

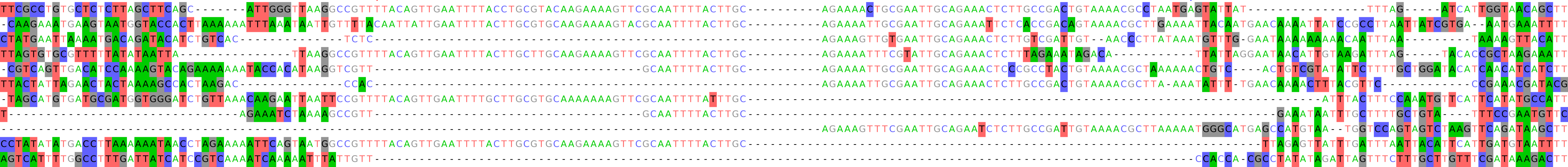
Start crop Point

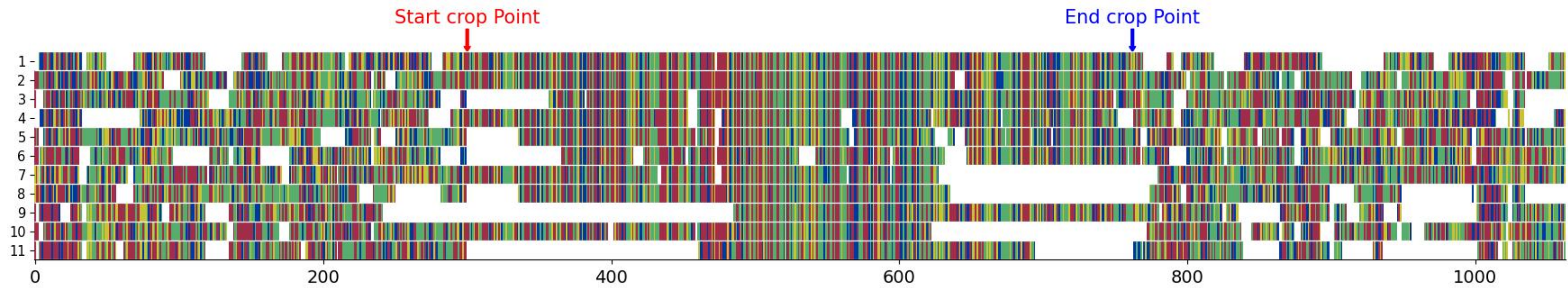


End crop Point



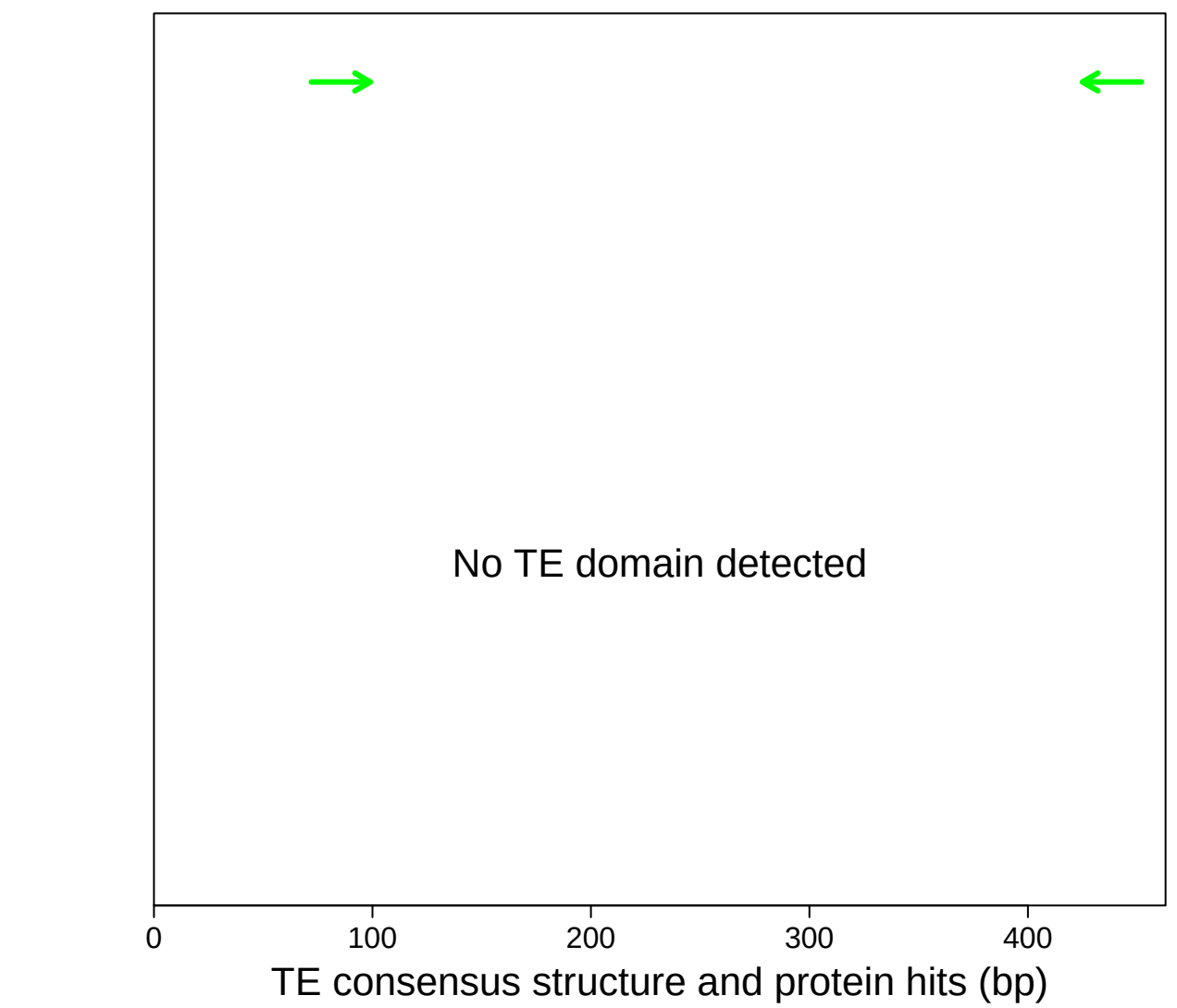
