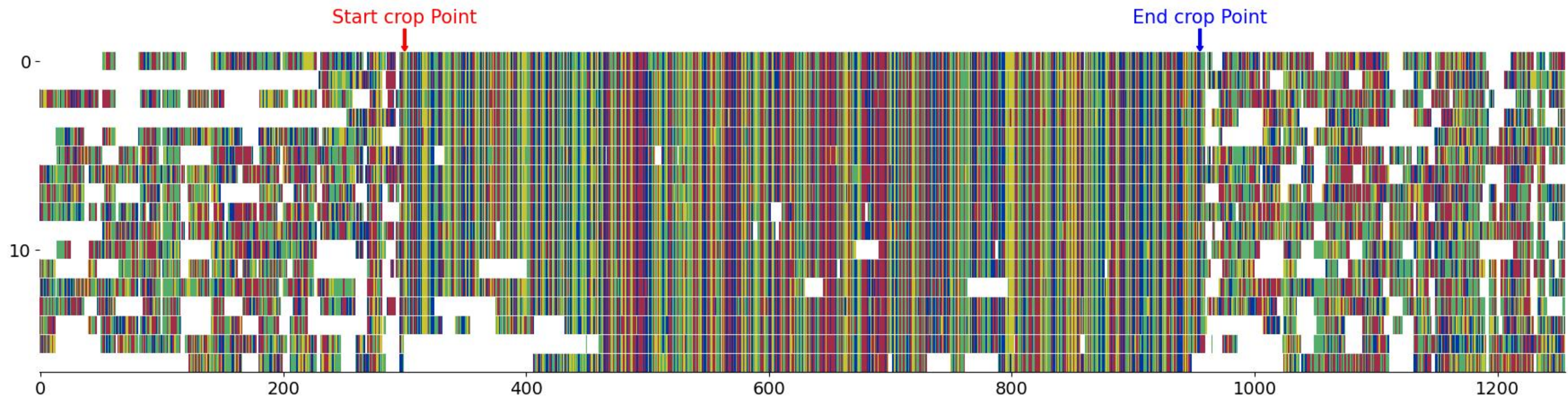
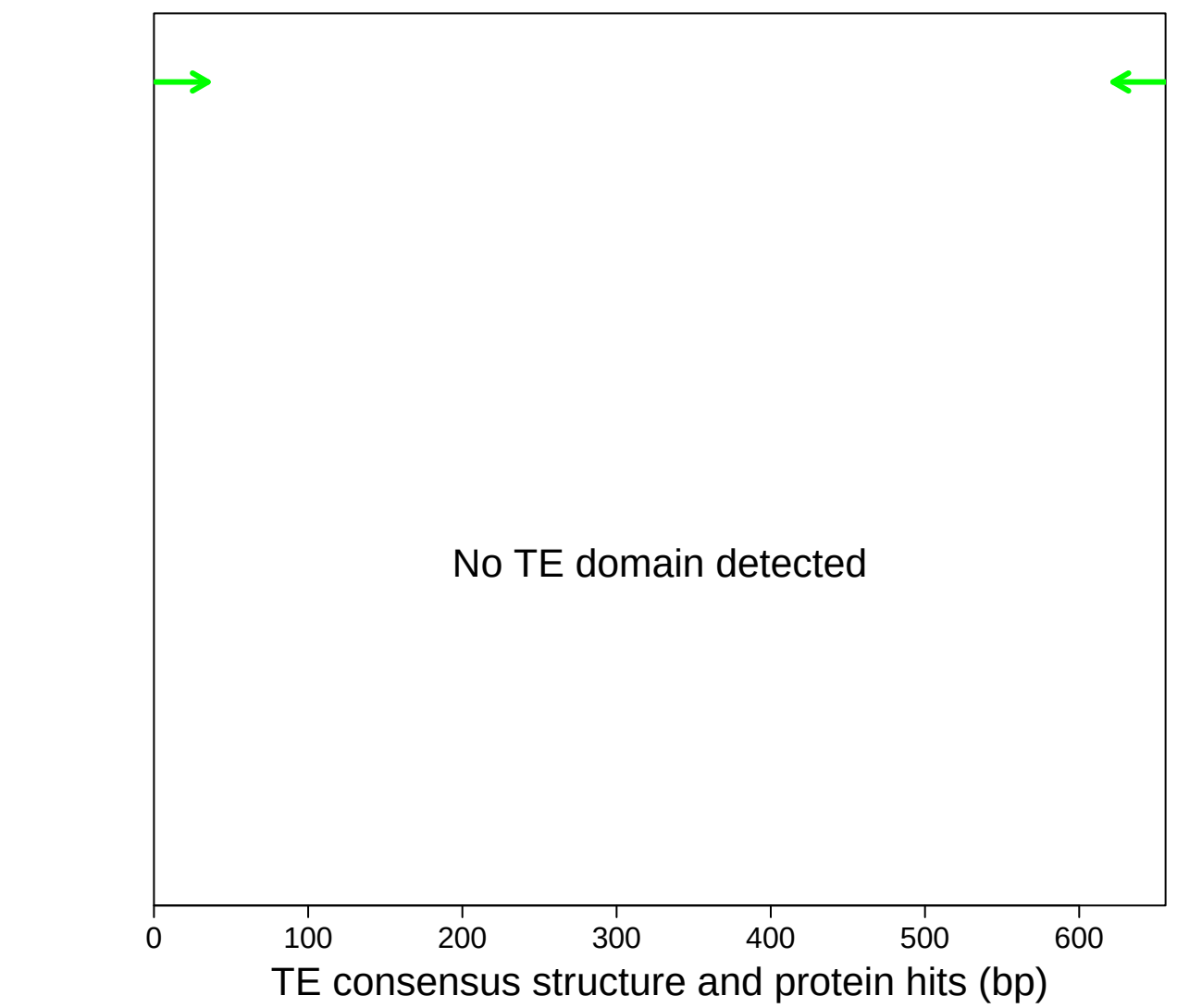
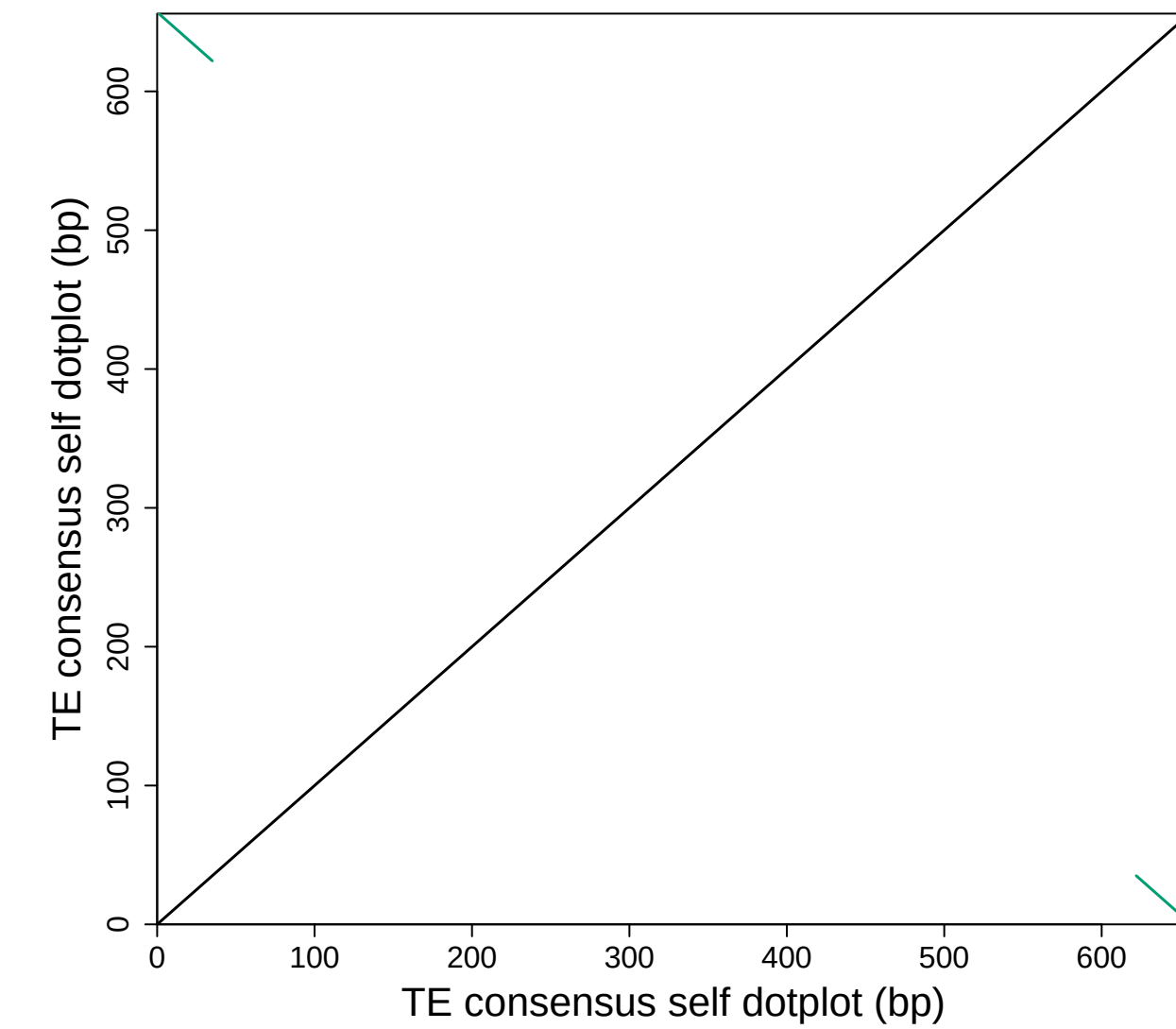
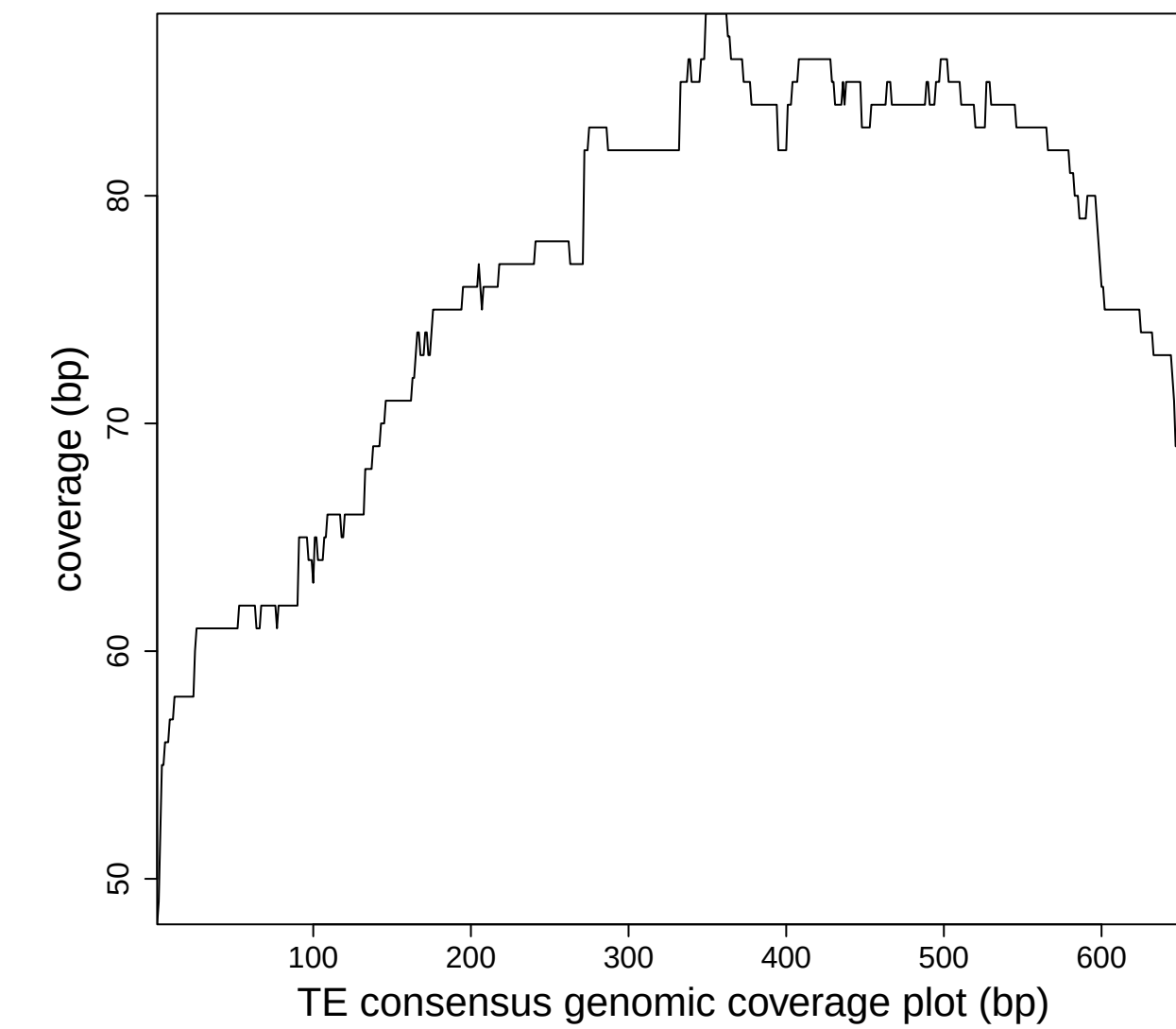
