

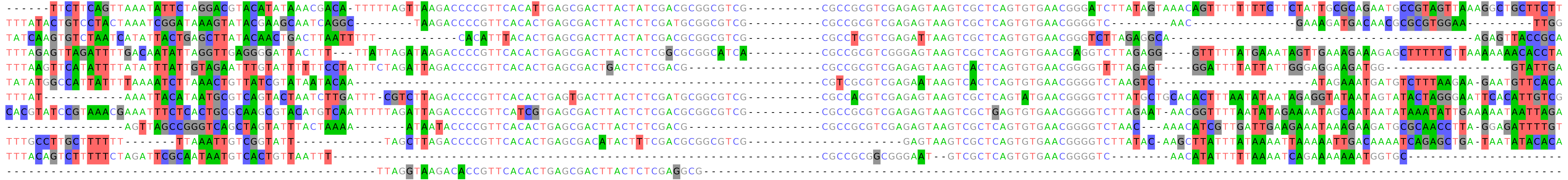
Start crop Point

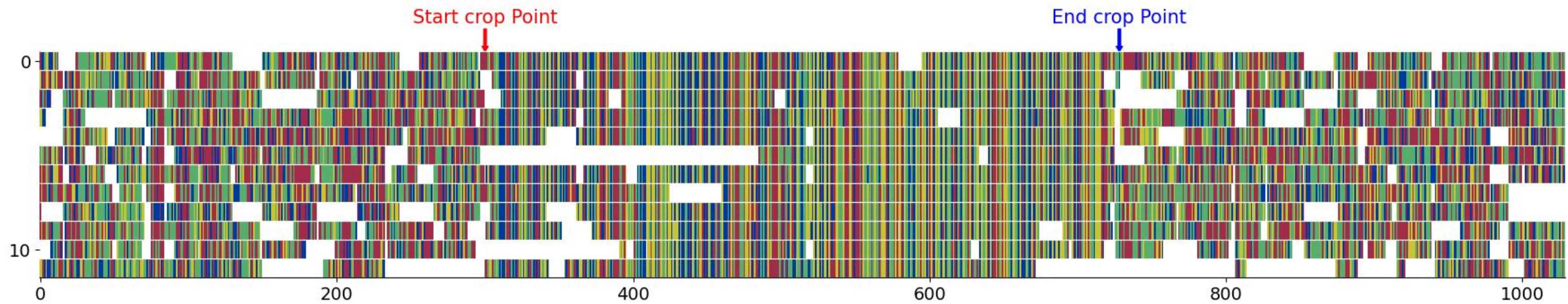


End crop Point



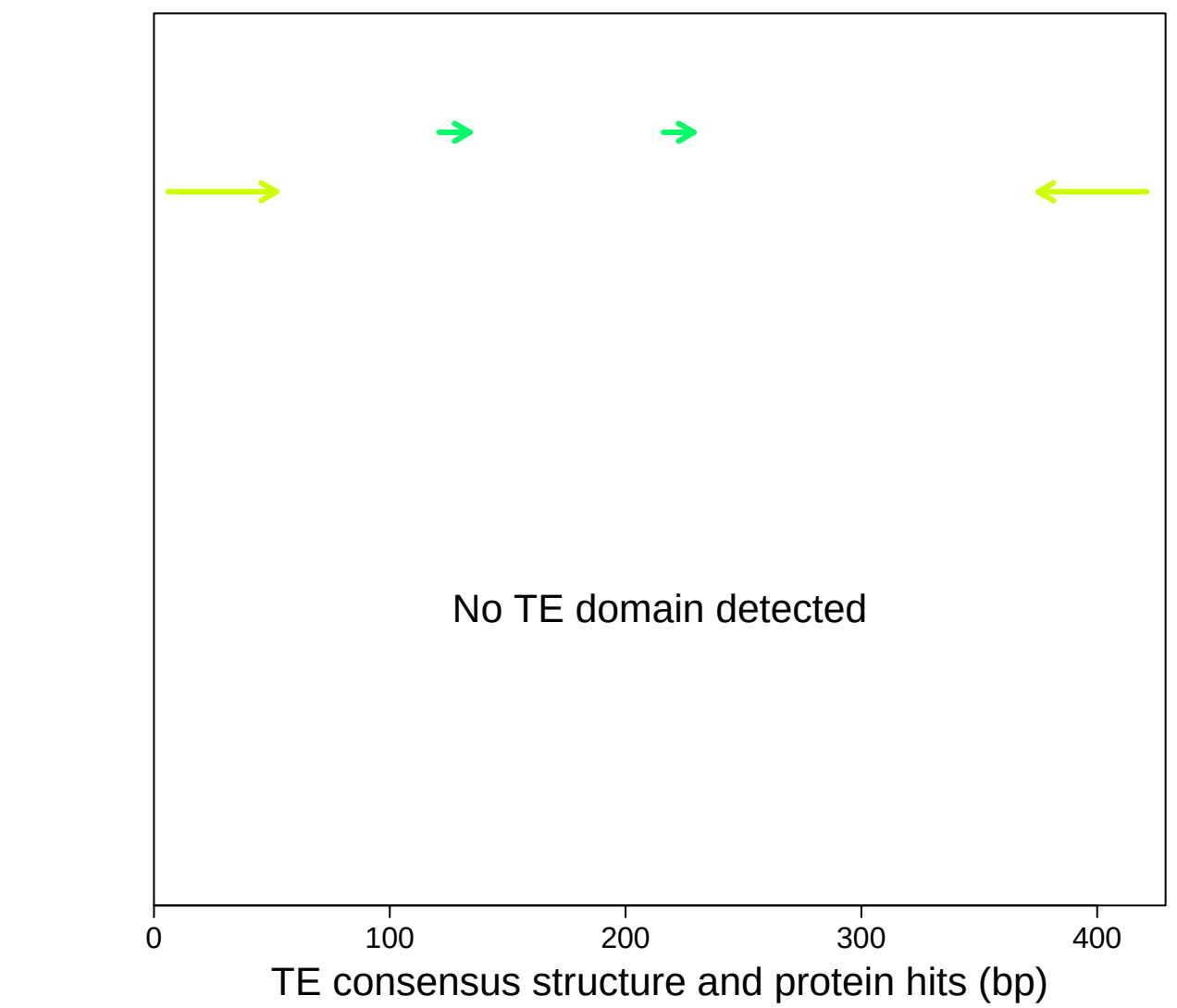
