

Start crop Point

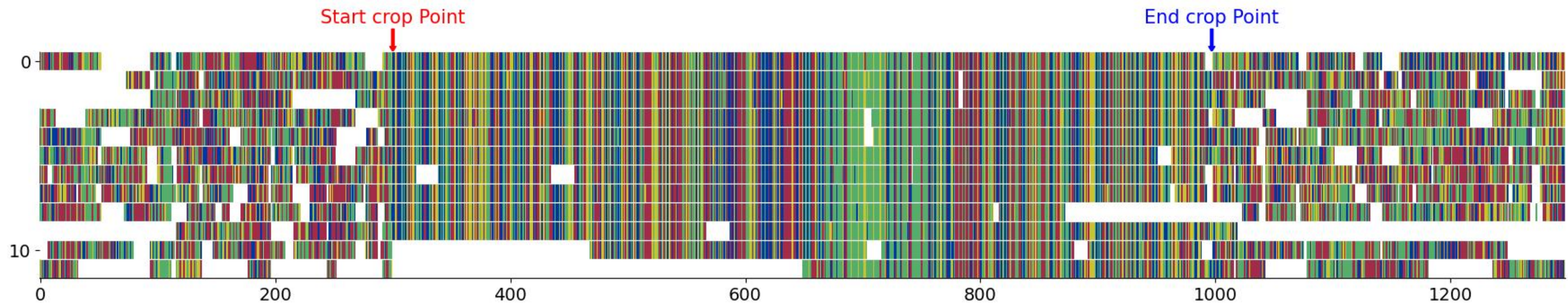


End crop Point



