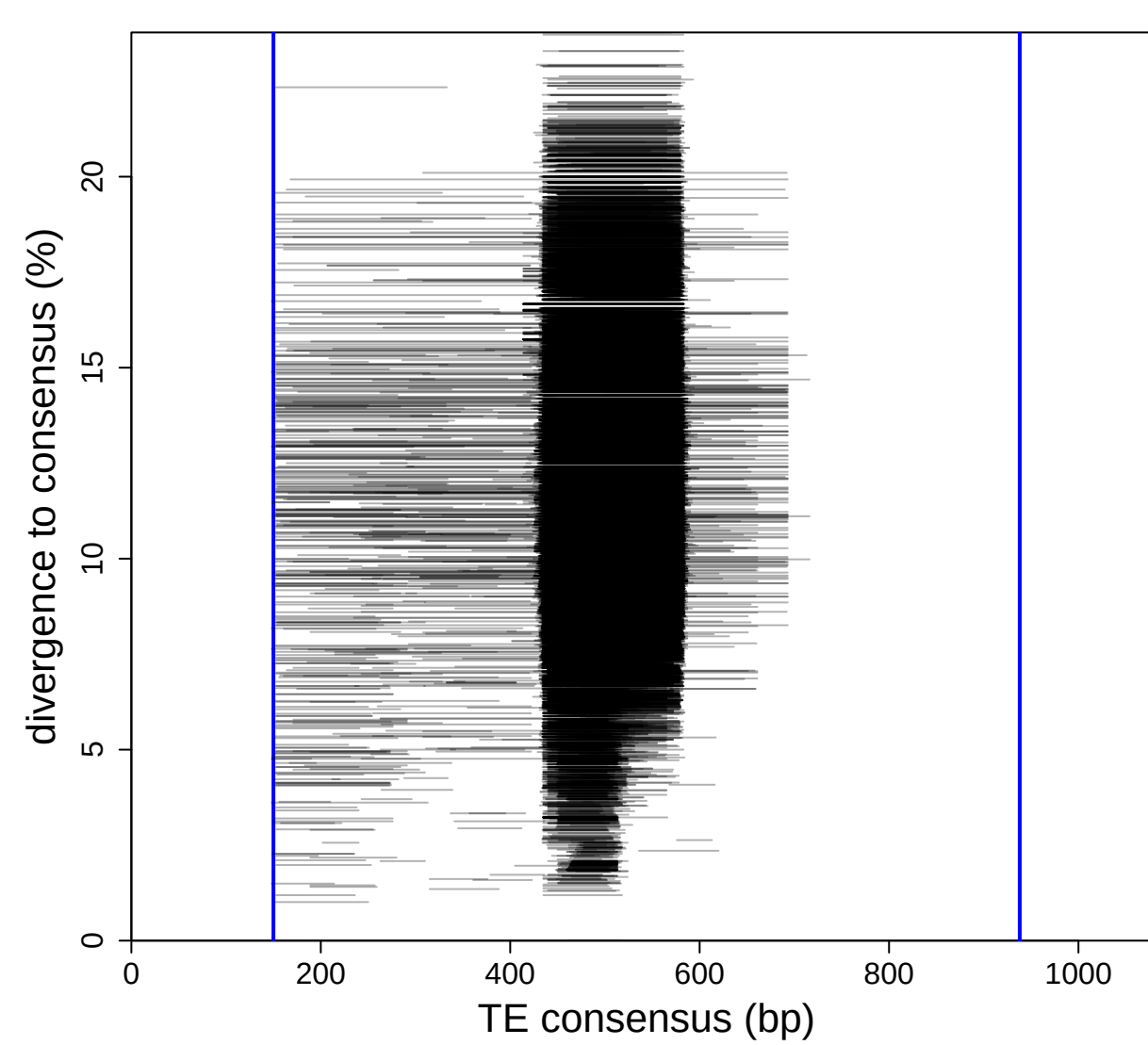
l.b.bed uf.bed g_3.bed fm_1.bed 0 0 bcIn.fa aln.fa cl.fa gs.fa
size: 639bp; fragments: 