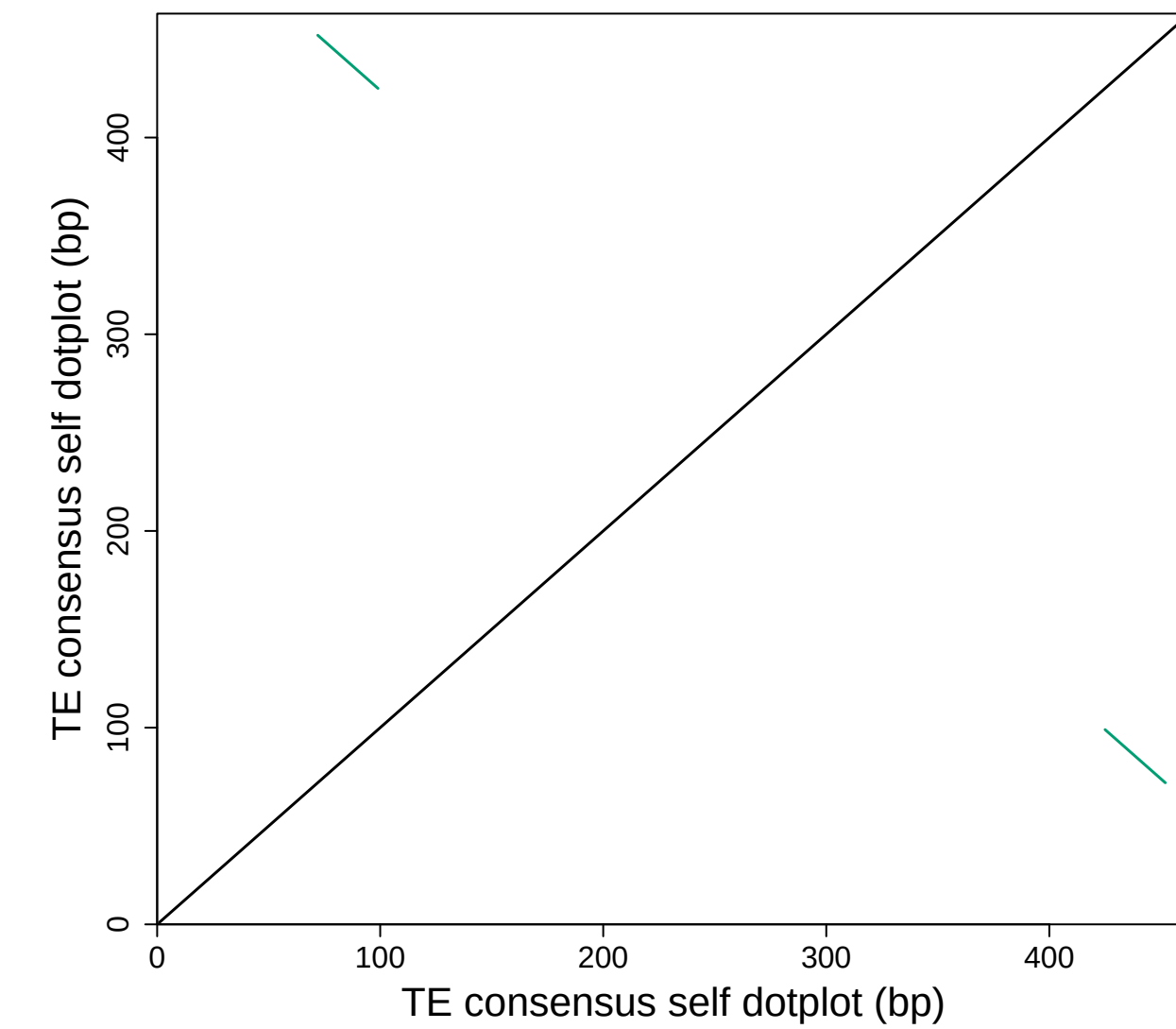
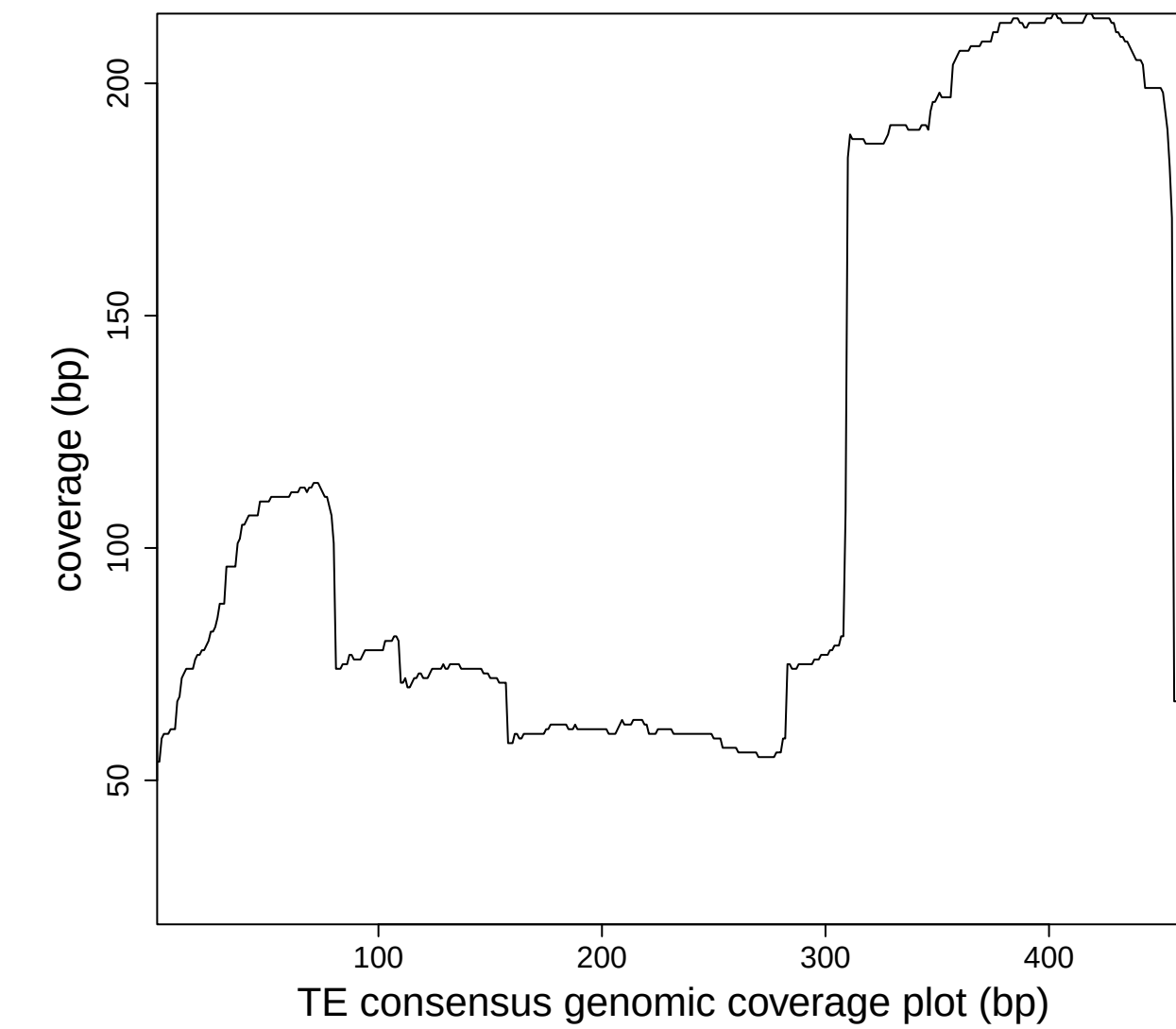
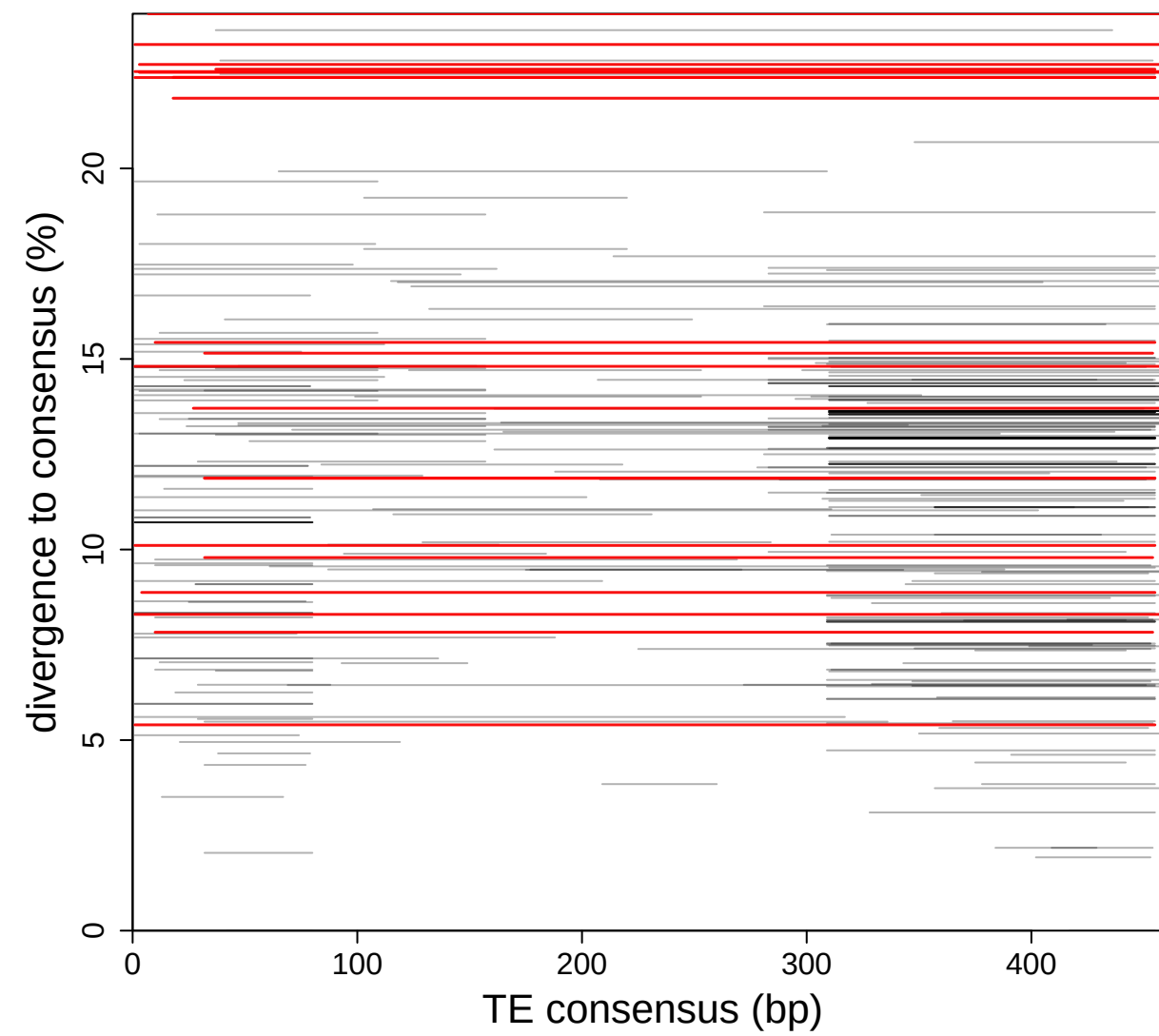