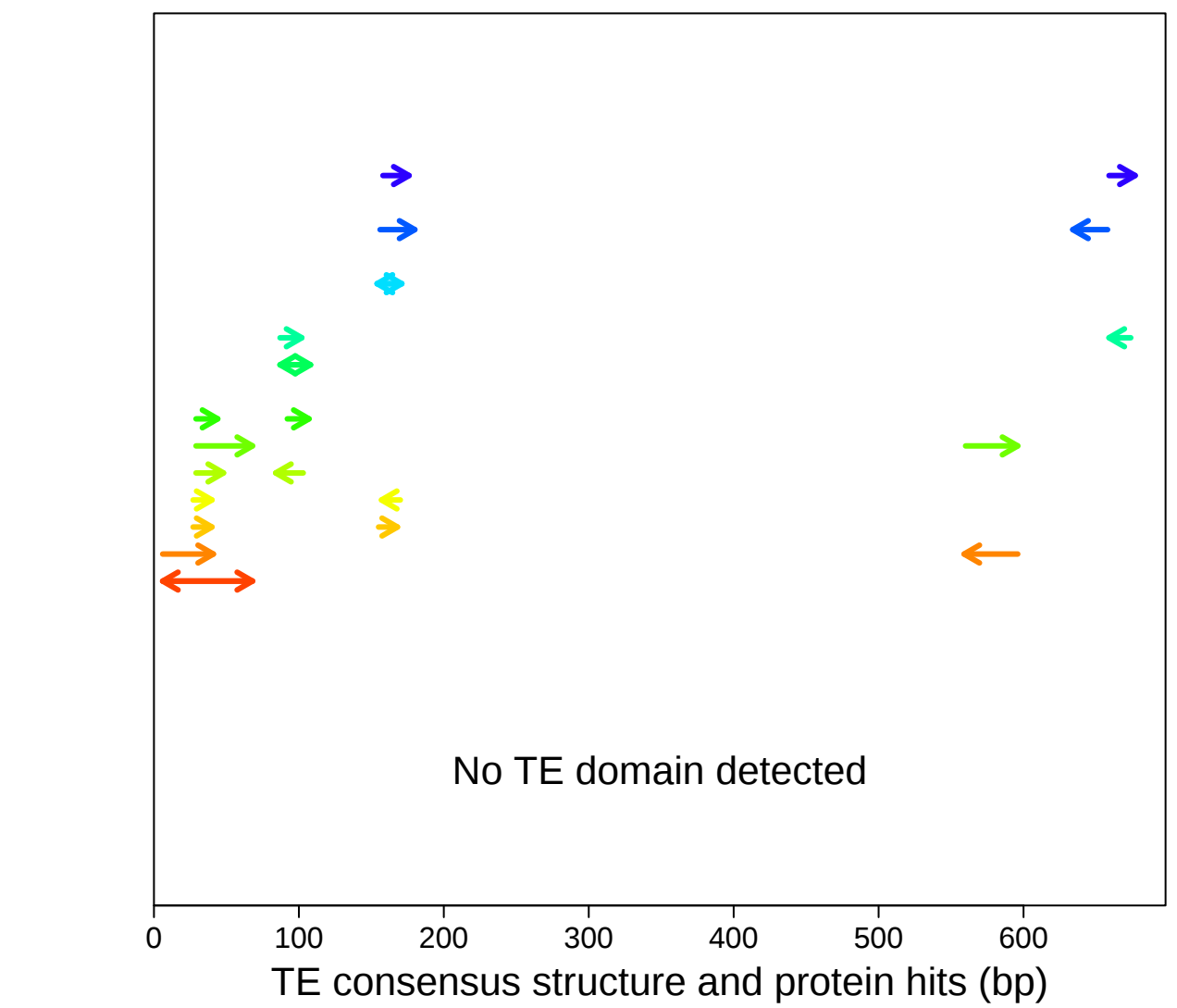
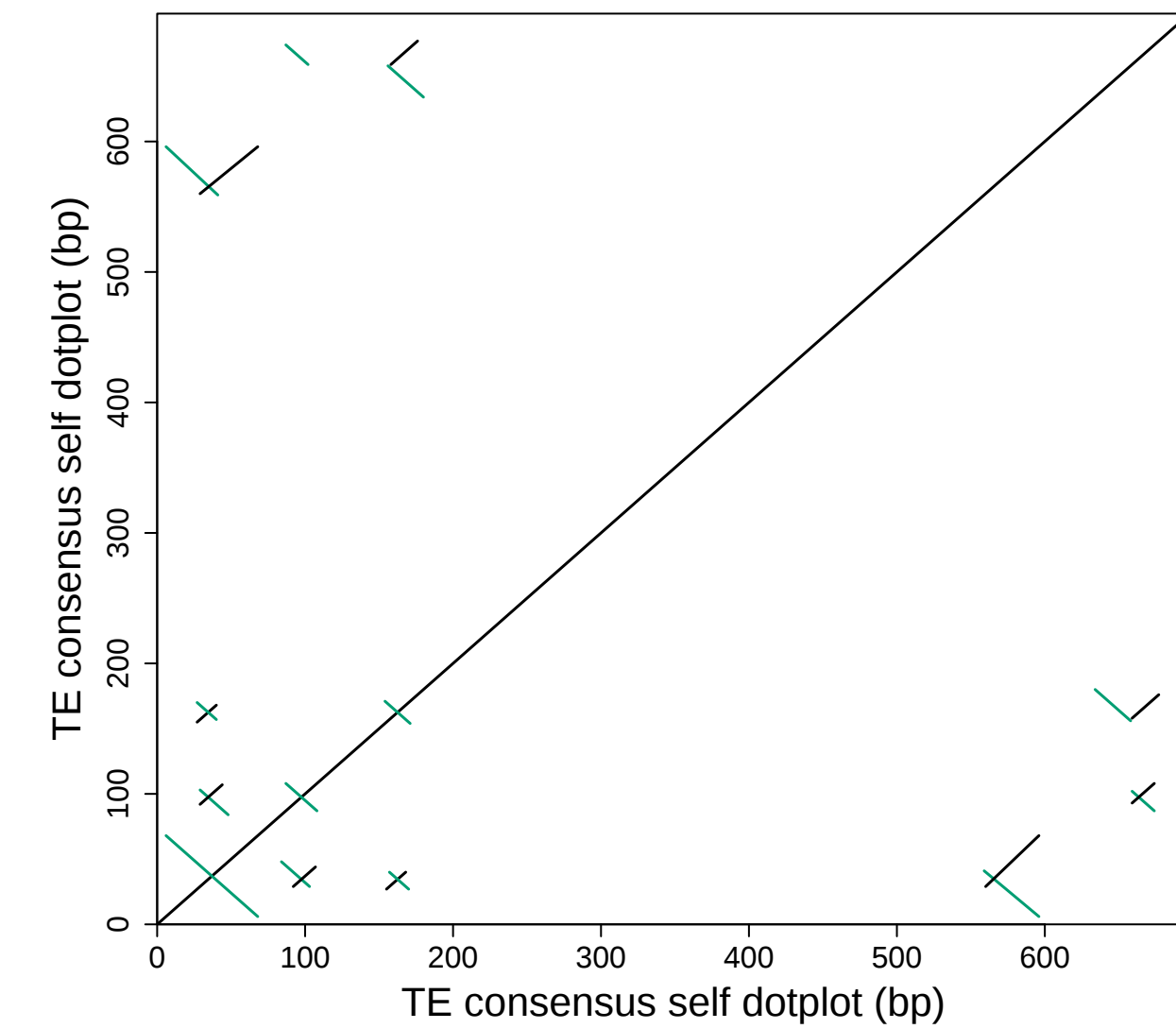