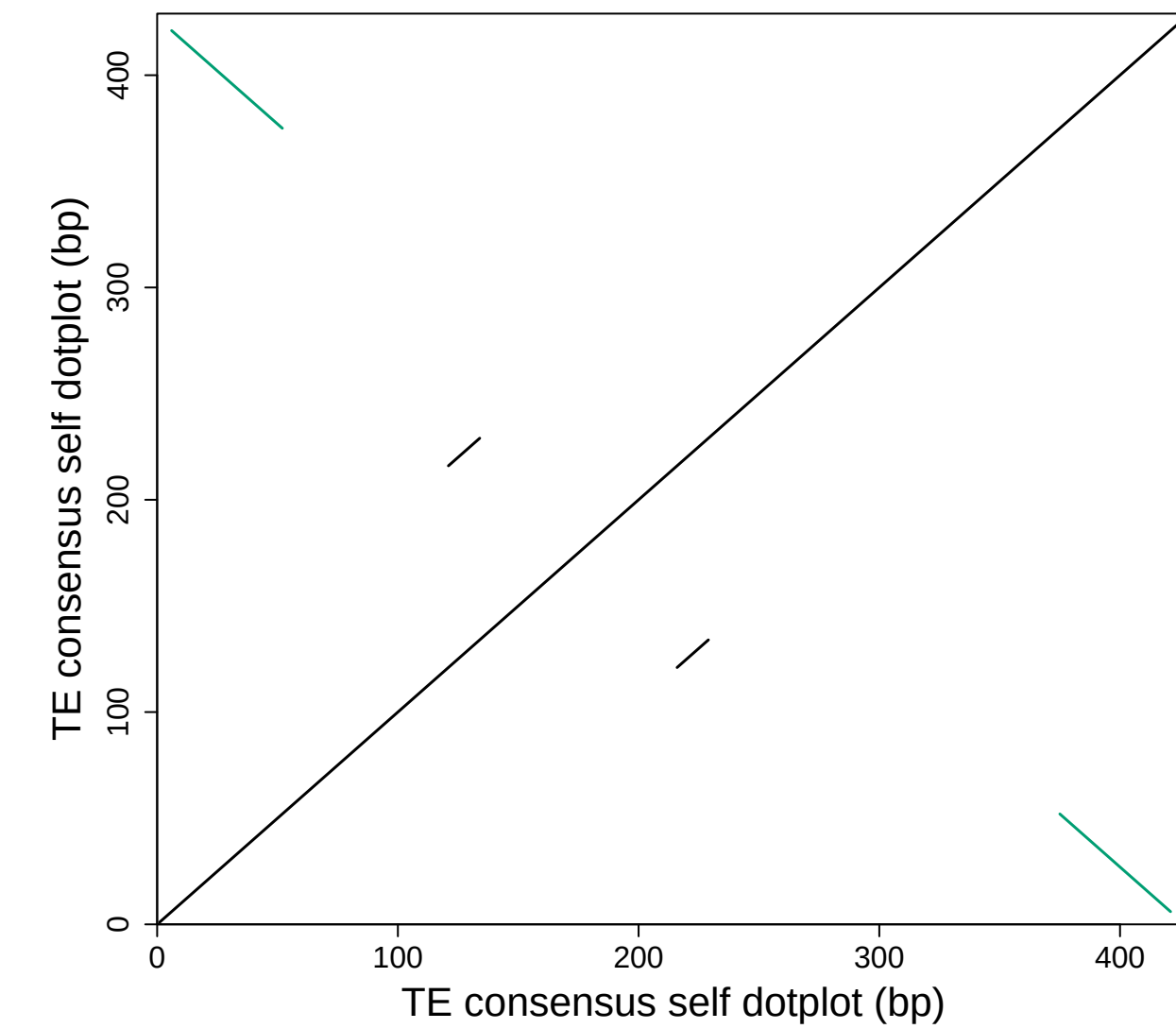
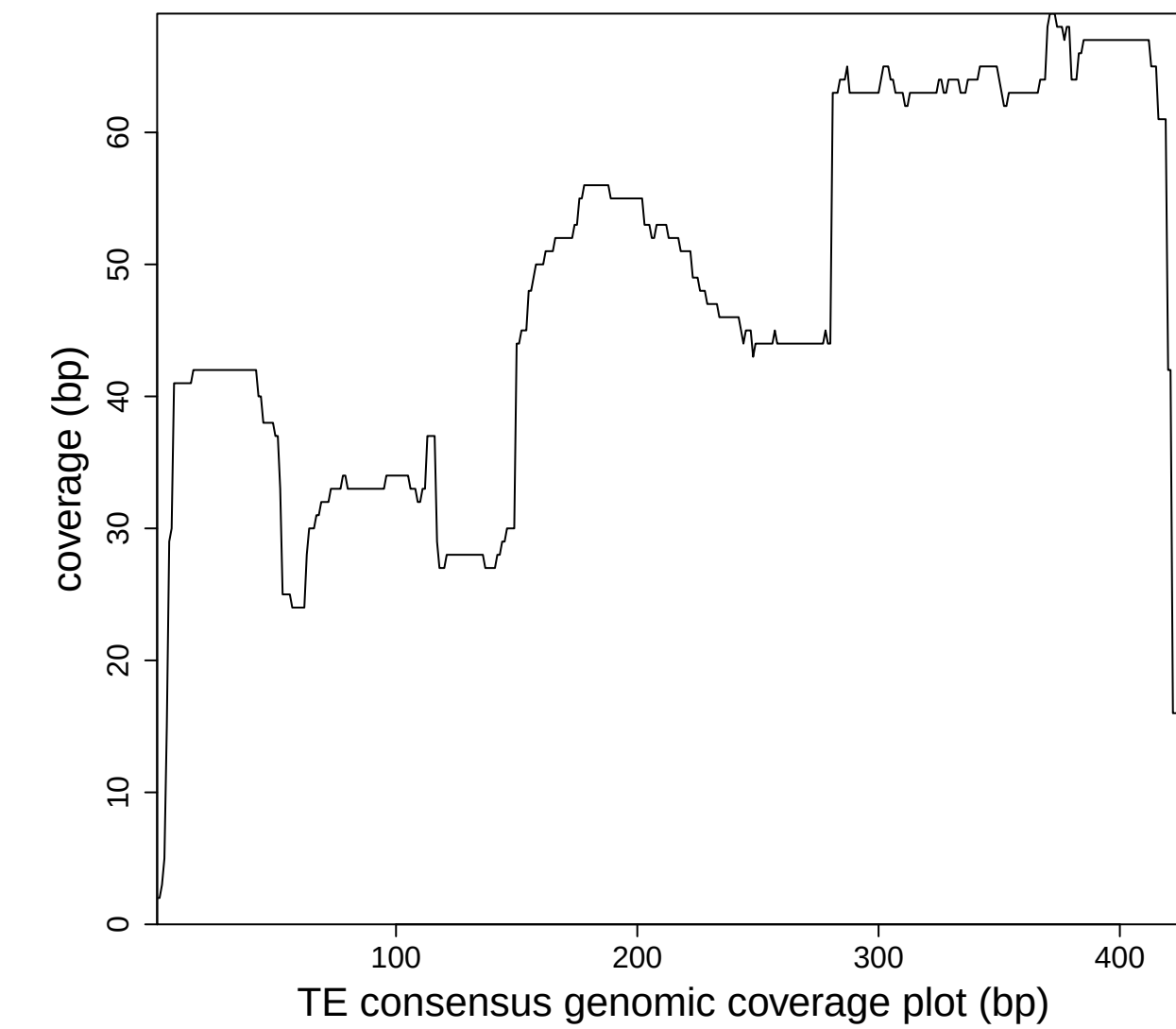
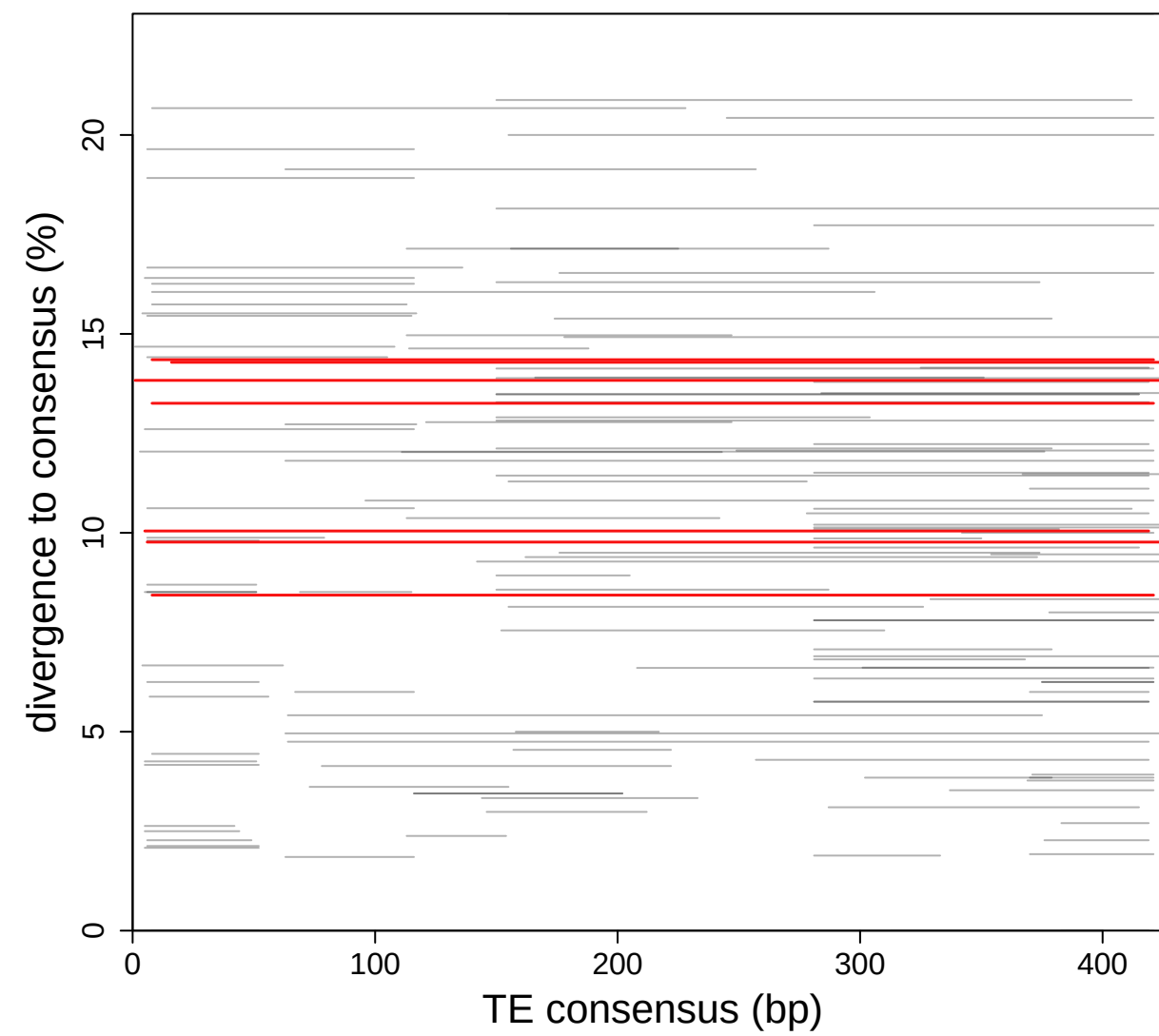
MSA length = 429