33400; full length: 1 (>=575.1bp)

After TEtrimmer 639 bp



5.fasta.b.bed uf.bed g 3.bed fm 1.bed 0 0 bcln.fa aln.fa cl.fa
size: 1088bp; fragments: 30785; full length: 0 (>=979.2bp)

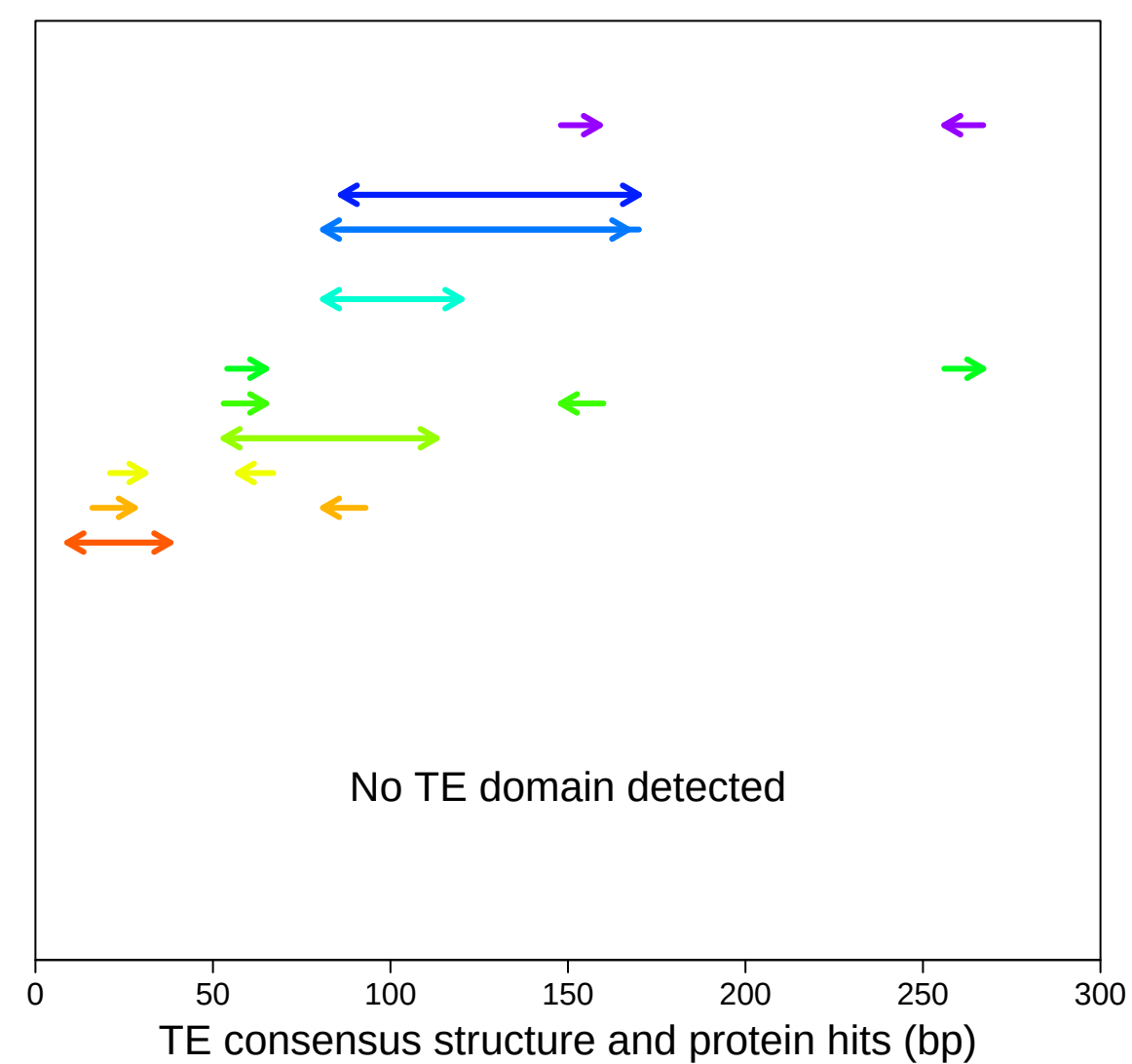