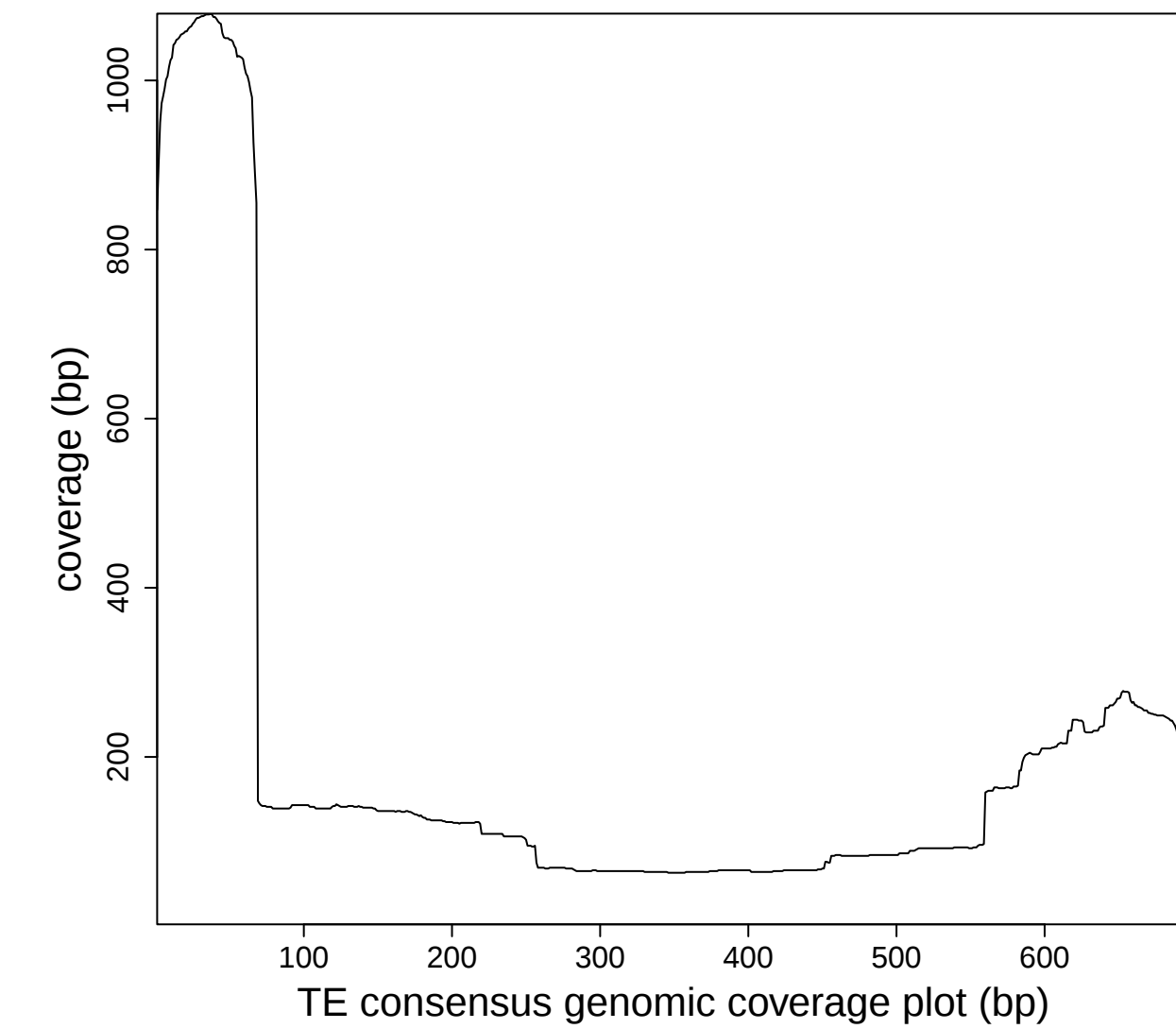
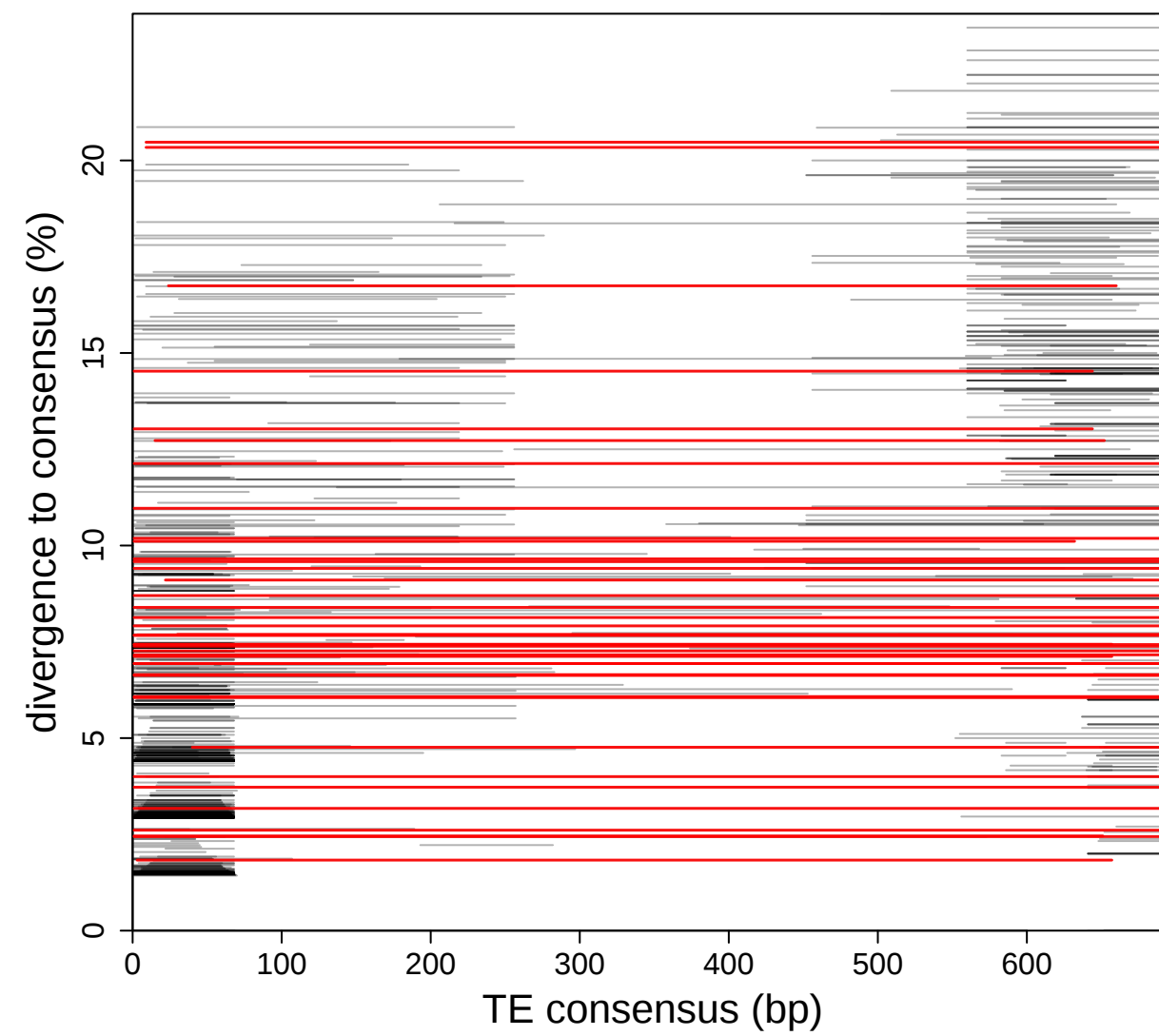