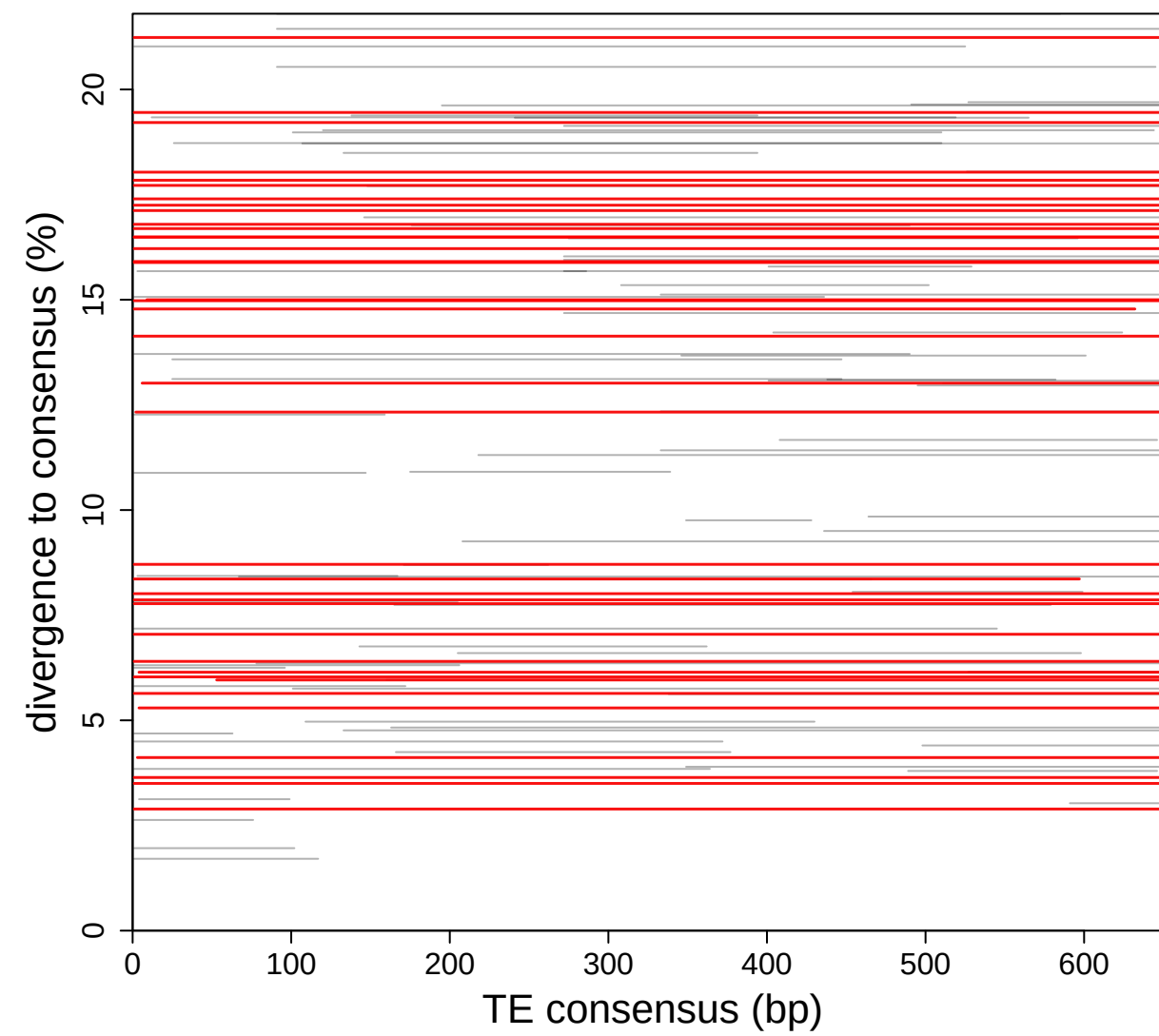




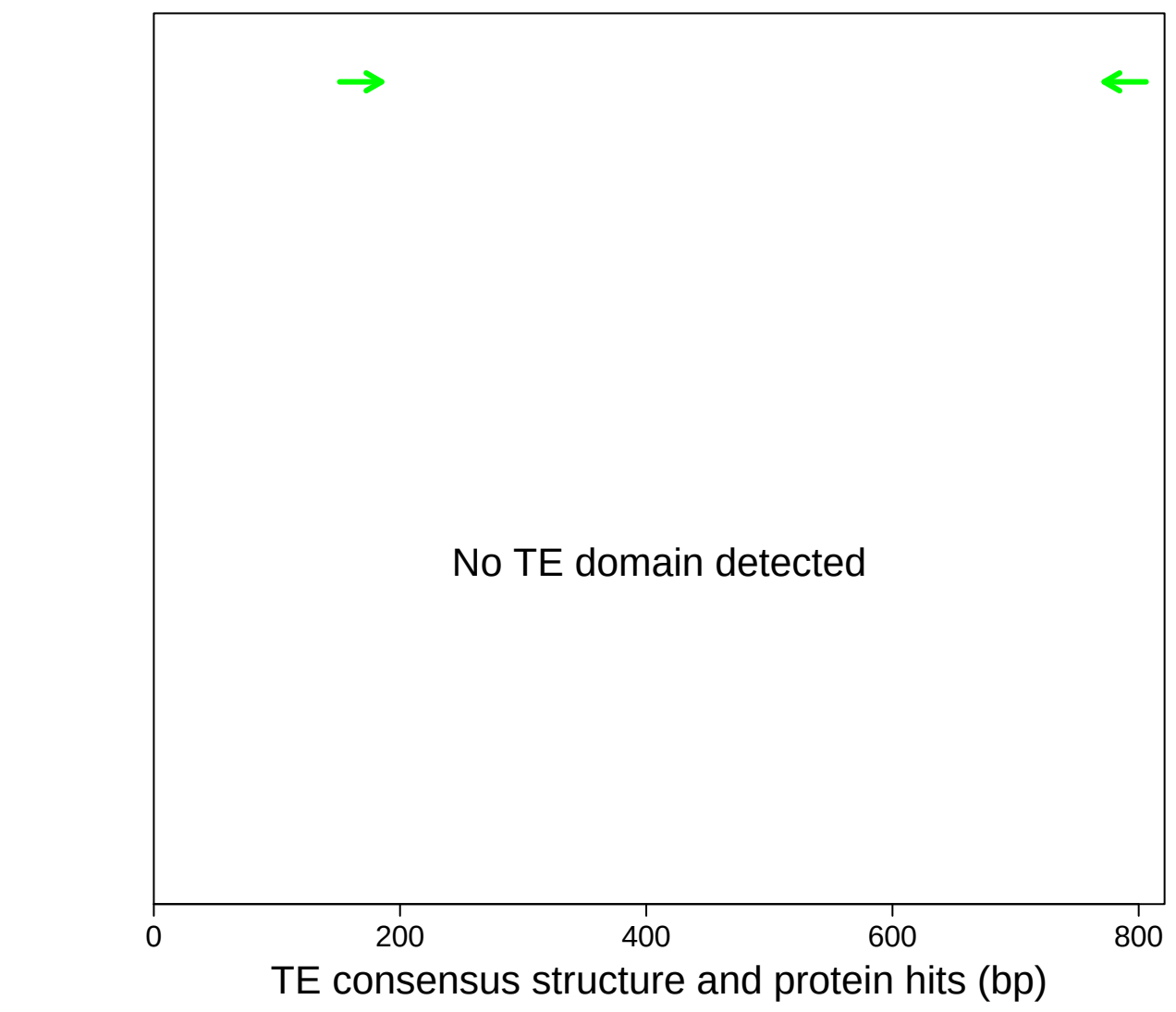
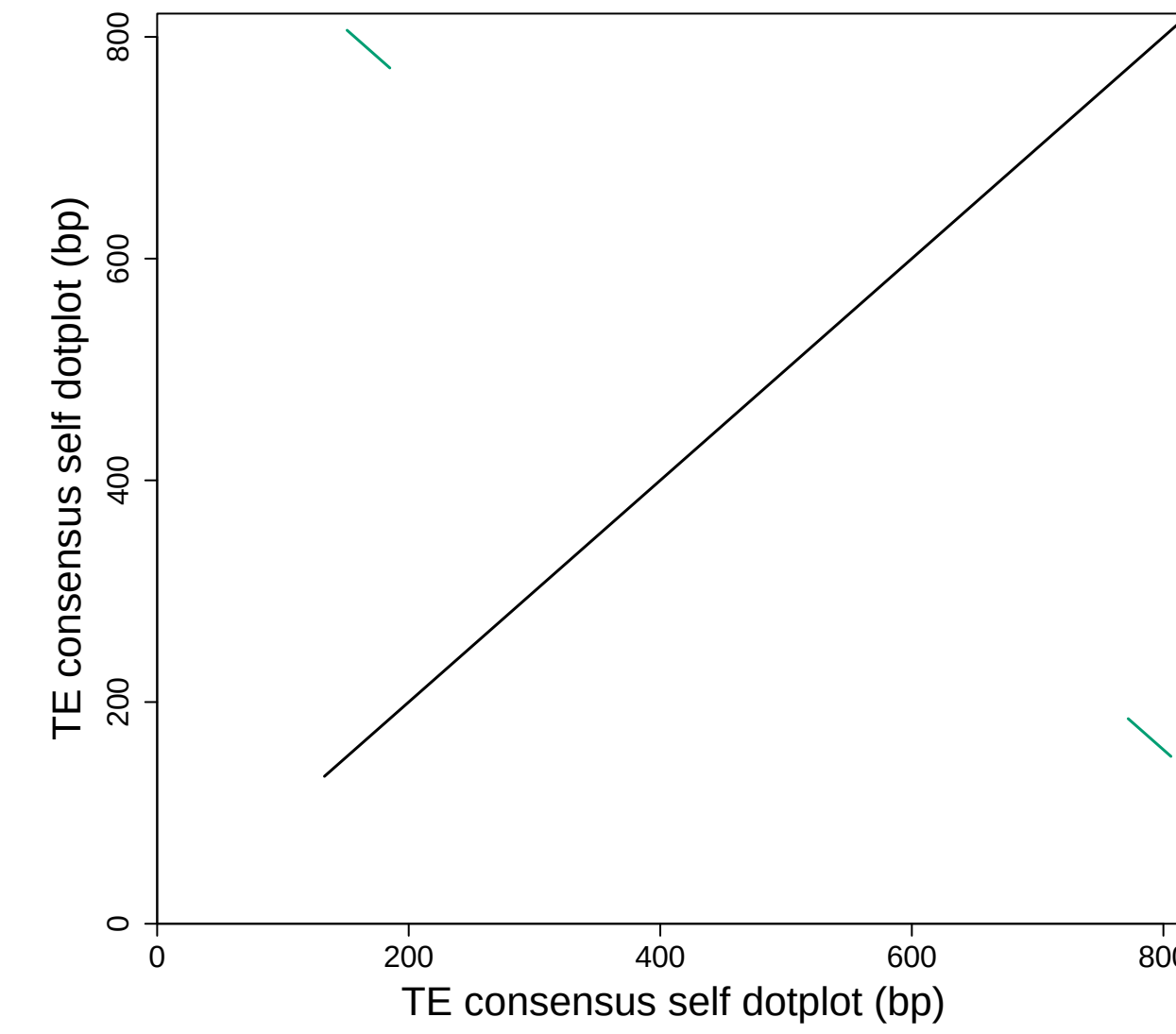