MSA length = 463





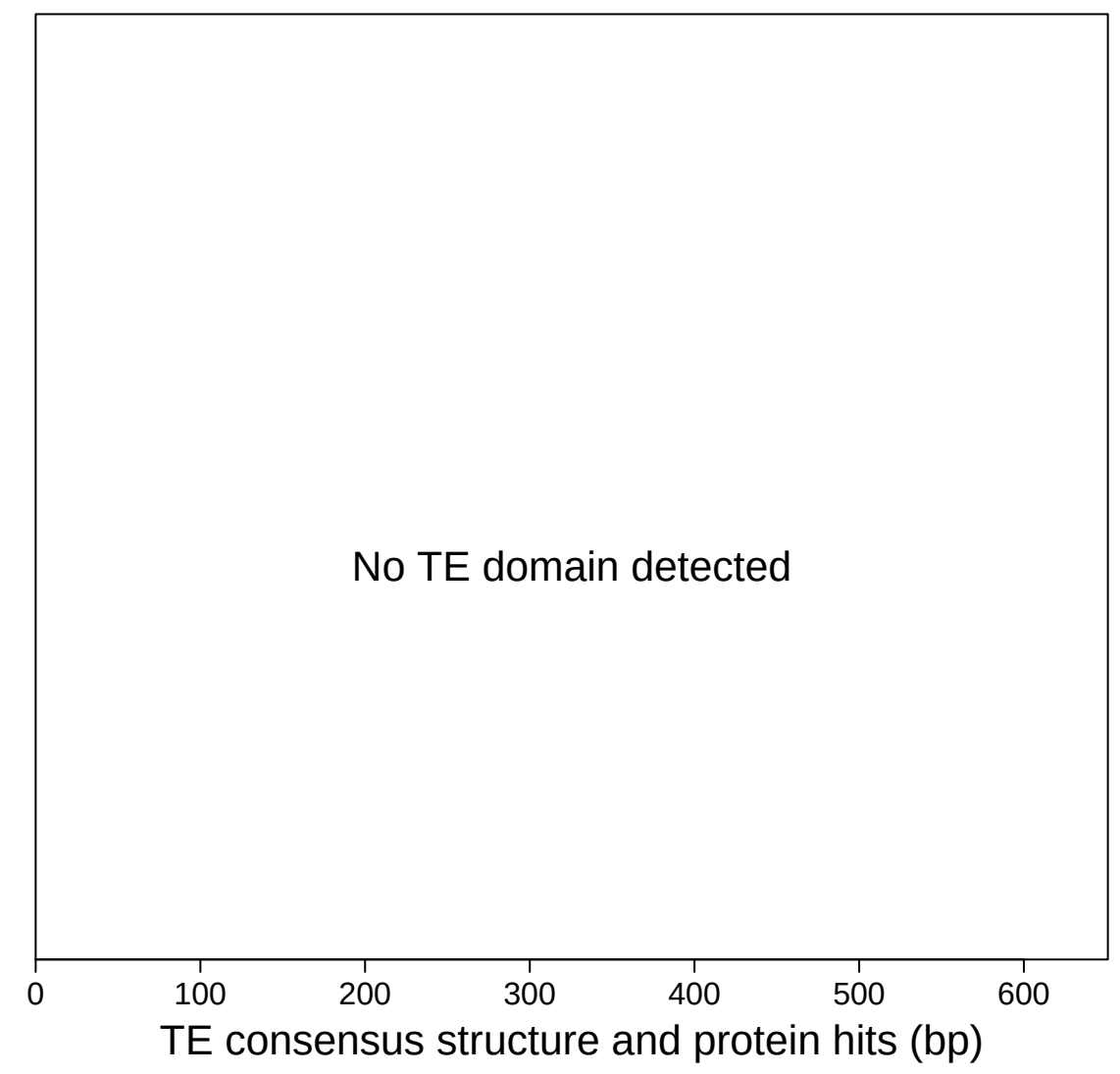
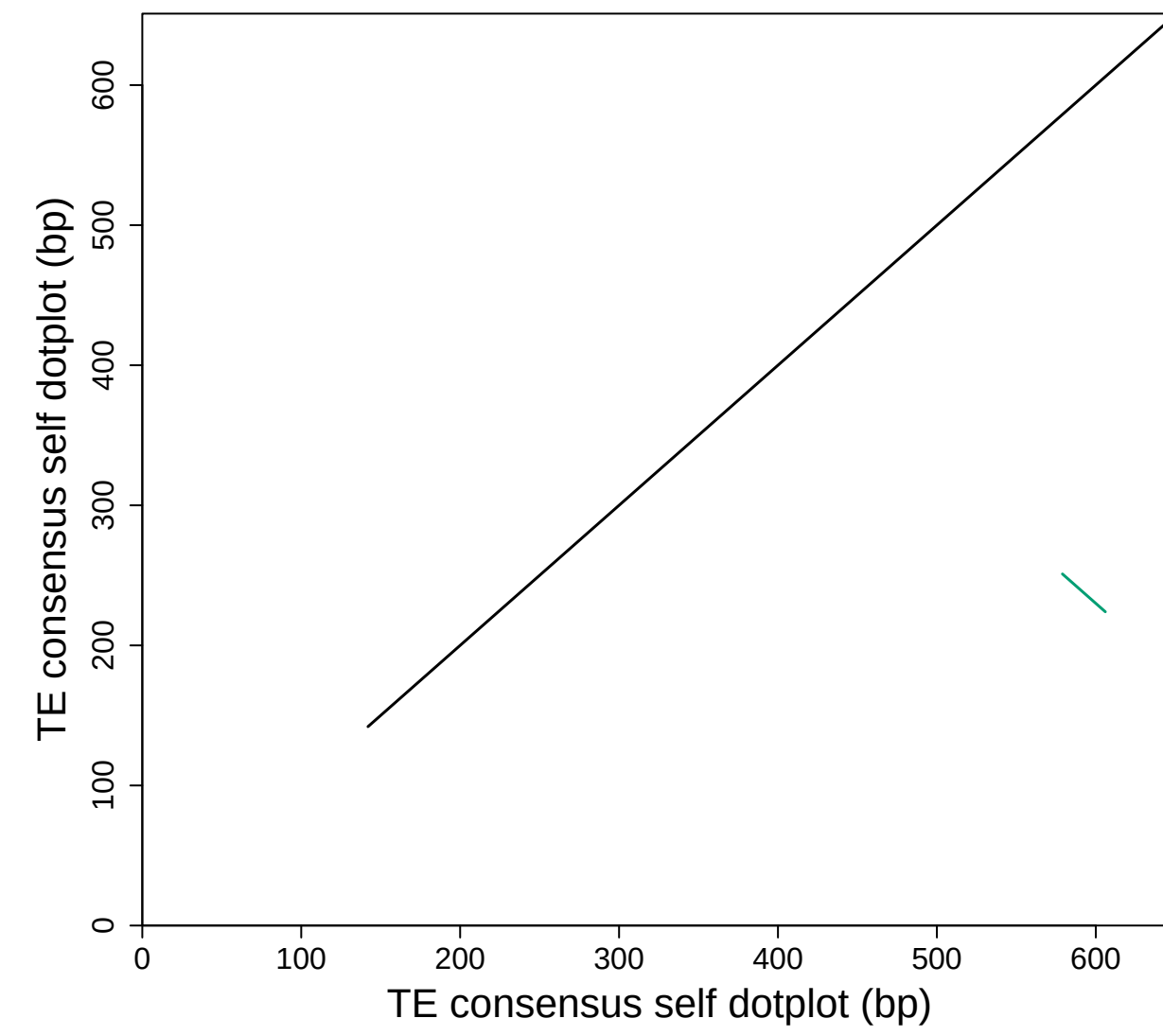
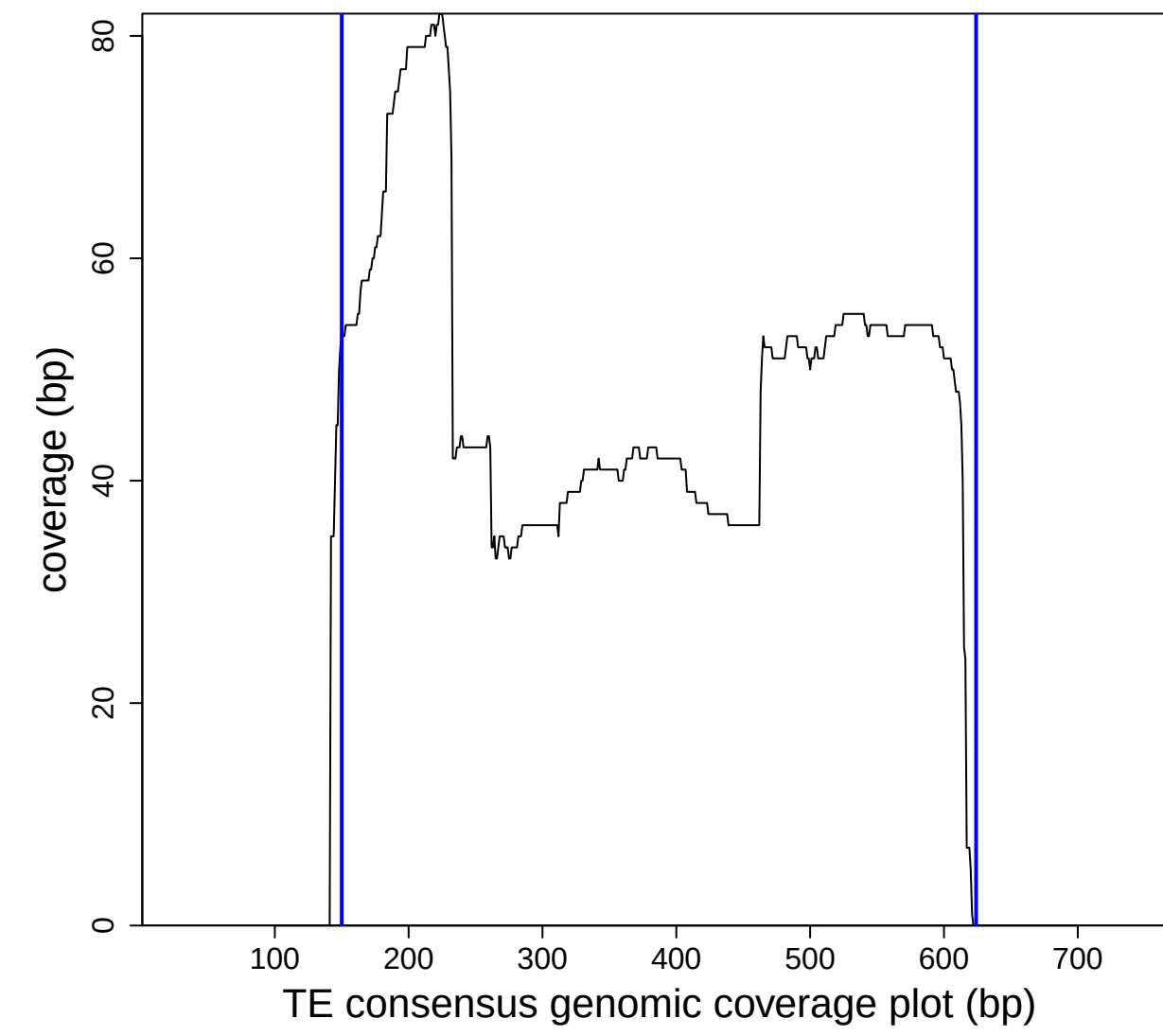