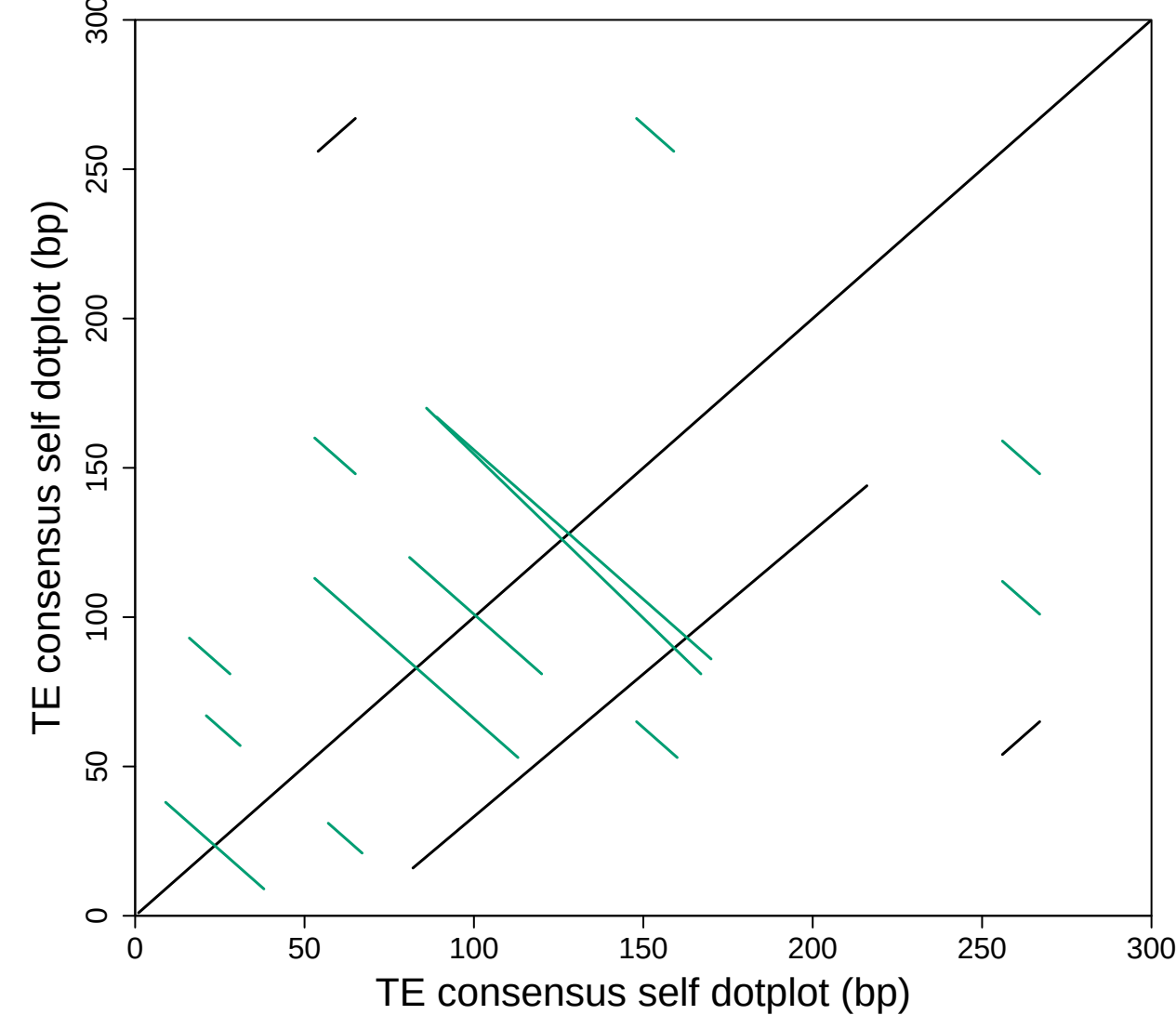
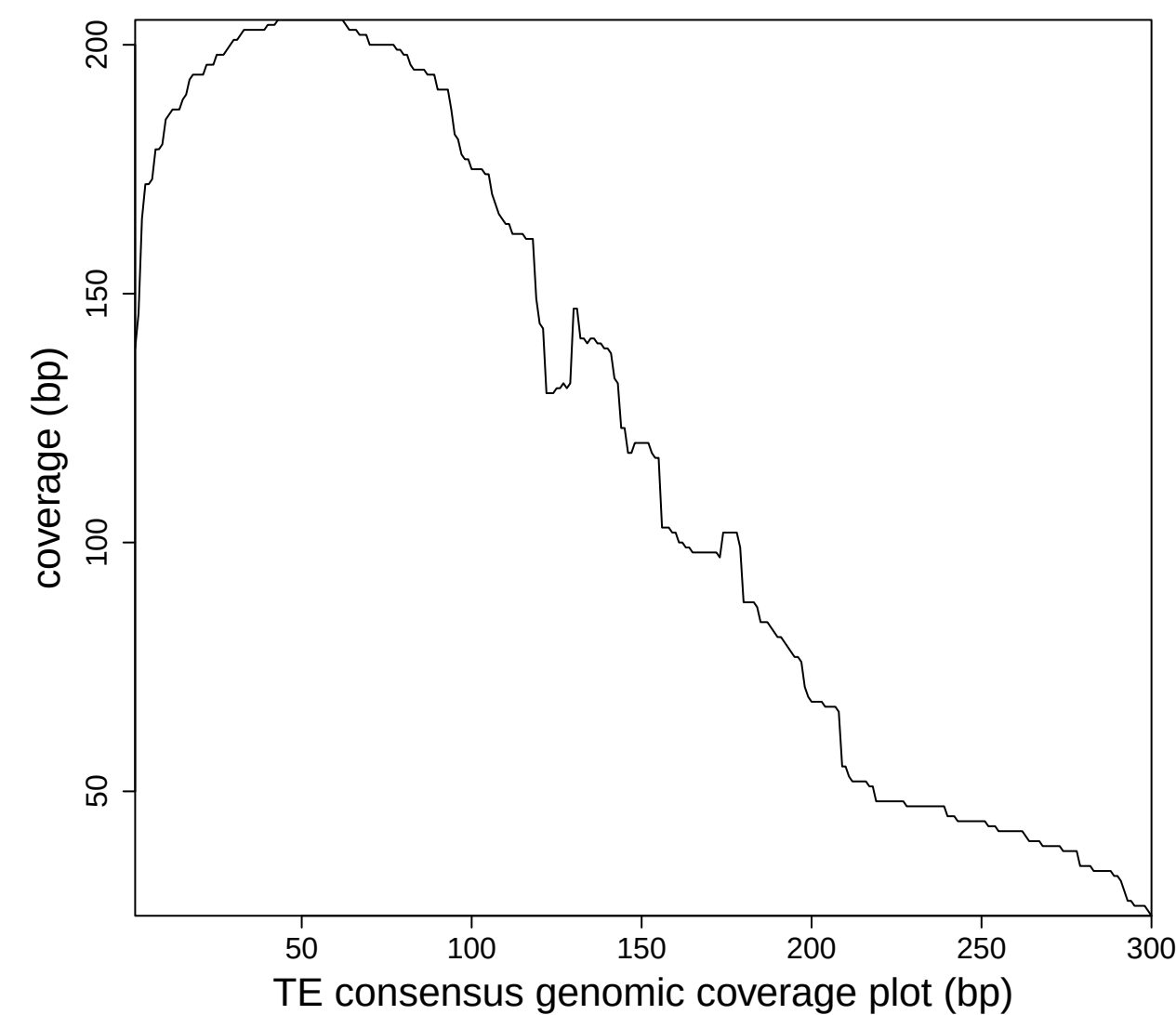
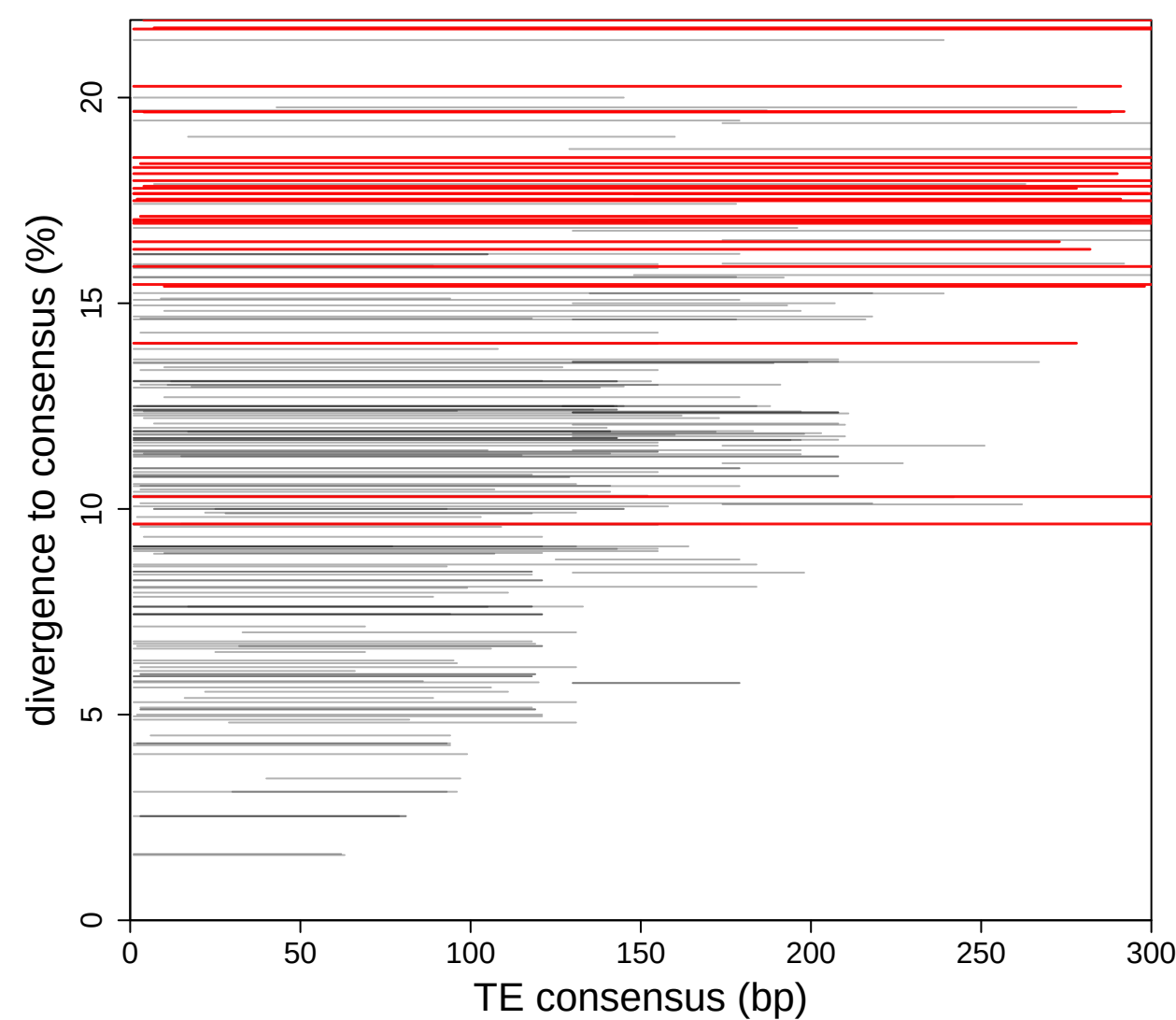