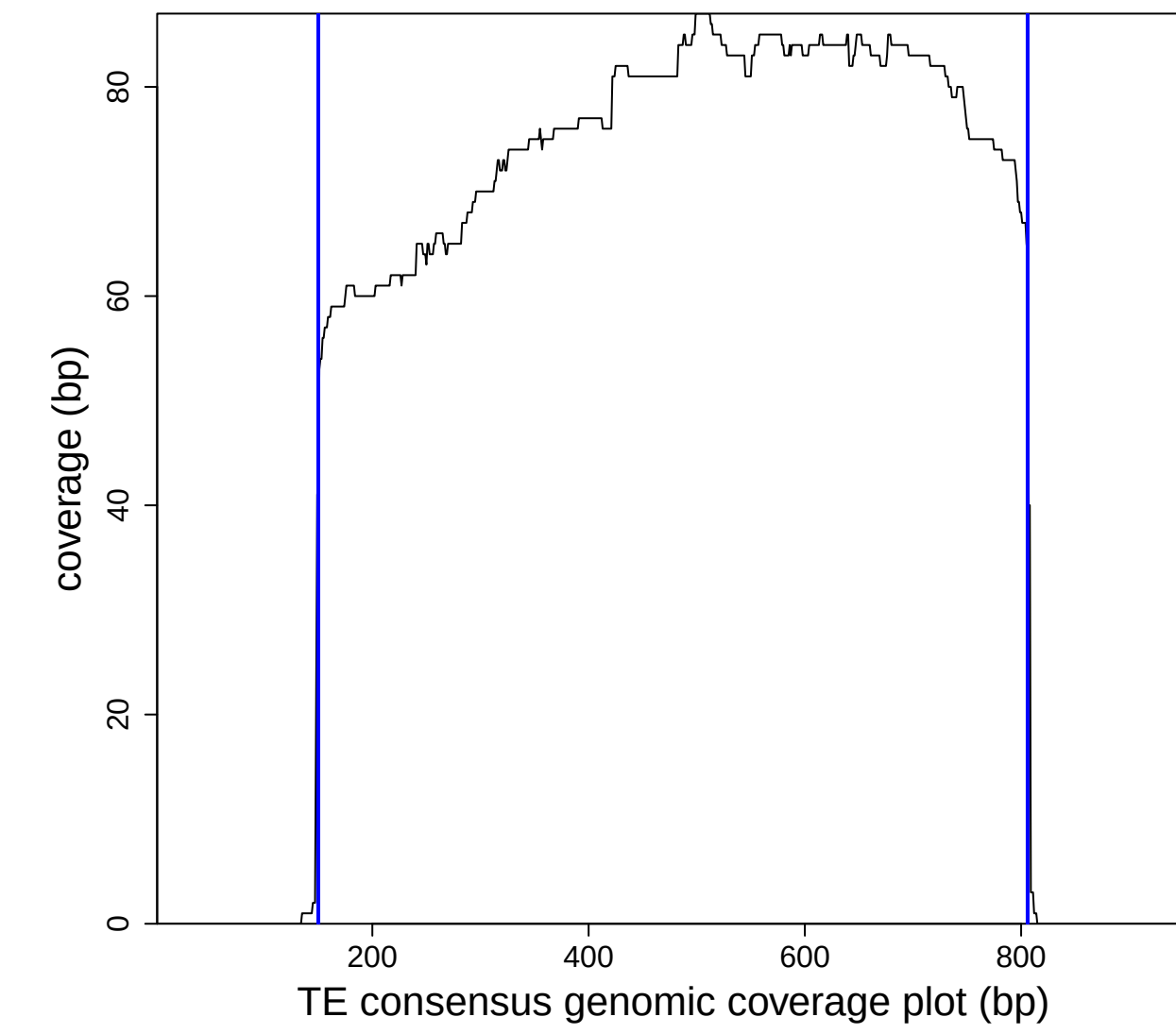
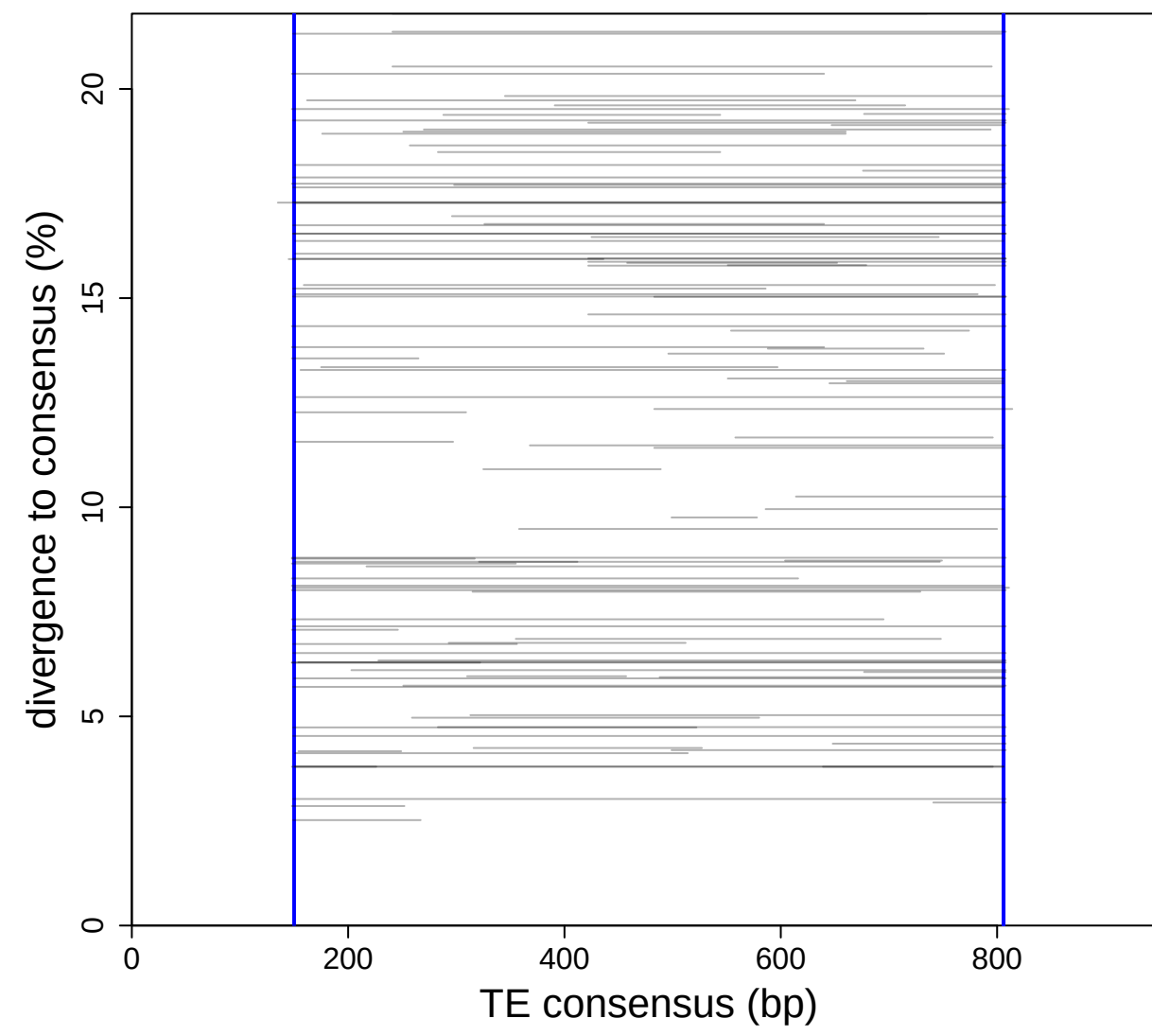
a.b.bed_uf.bed_g_2.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.f