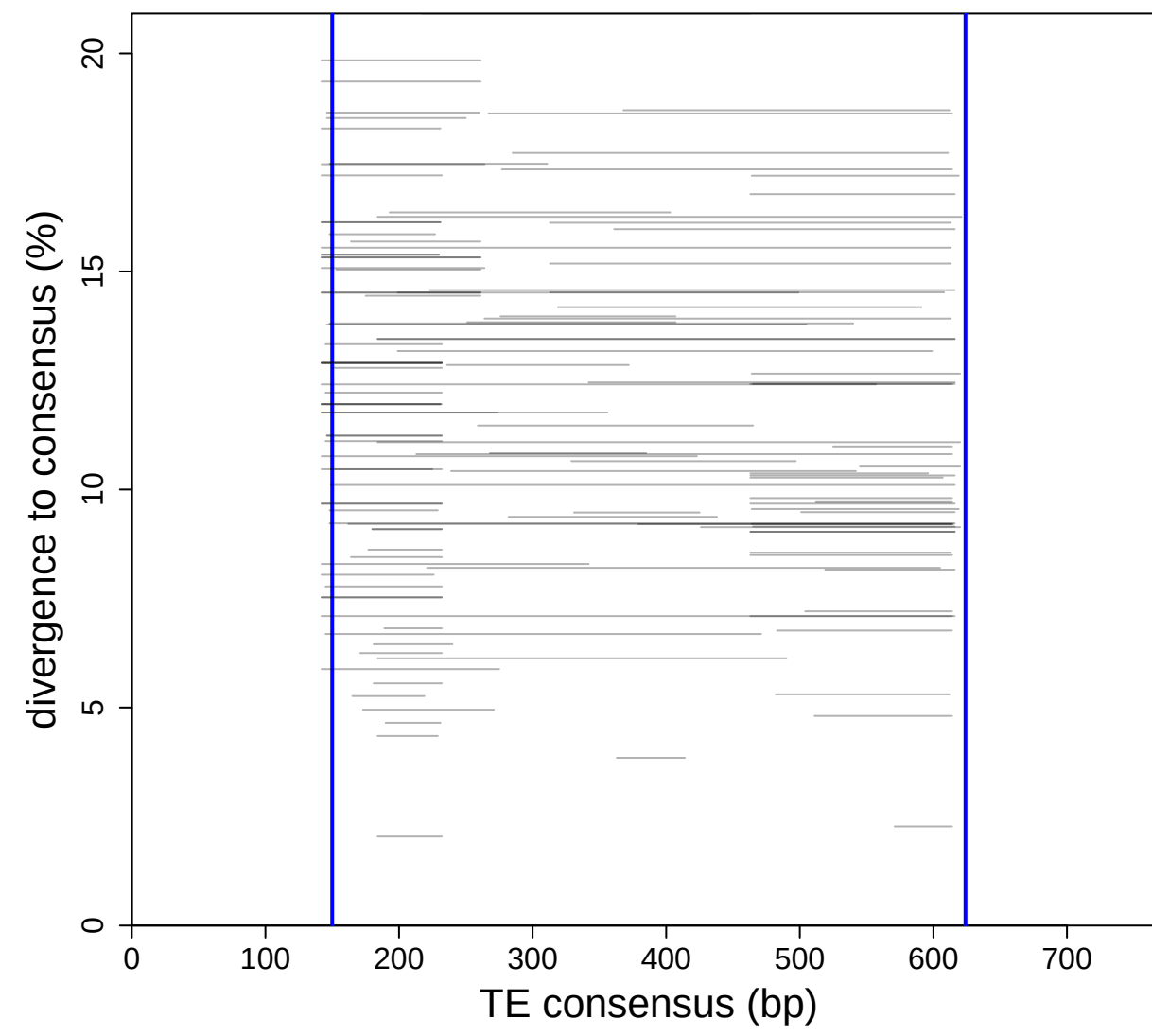
a.b.bed_uf.bed_g_2.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.f
size: 463bp; fragments: 318; full length: 21 (≥ 416.7 bp)