MSA length = 698





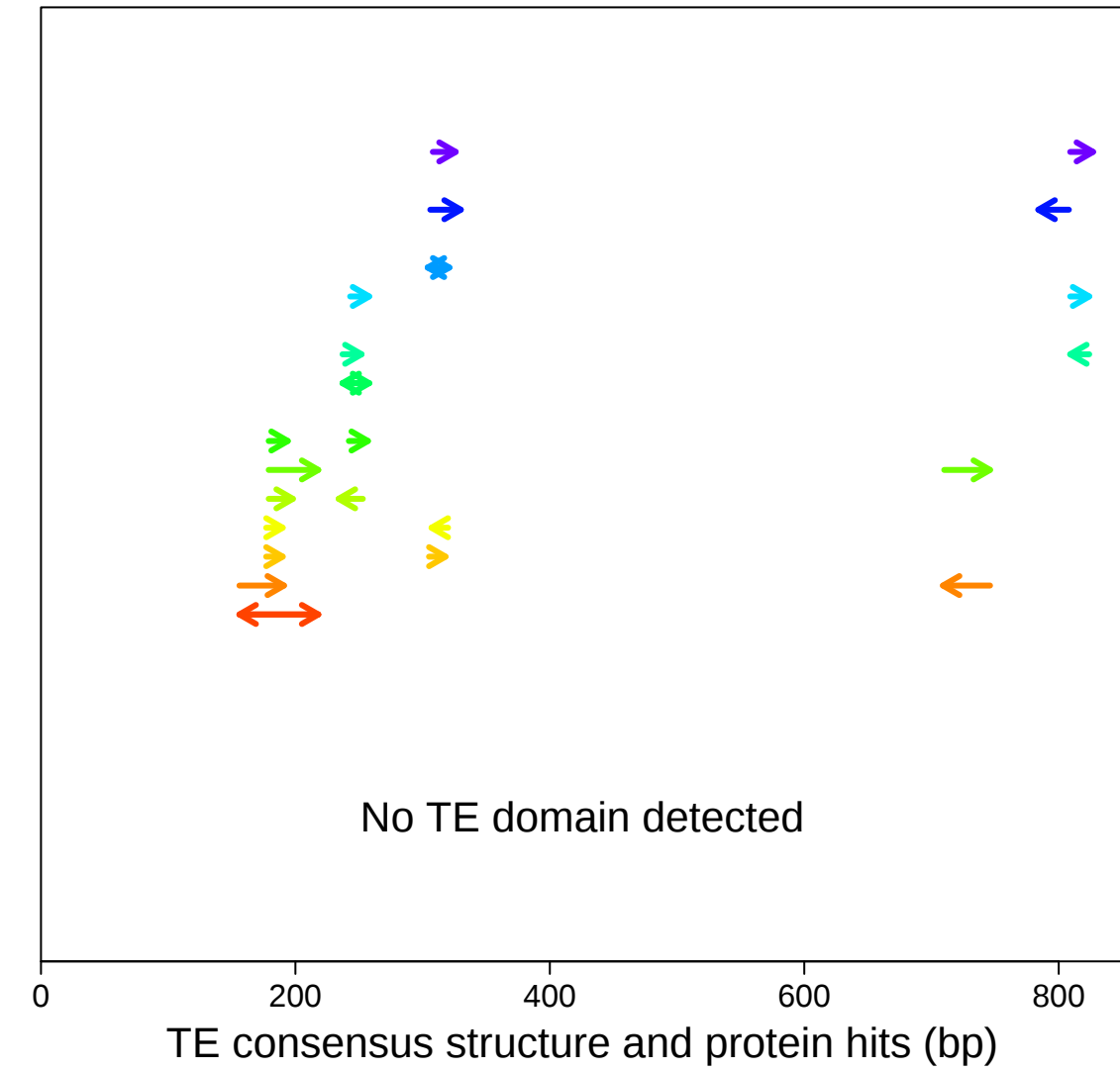