size: 656bp; fragments: 121; full length: 38 (≥ 590.4 bp)

After TEtrimmer 656 bp

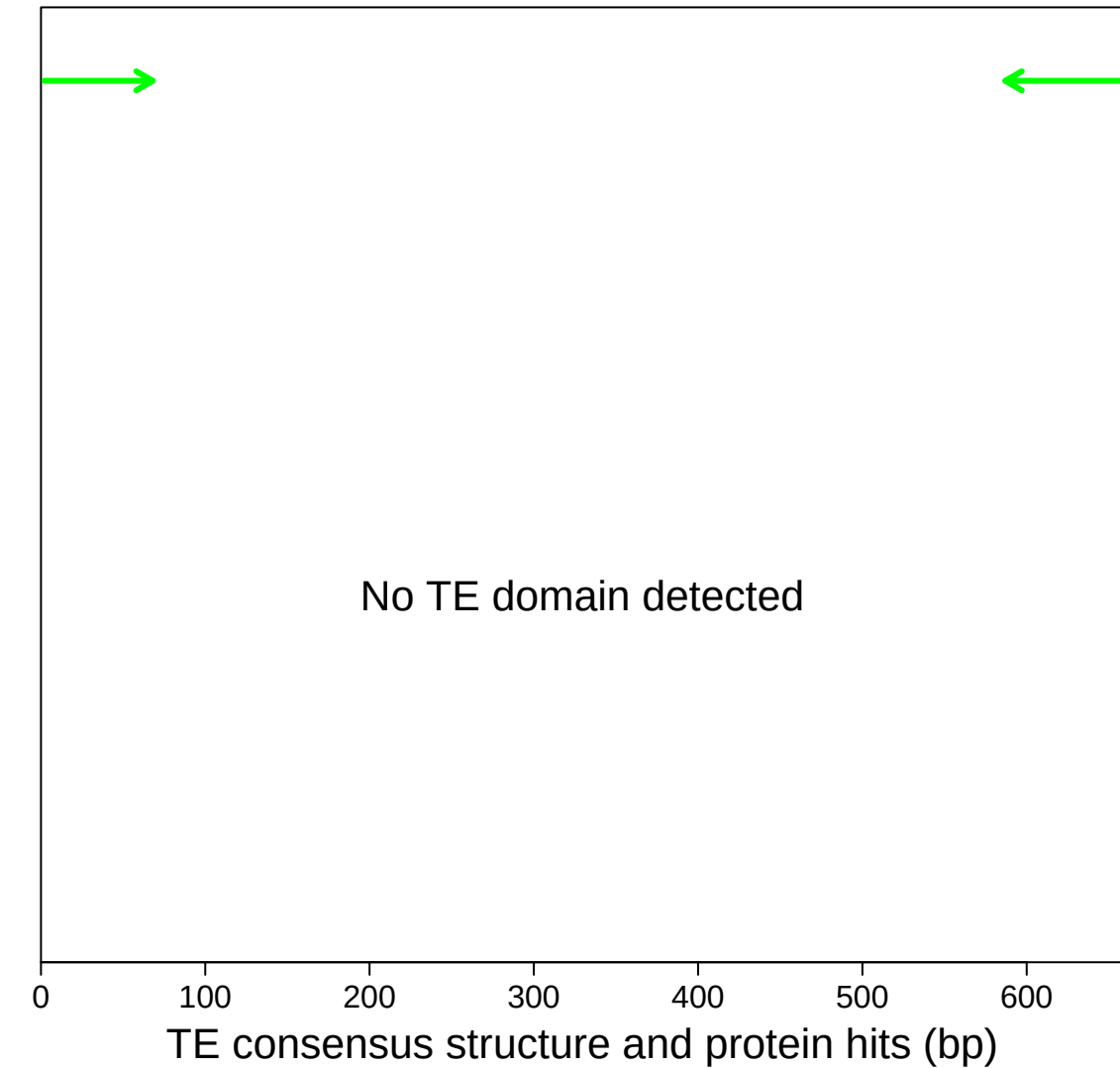