After TEtrimmer 463 bp

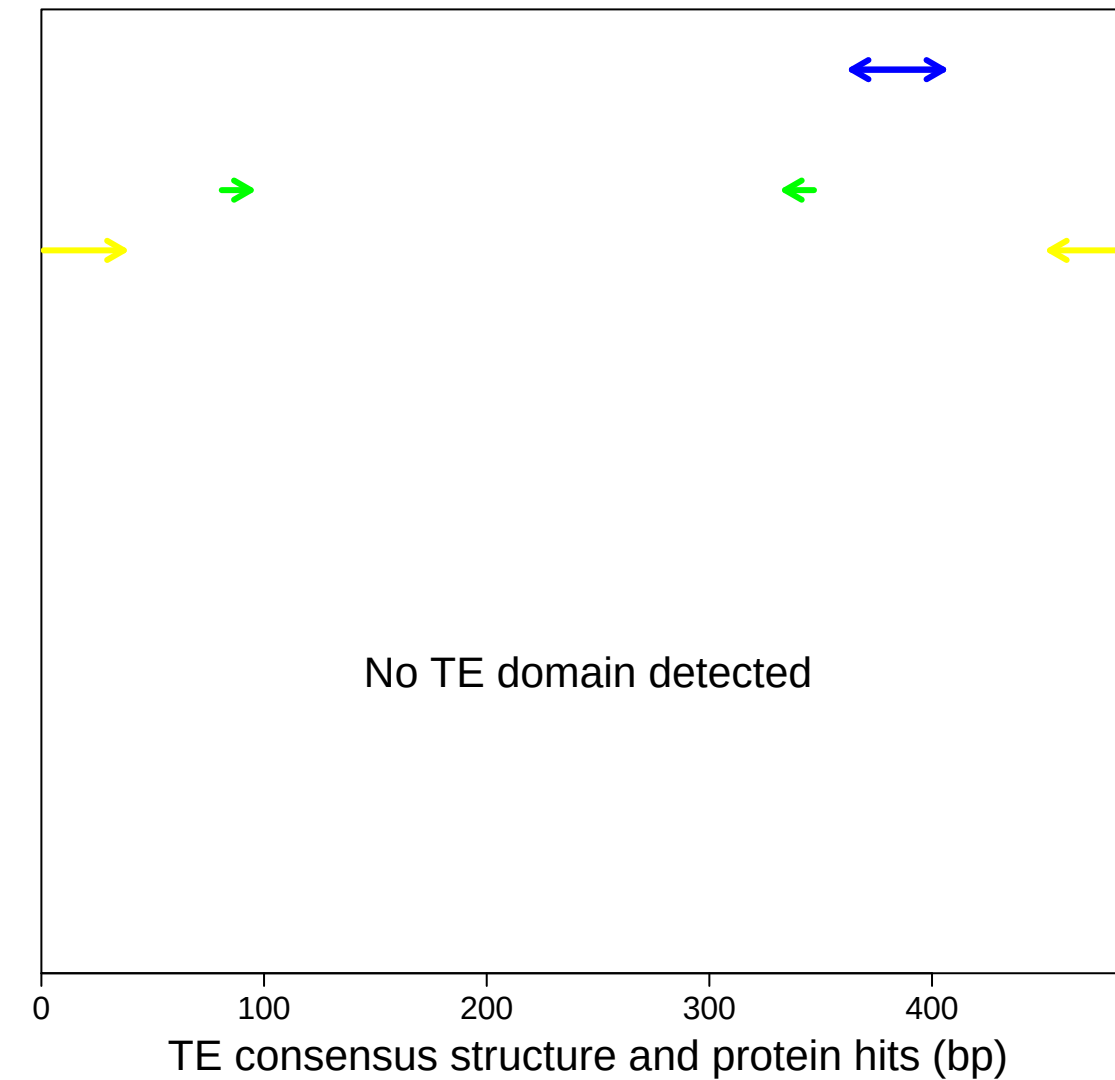
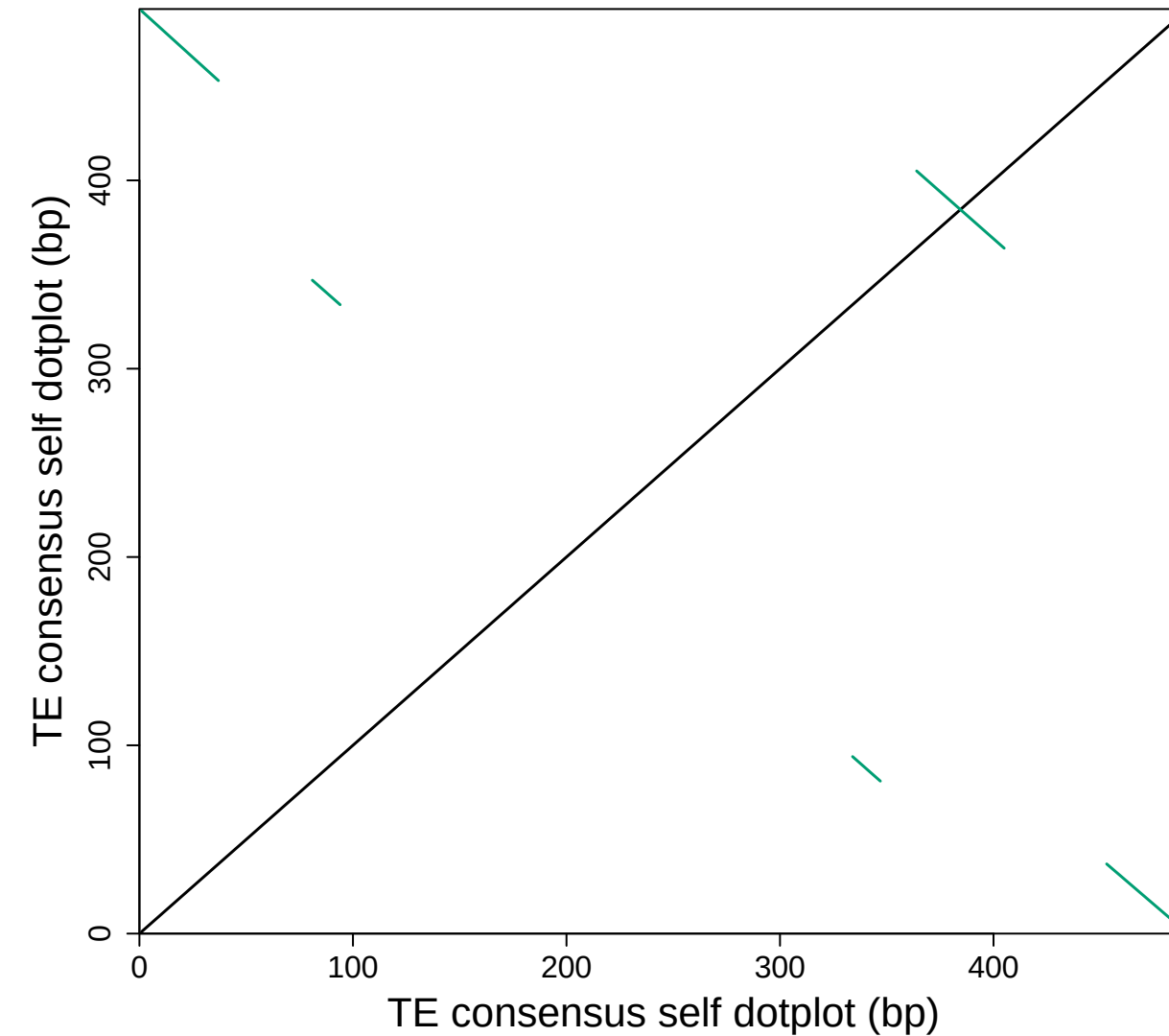
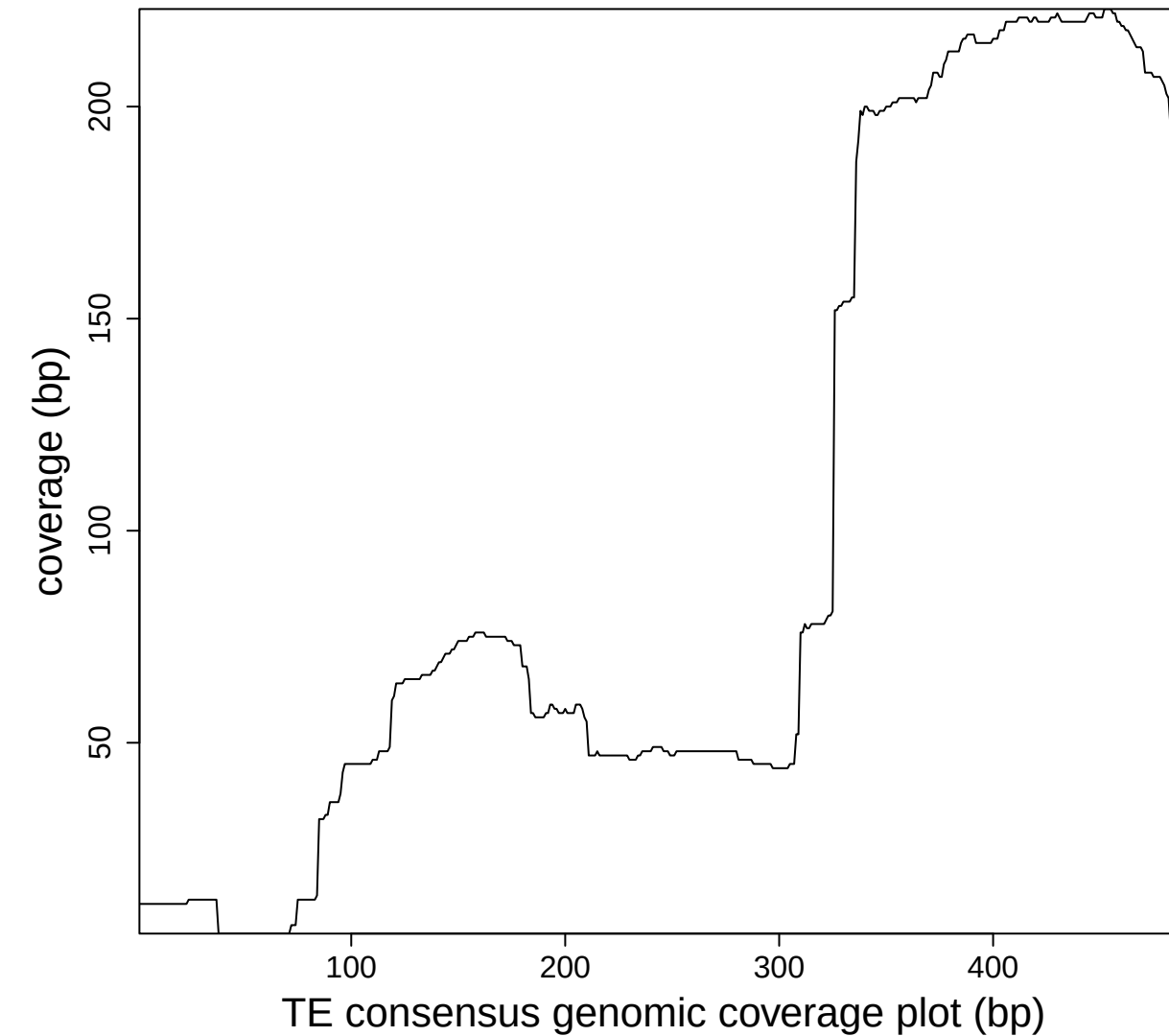