After TEtrimmer Extended plot Blue lines are boundaries



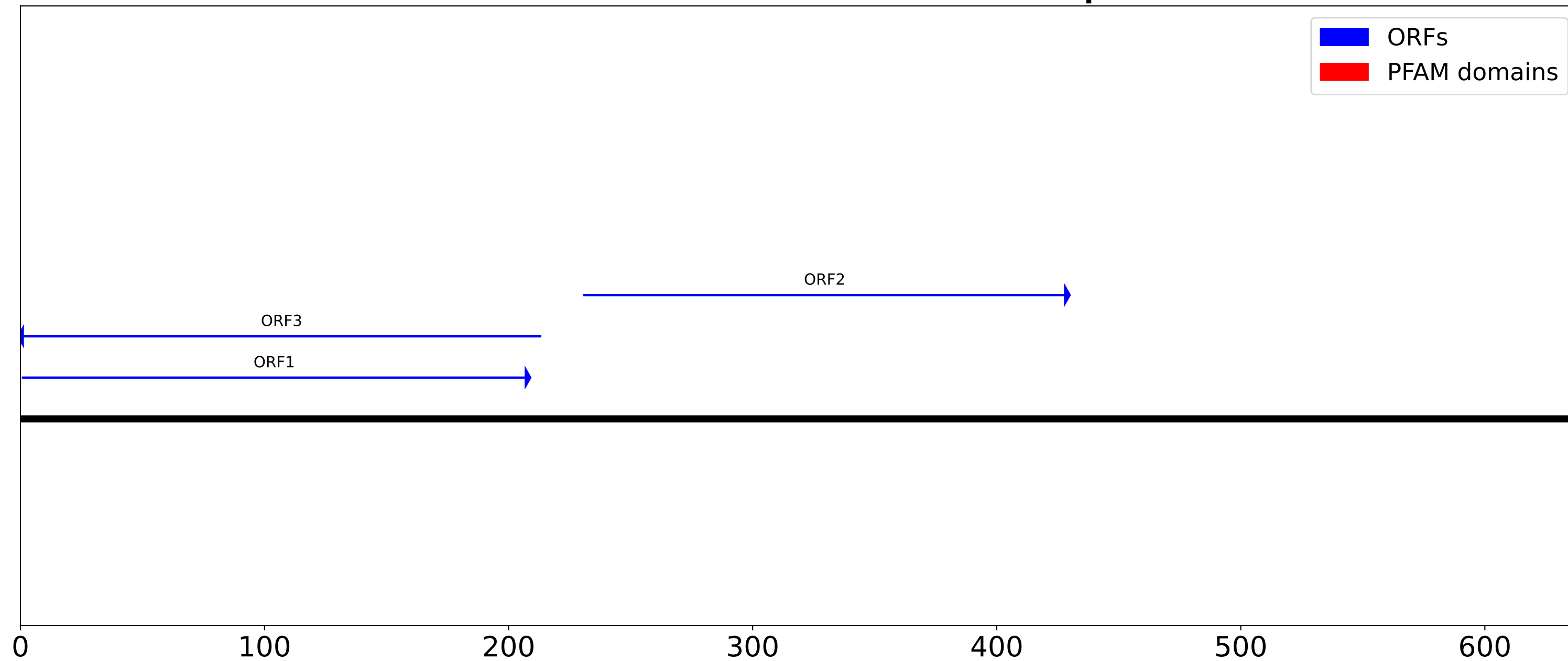
Before TEtrimmer 300 bp

TE: rnd_4_family_365