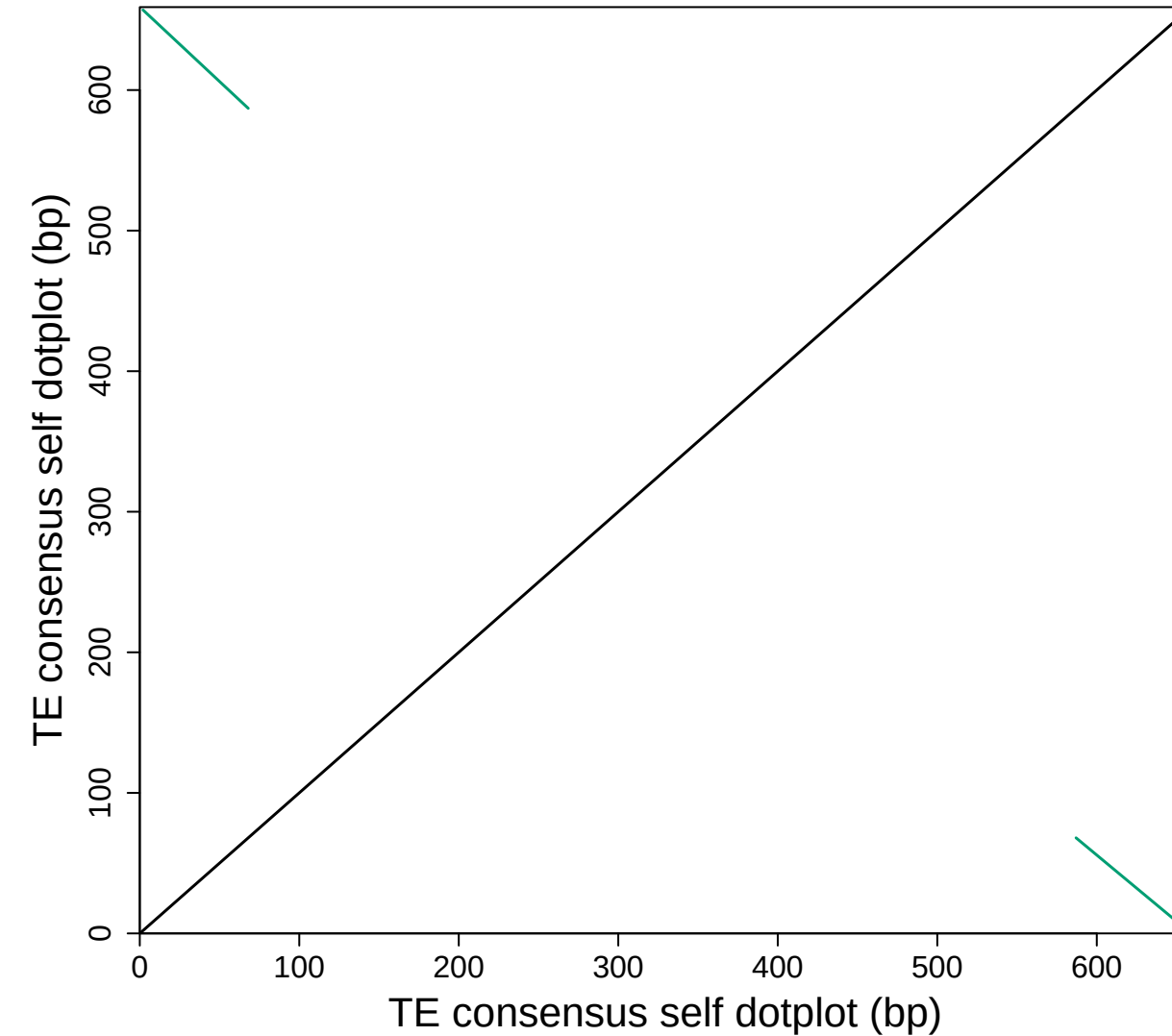
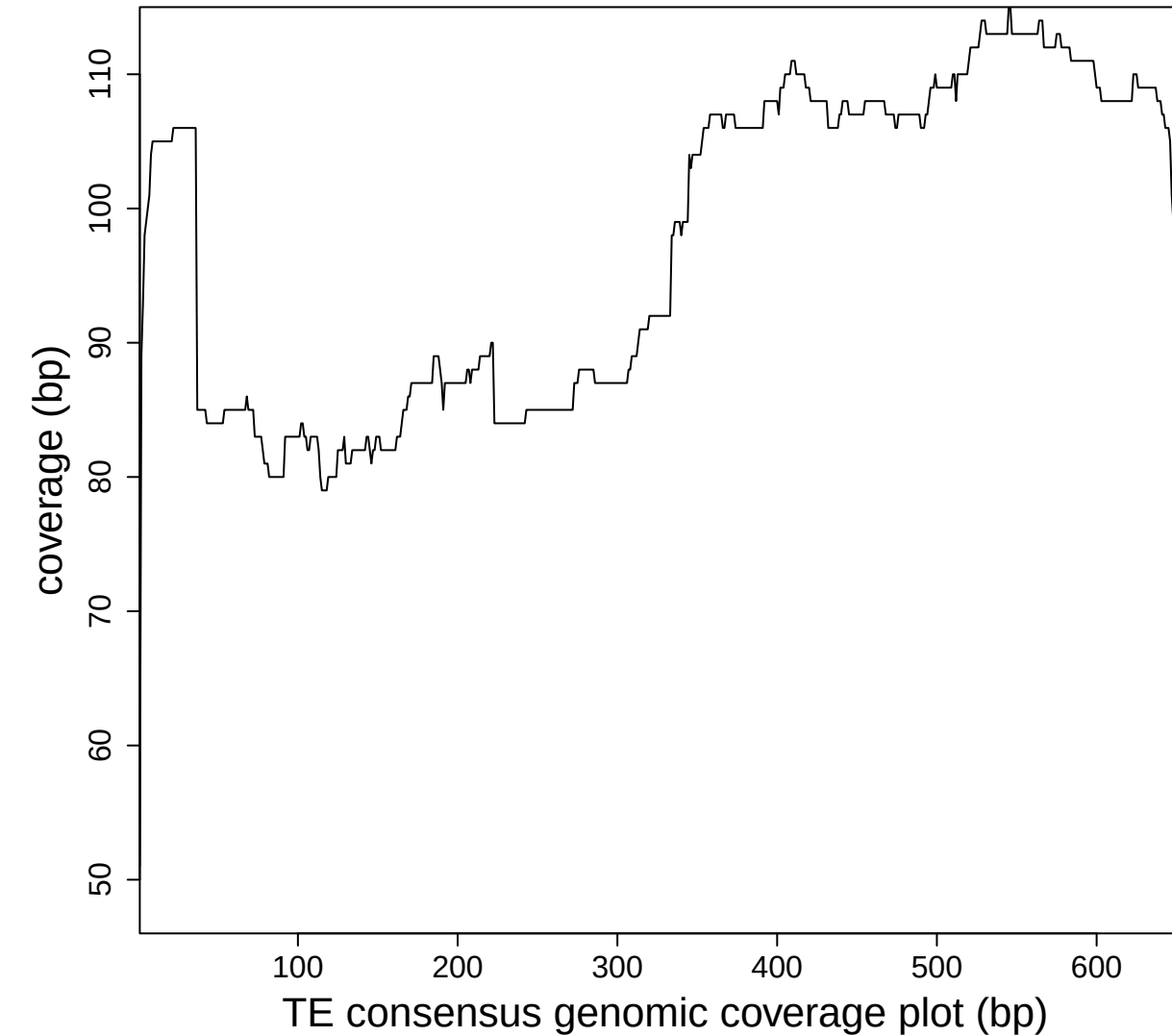
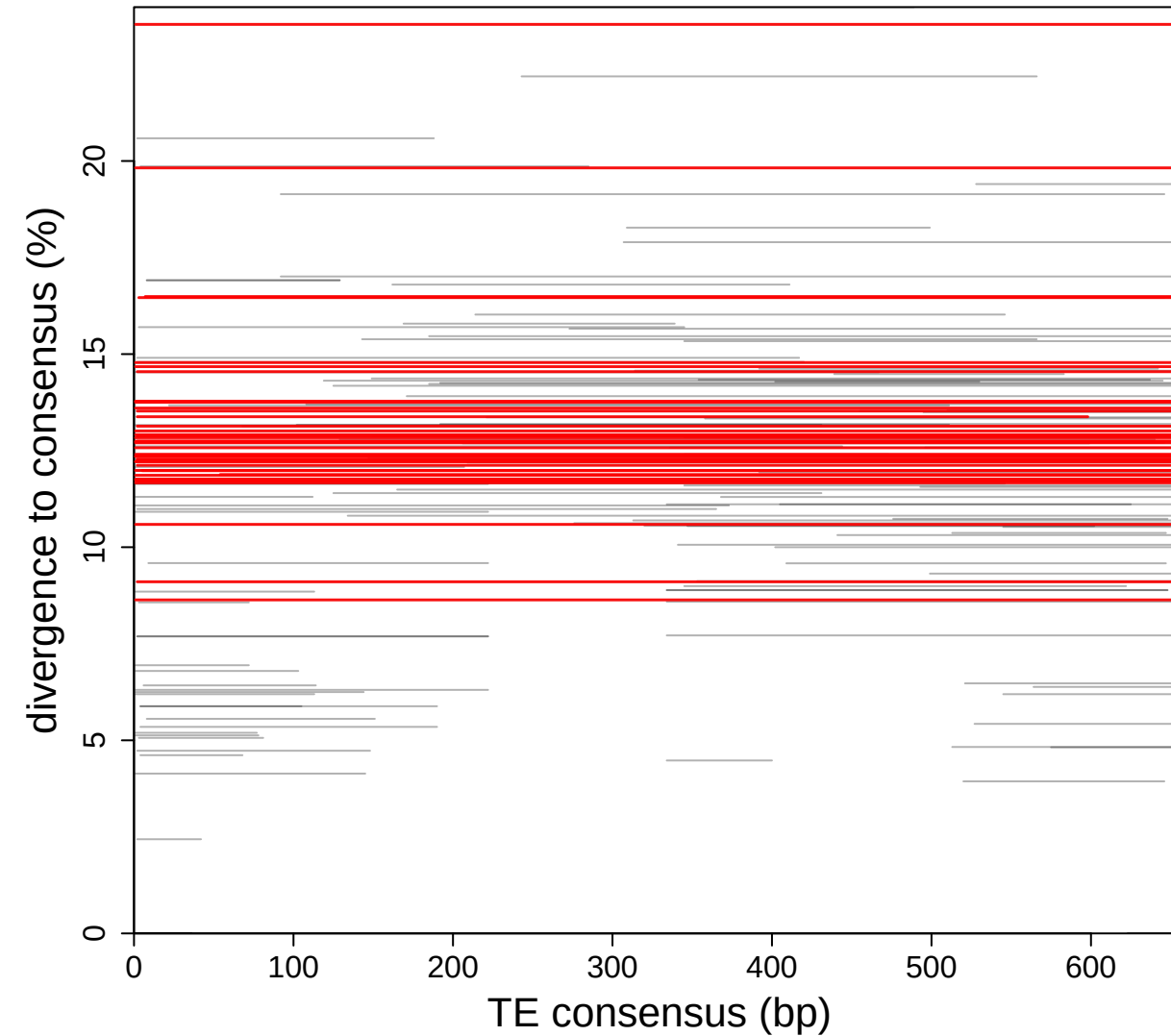


0.fasta.b.bed uf.bed g_2.bed