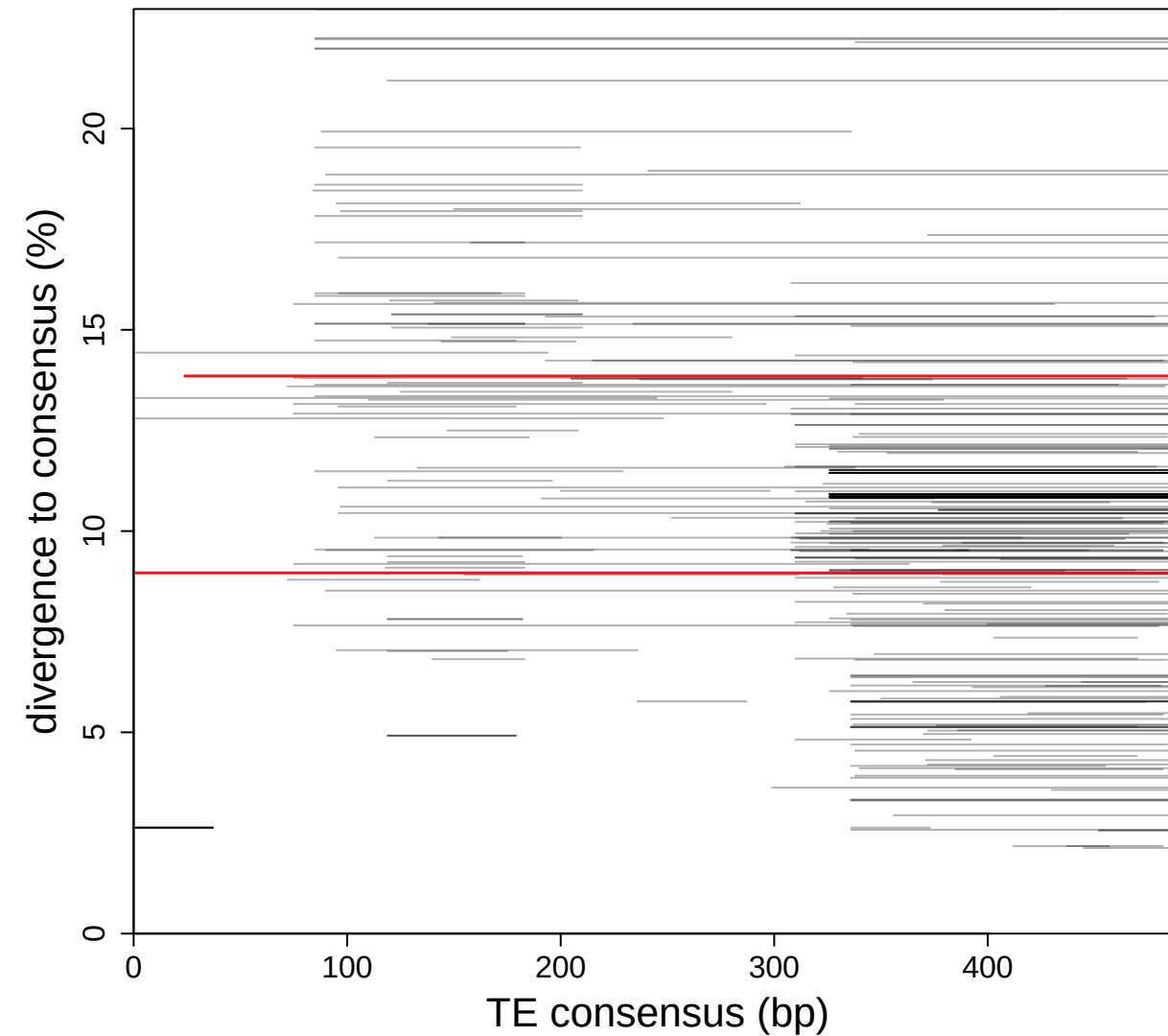


9.fasta.b.bed uf.bed g_2.bed fm_1.bed 0_0_bcln.fa aln.fa cl.f
size: 774bp; fragments: 