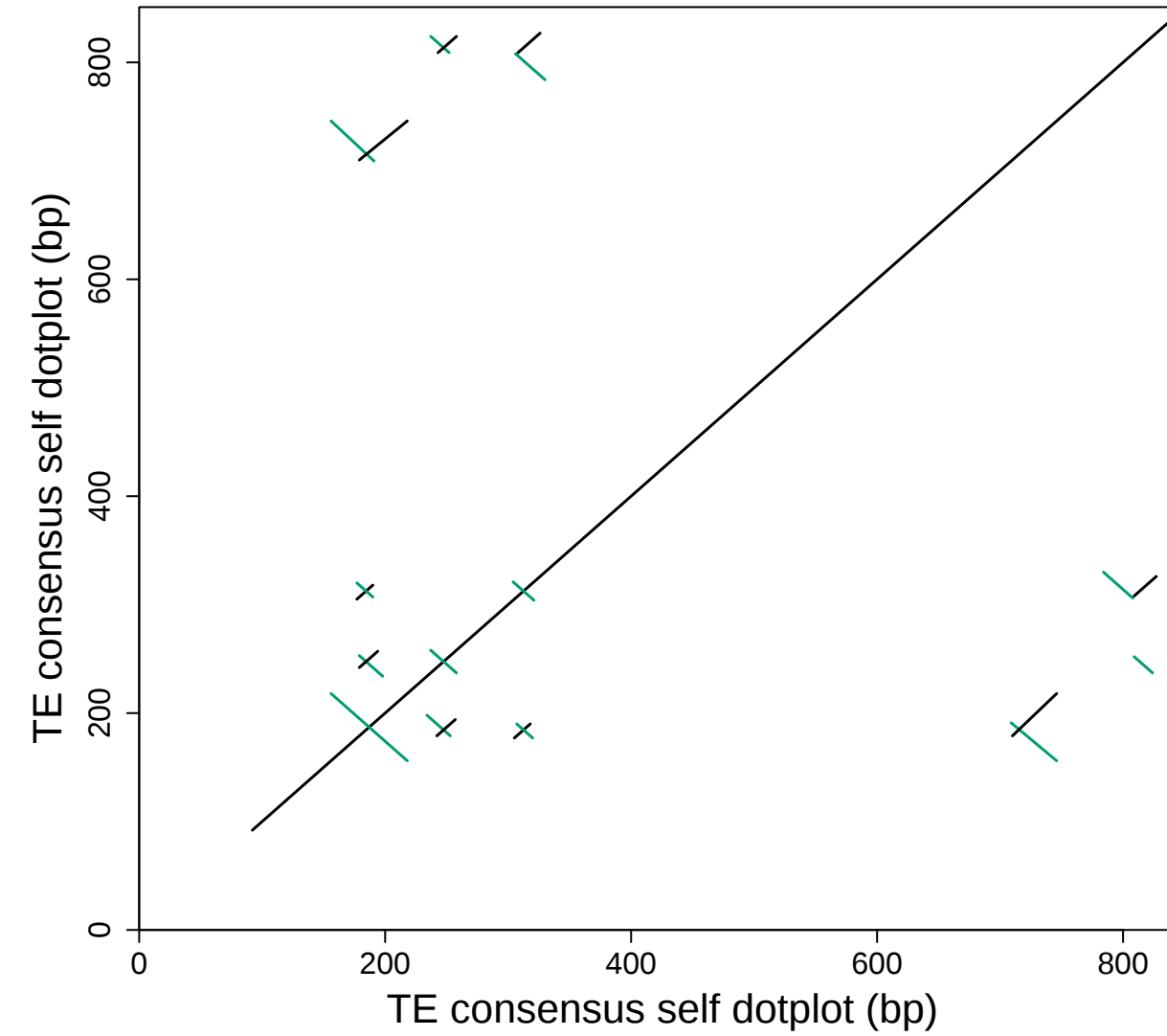
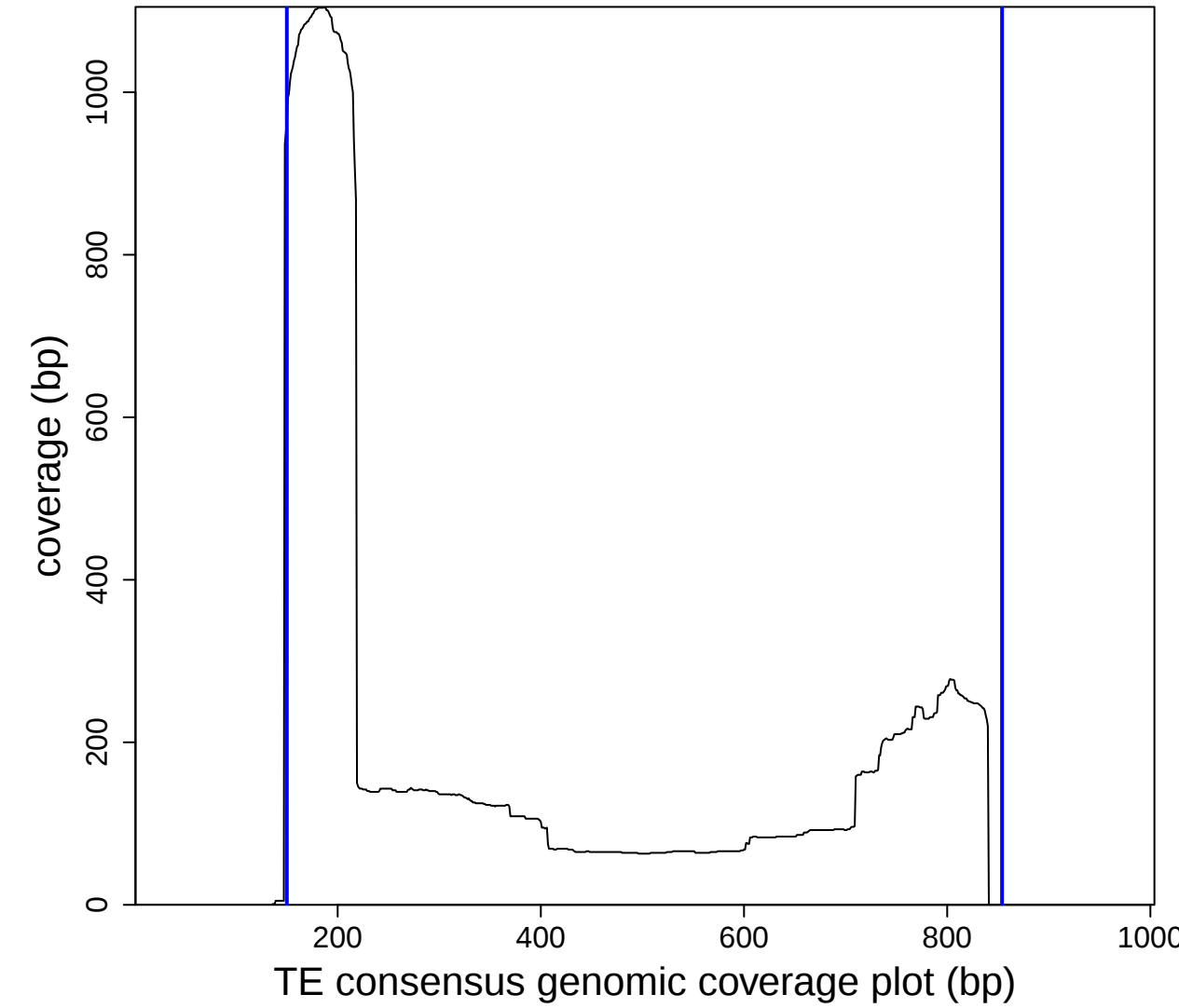