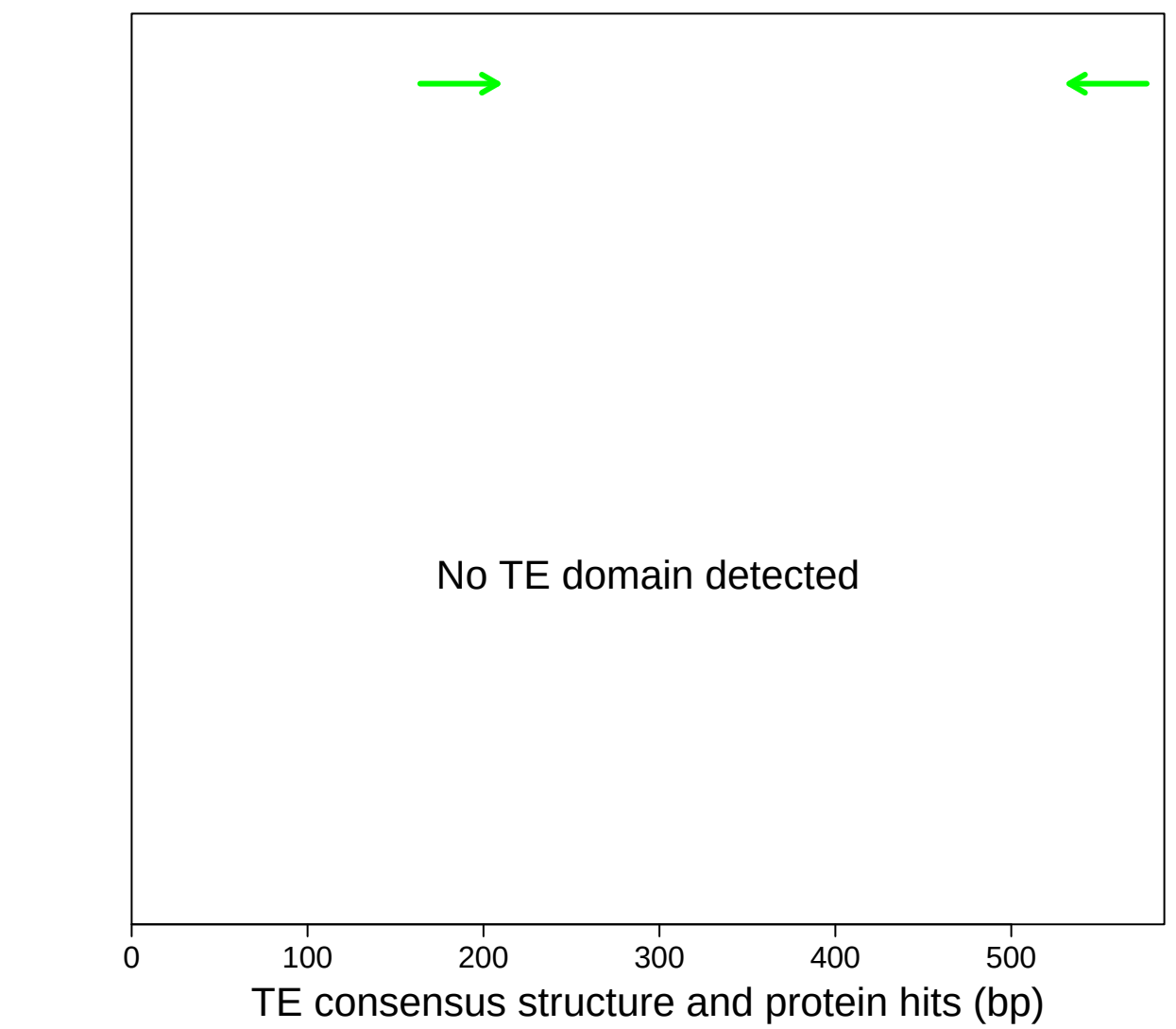
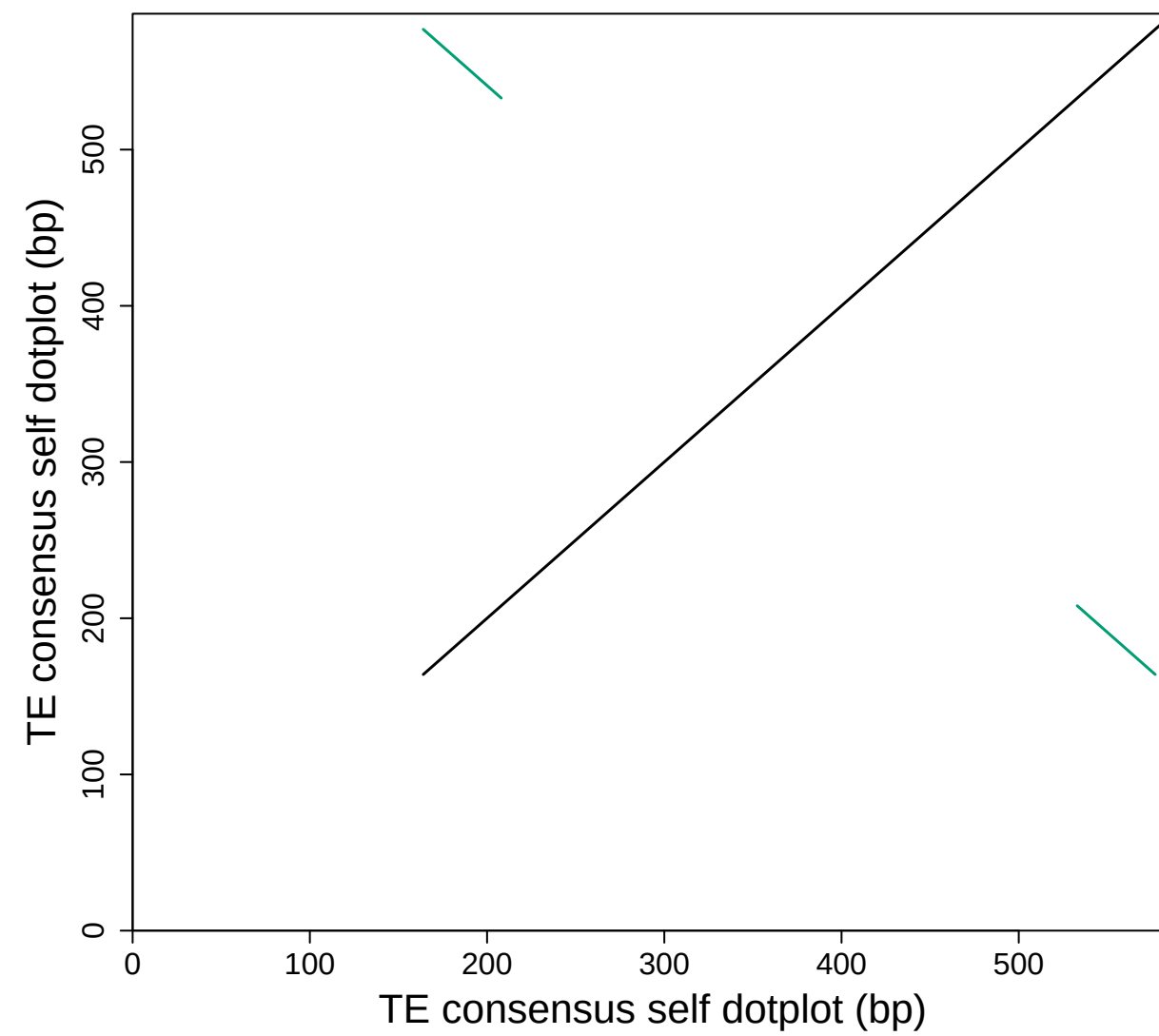
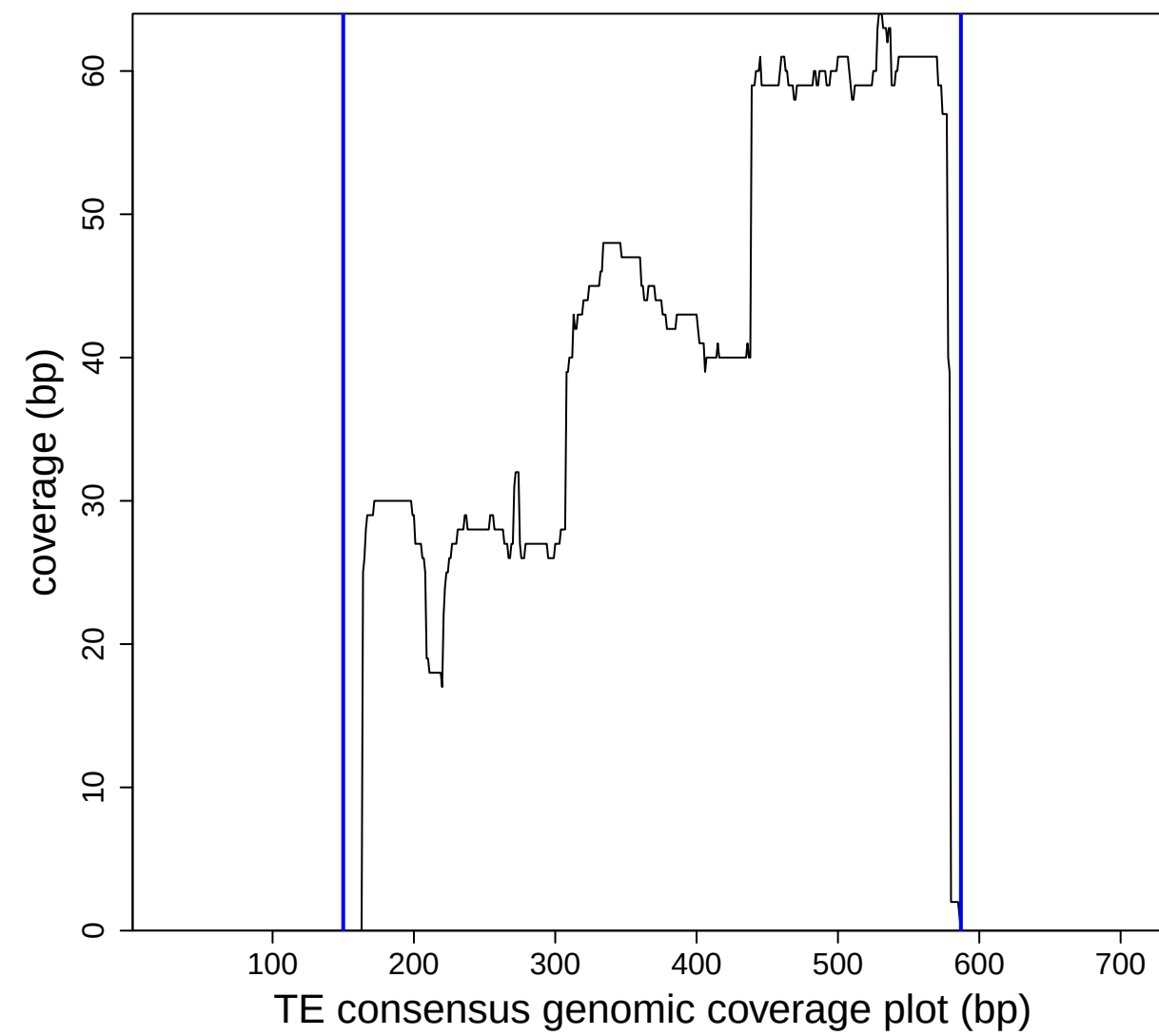
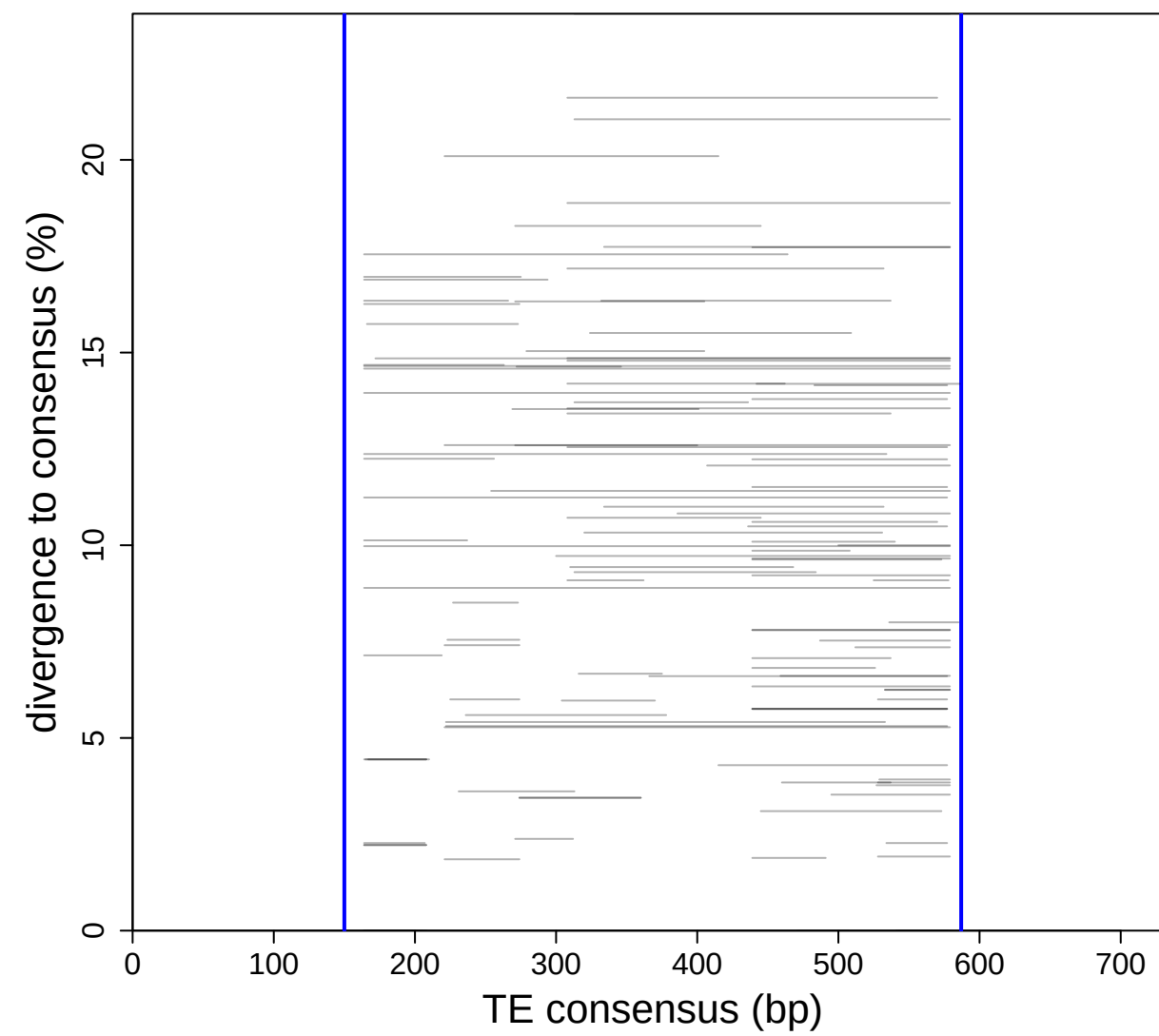