fm_1.bed 0_0_bcln.fa aln.fa cl.f
size: 956bp; fragments: 121; full length: 0 (>=860.4bp)

After TEtrimmer Extended plot Blue lines are boundaries

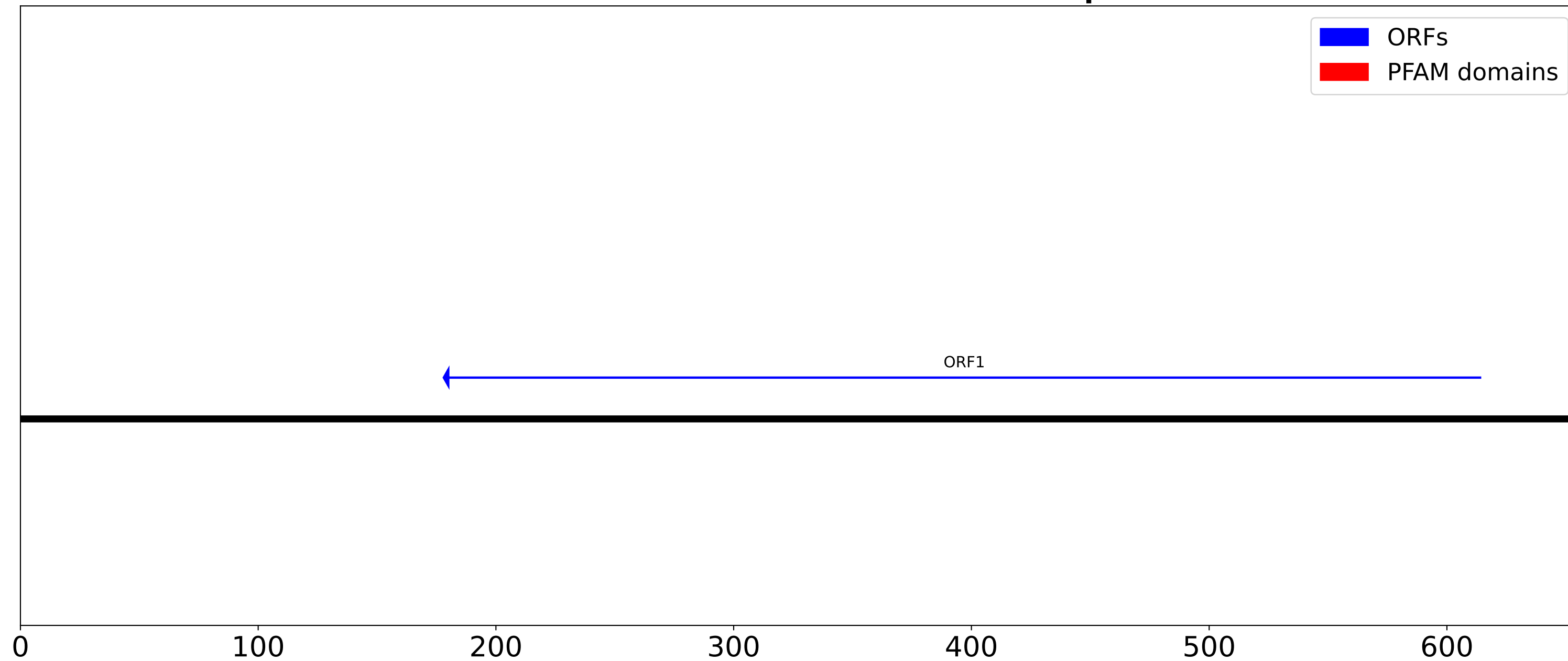


TE: rnd_5_family_3880
size: 659bp; fragments: 190; full length: 44 (≥ 593.1 bp)