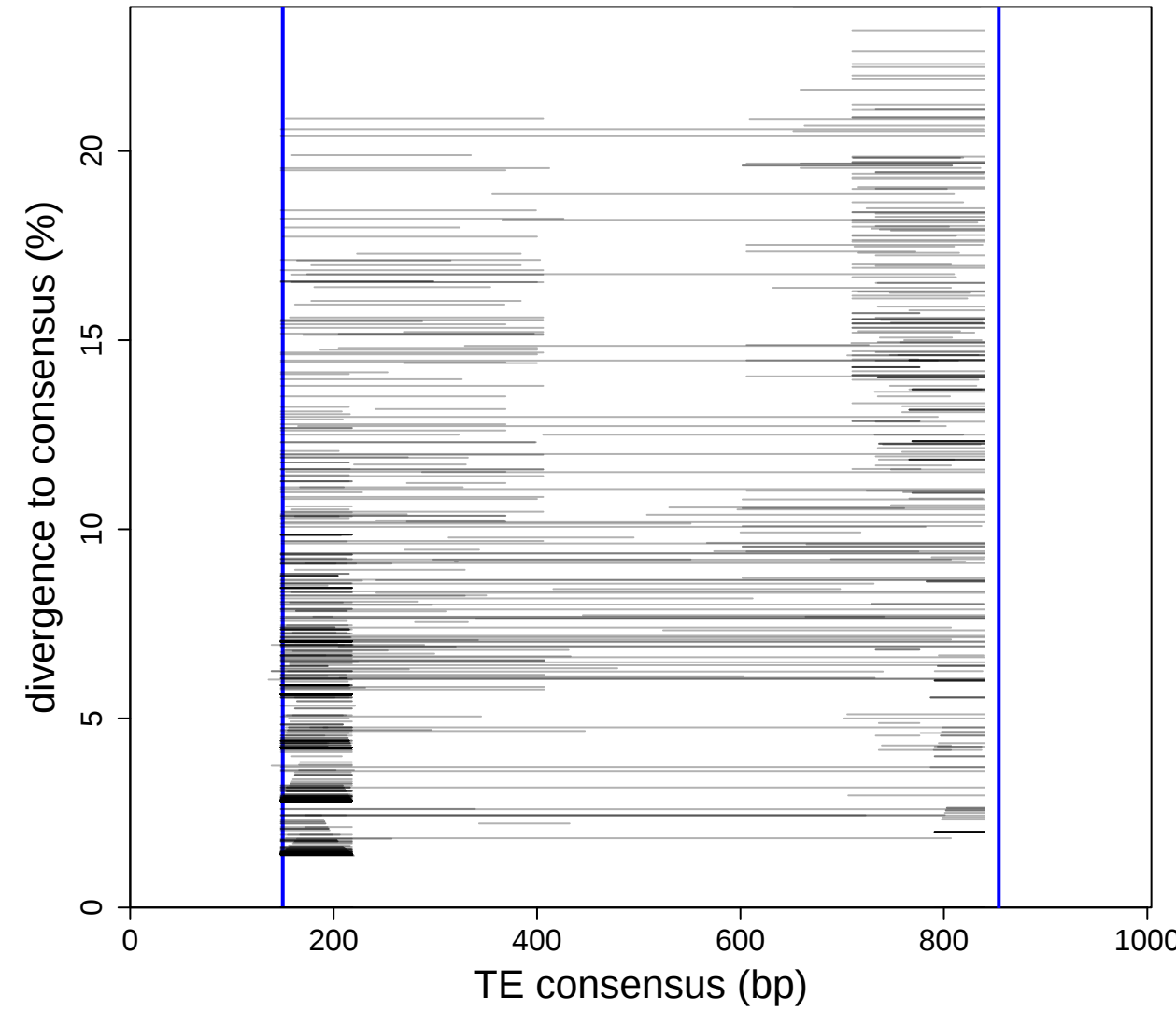
a.b.bed_uf.bed_g_3.