a.b.bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.f
size: 429bp; fragments: 143; full length: 7 (>=386.1bp)

After TEtrimmer 429 bp

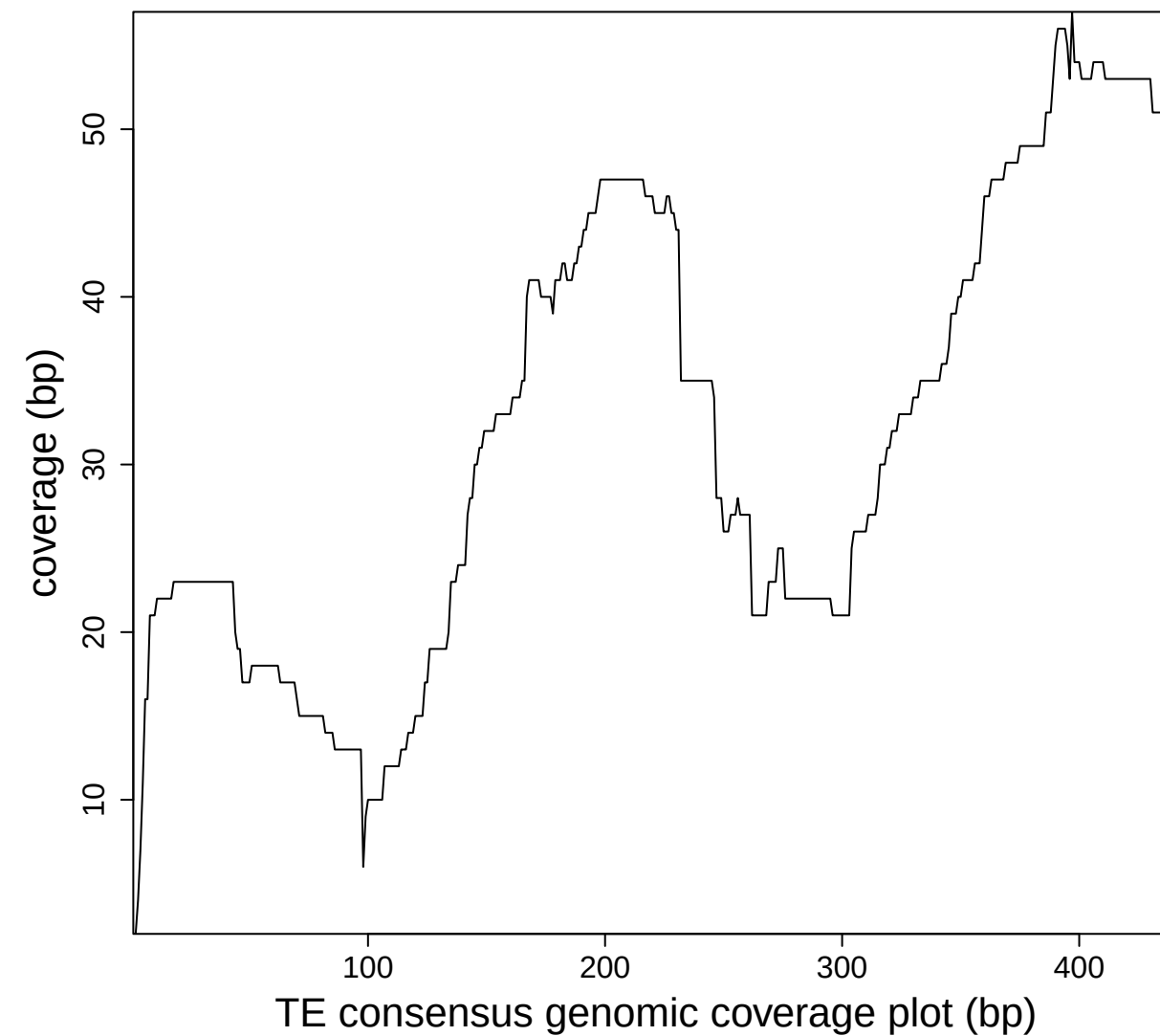
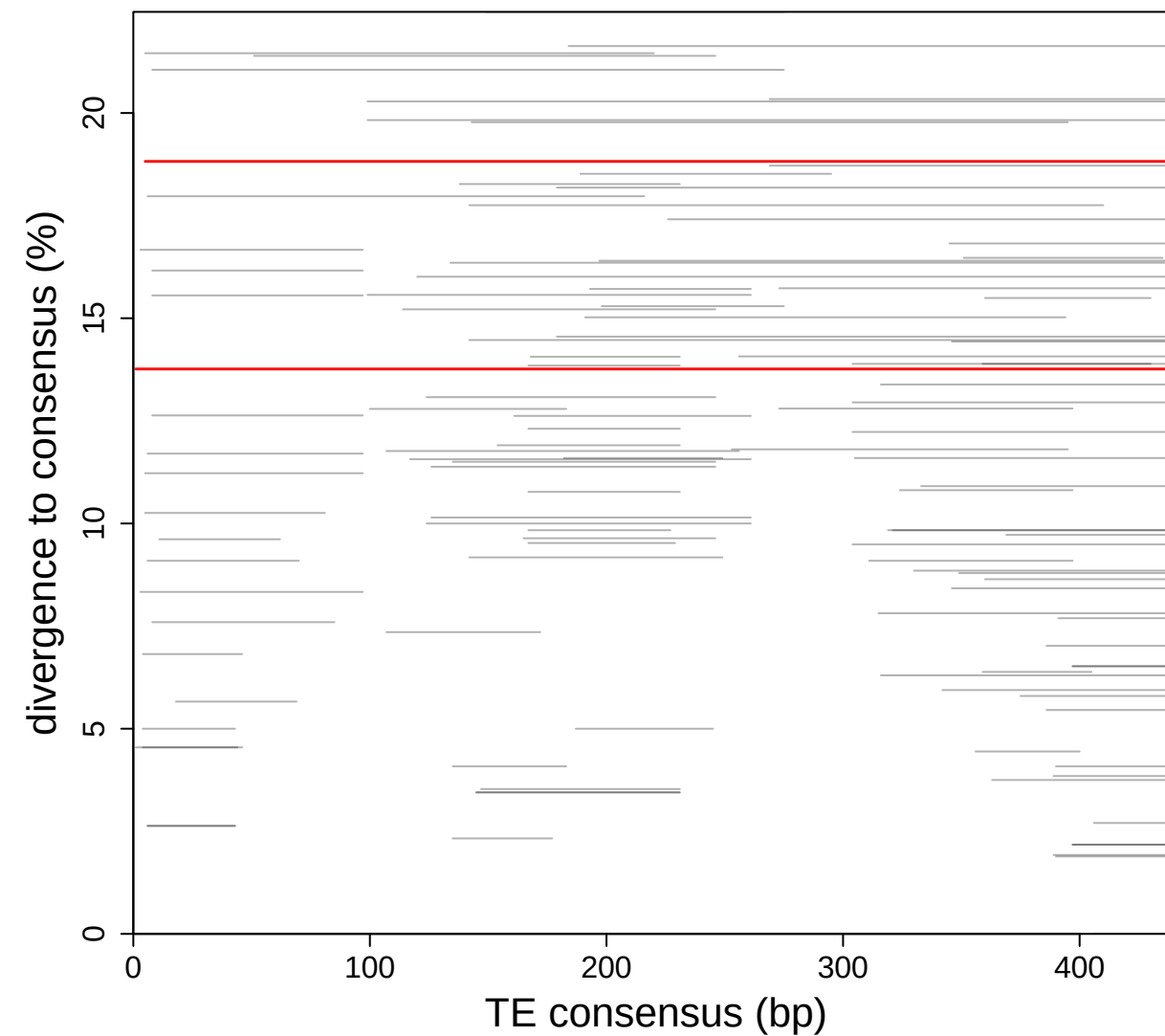


6.fasta.b.bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcIn.fa_aln.fa_cl.f