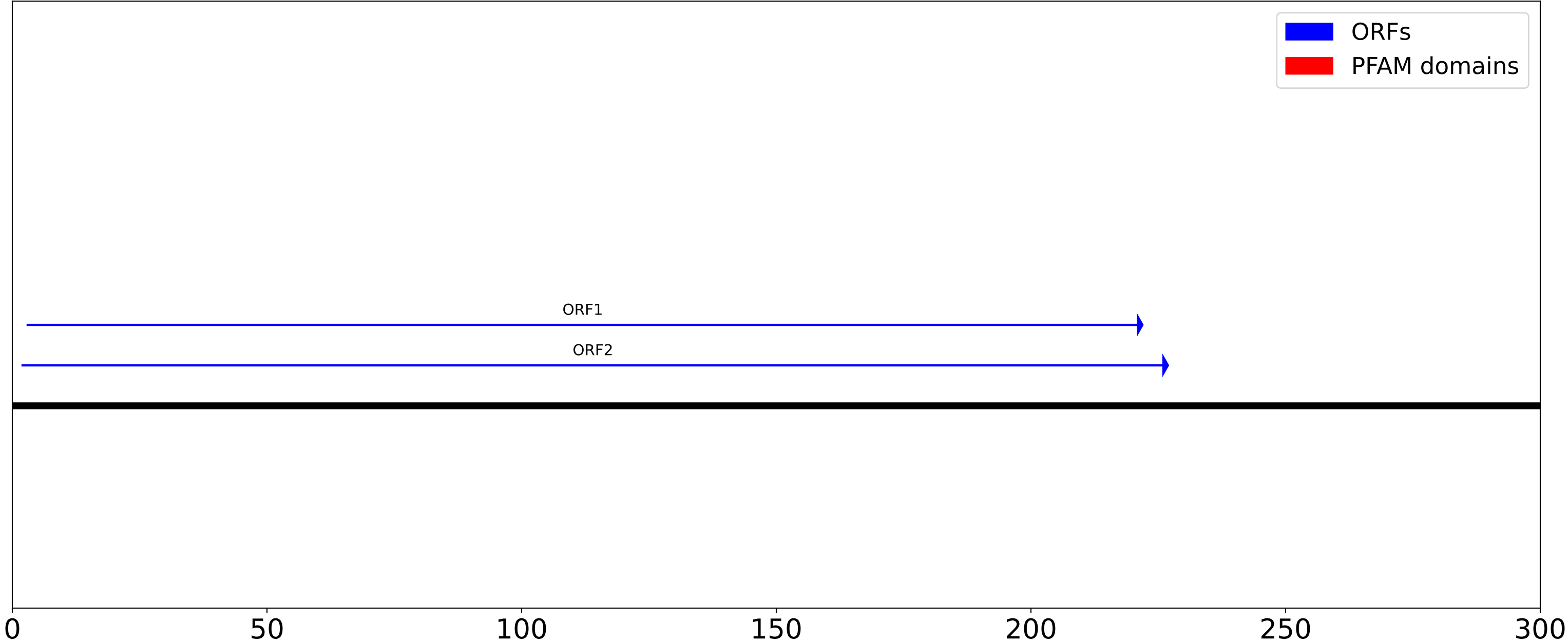
size: 300bp; fragments: 234; full length: 29 (≥ 270 bp)