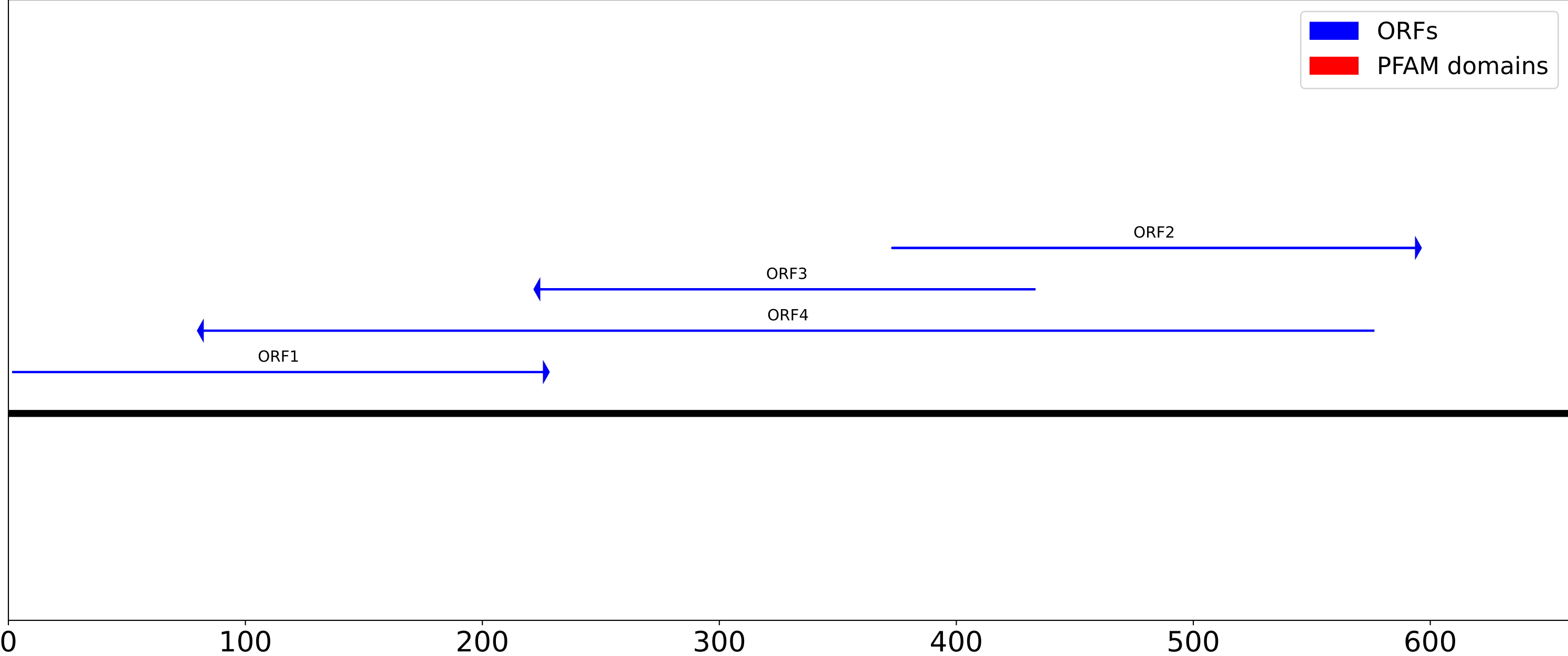


Before TEtrimmer 659 bp

After TEtrimmer ORF and PFAM domain plot



Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