size: 737bp; fragments: 123; full length: 0 (>=663.3bp)

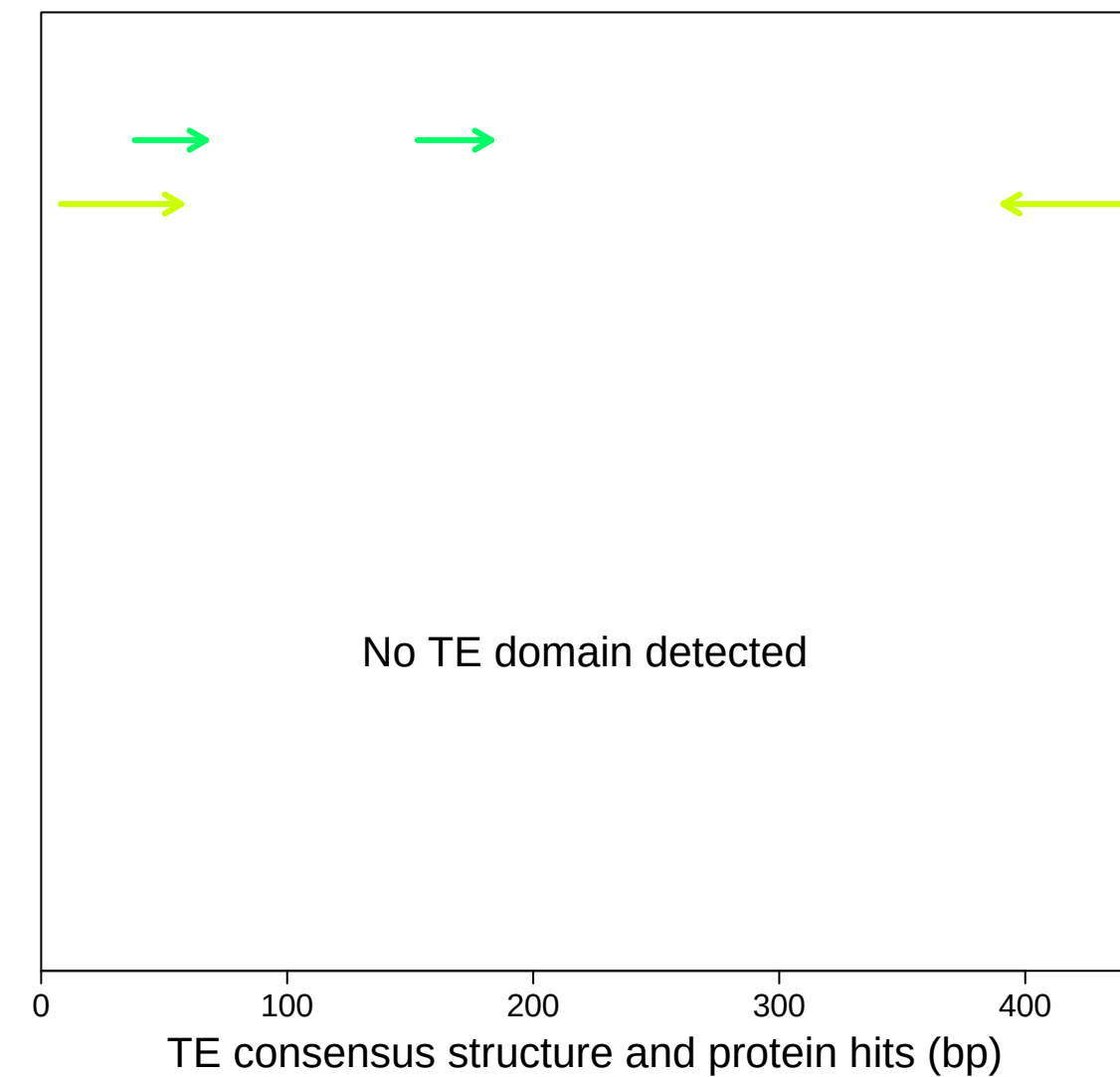
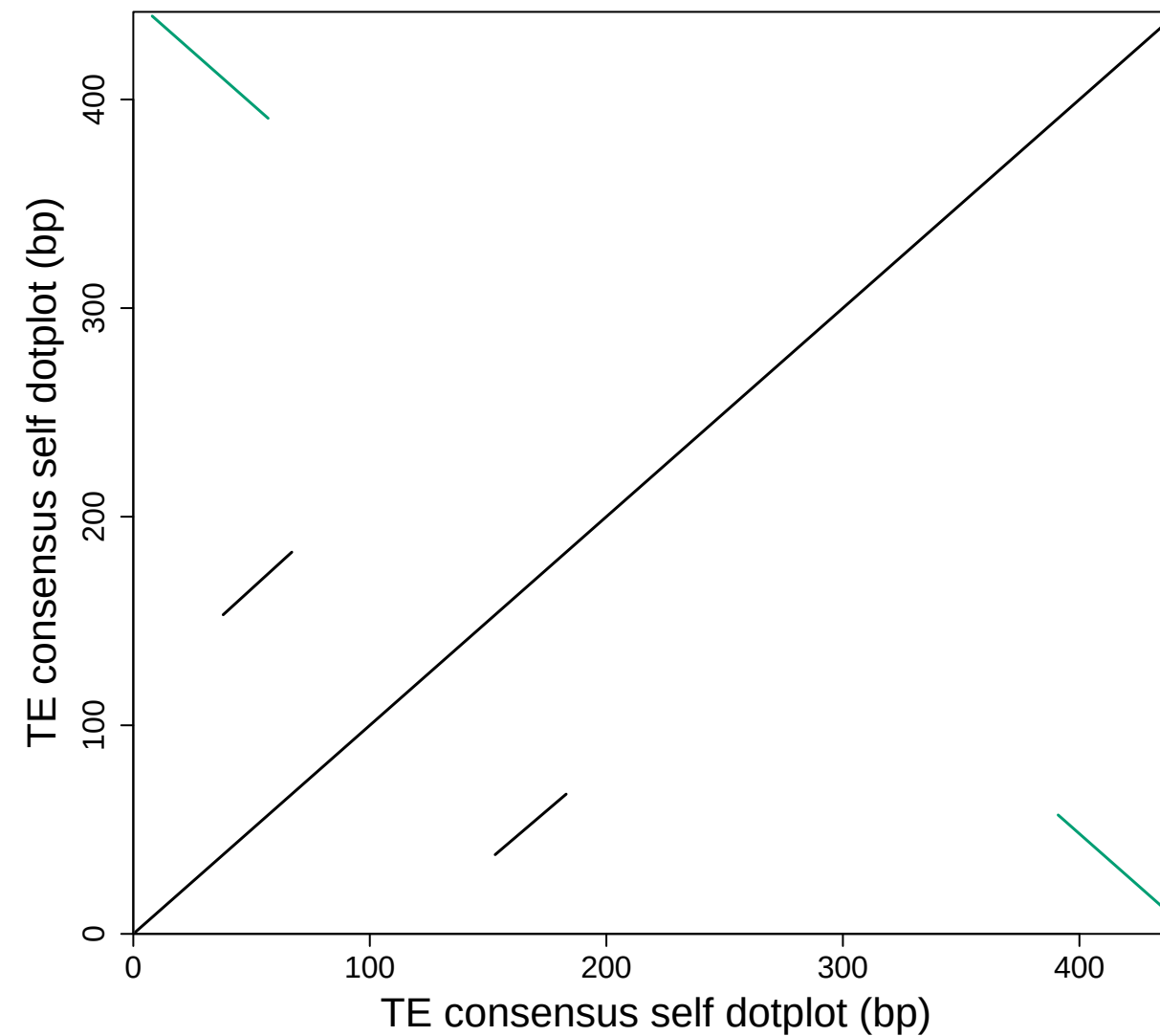
After TEtrimmer Extended plot