After TEtrimmer ORF and PFAM domain plot



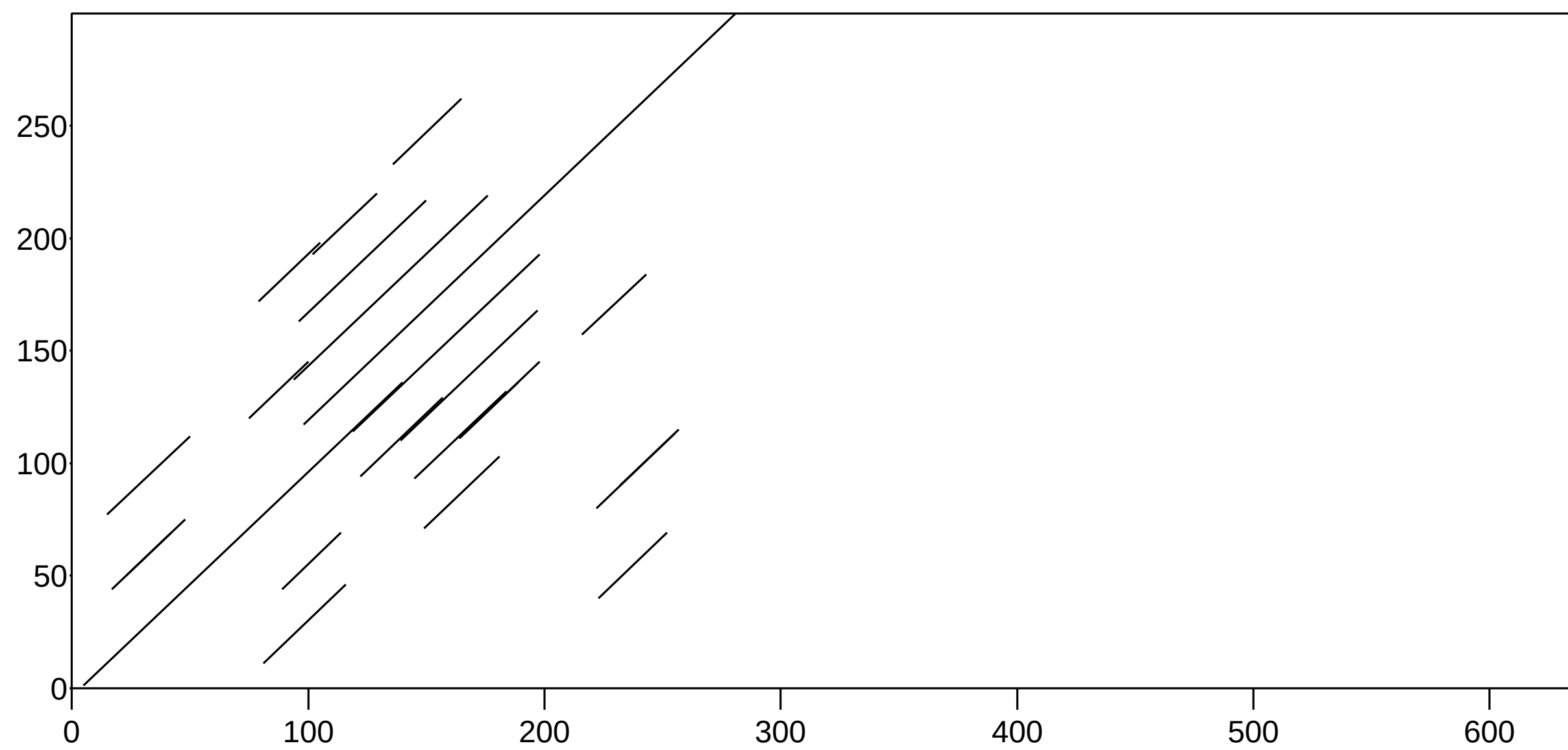
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)