134; full length: 0 (>=696.6bp)

After TEtrimmer Extended plot Blue lines are boundaries

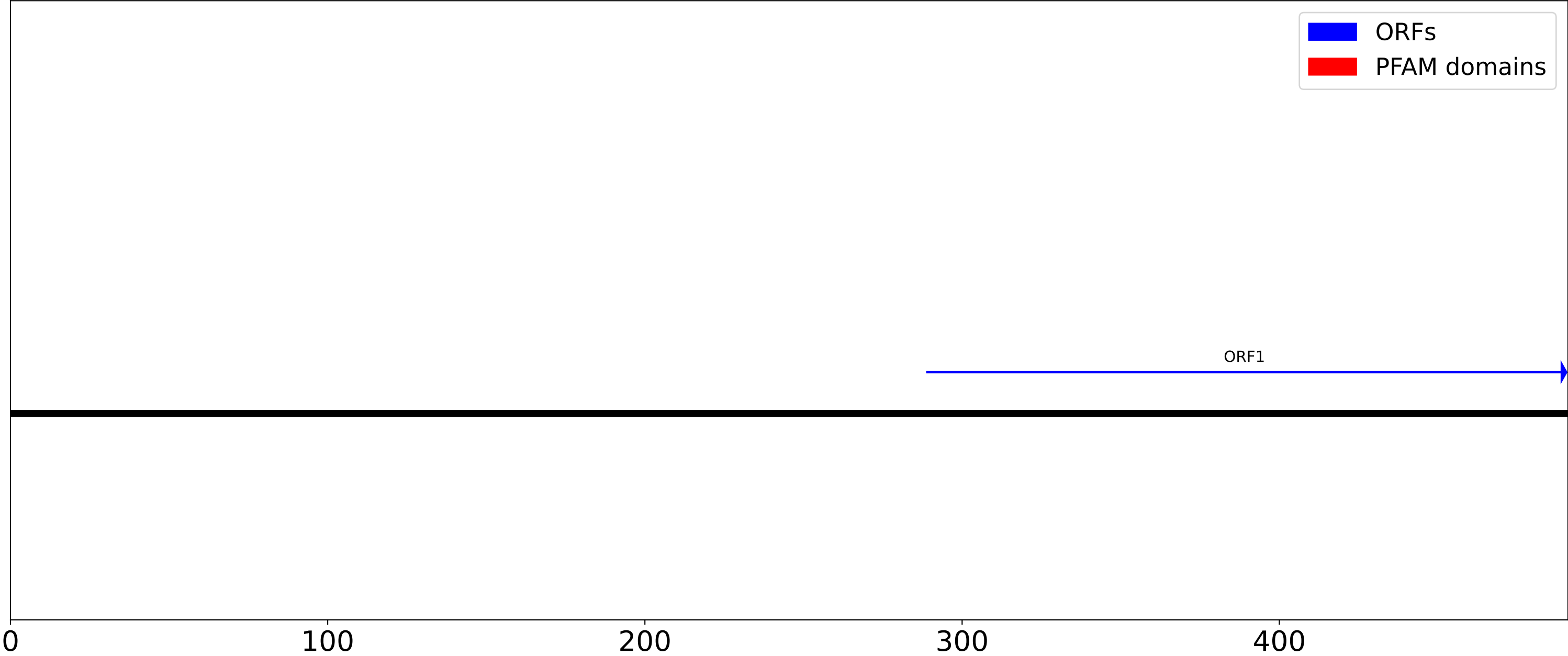


TE: rnd_5_family_4029
size: 491bp; fragments: 296; full length: 2 (≥ 441.9 bp)



Before TEtrimmer 491 bp

Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