Blue lines are boundaries



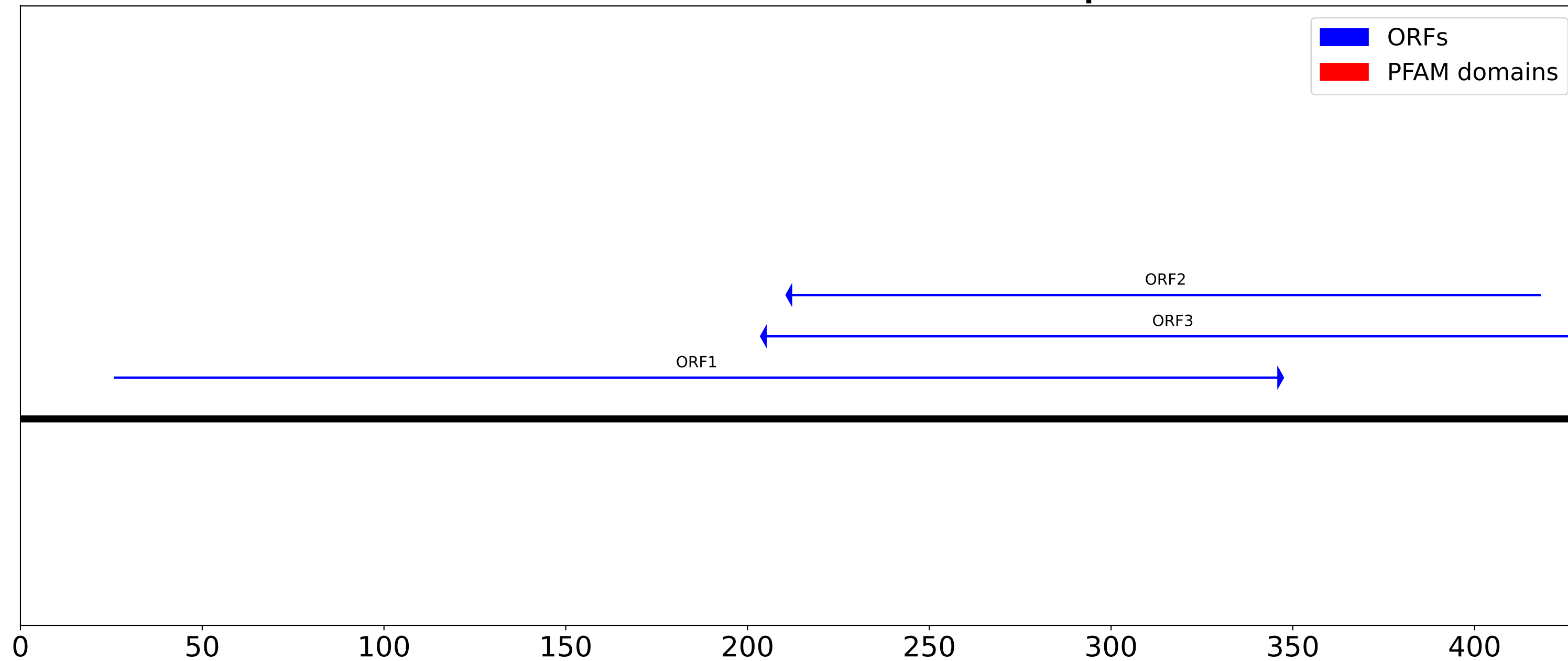
TE: rnd_4_family_4536
size: 442bp; fragments: 117; full length: 2 (≥ 397.8 bp)



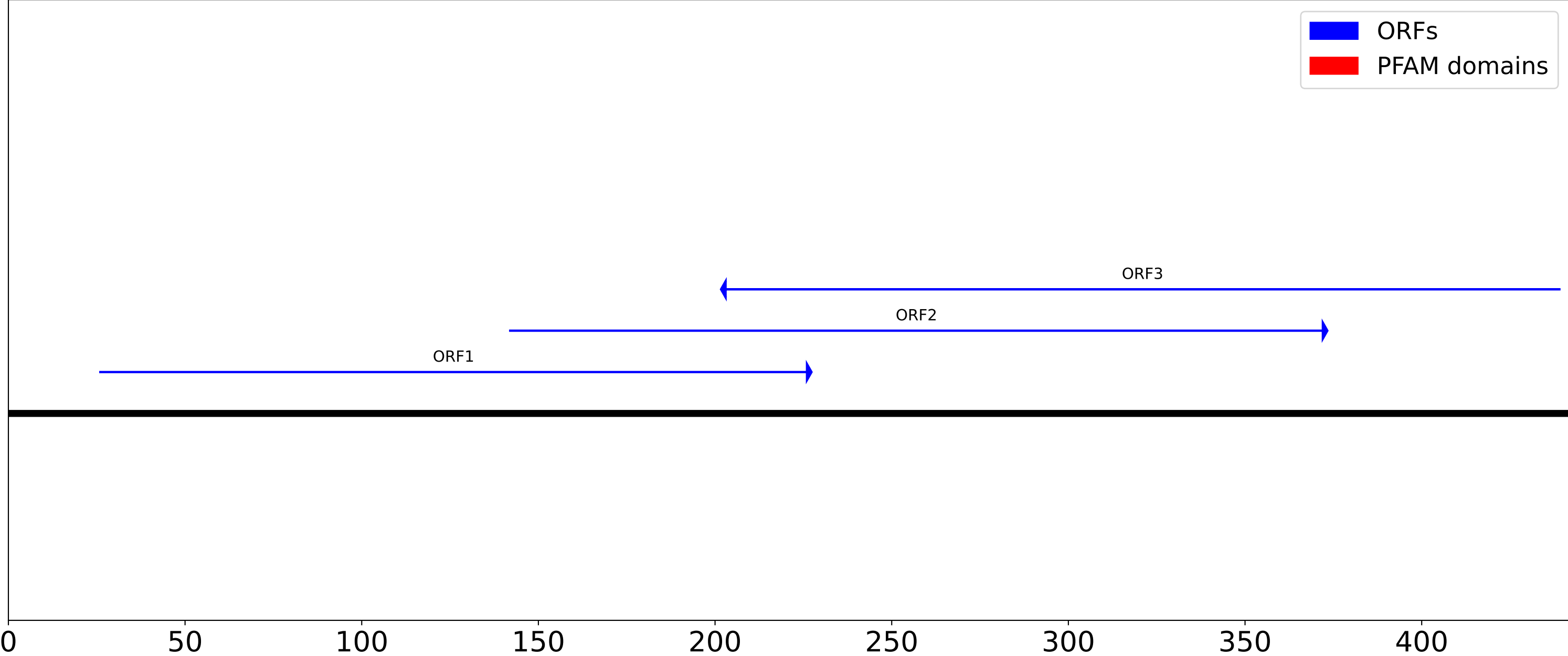
Before TEtrimmer 442 bp



After TEtrimmer ORF and PFAM domain plot



Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