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.f
size: 698bp; fragments: 1359; full length: 43 (>=628.2bp)

After TEtrimmer 698 bp

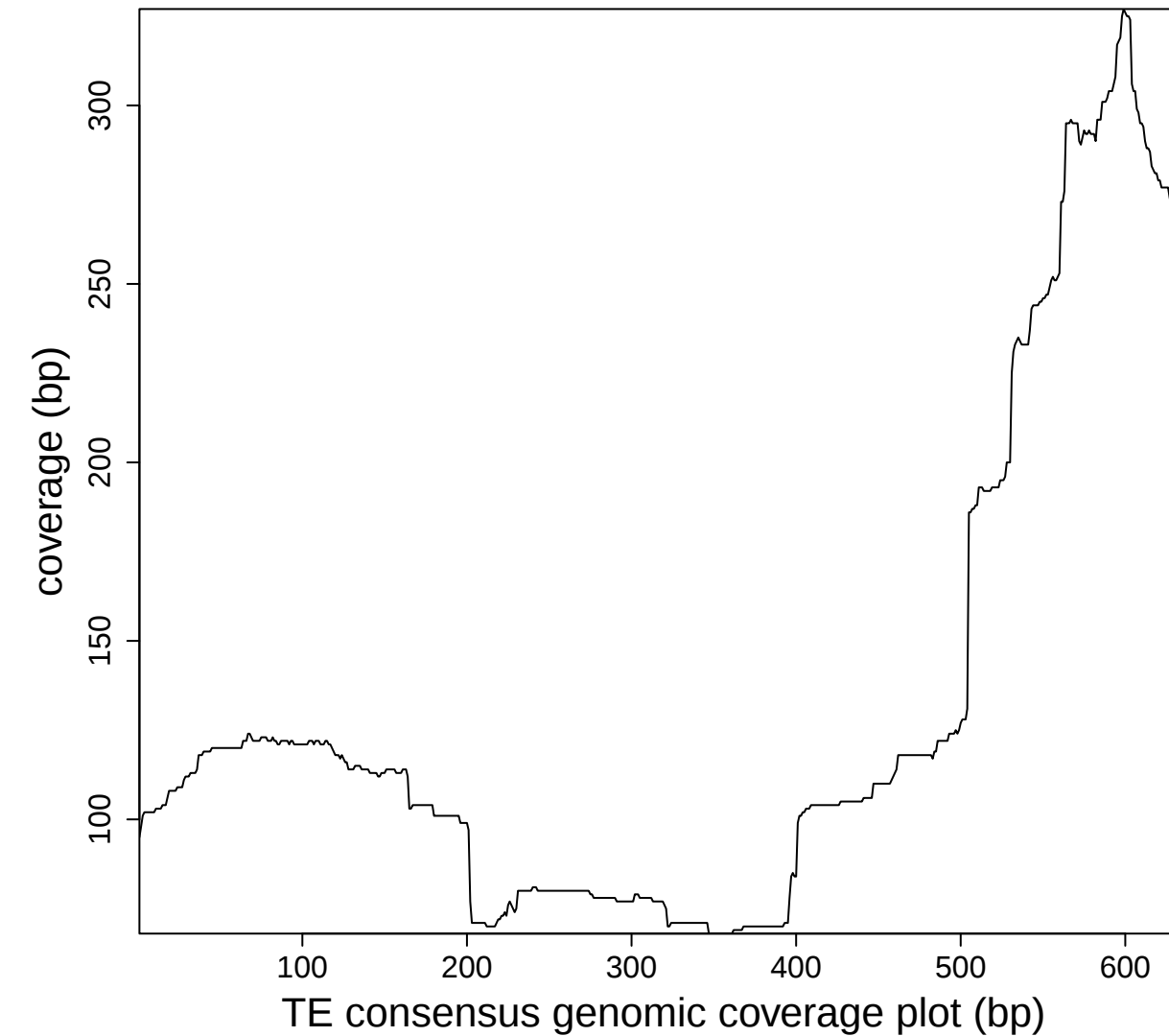
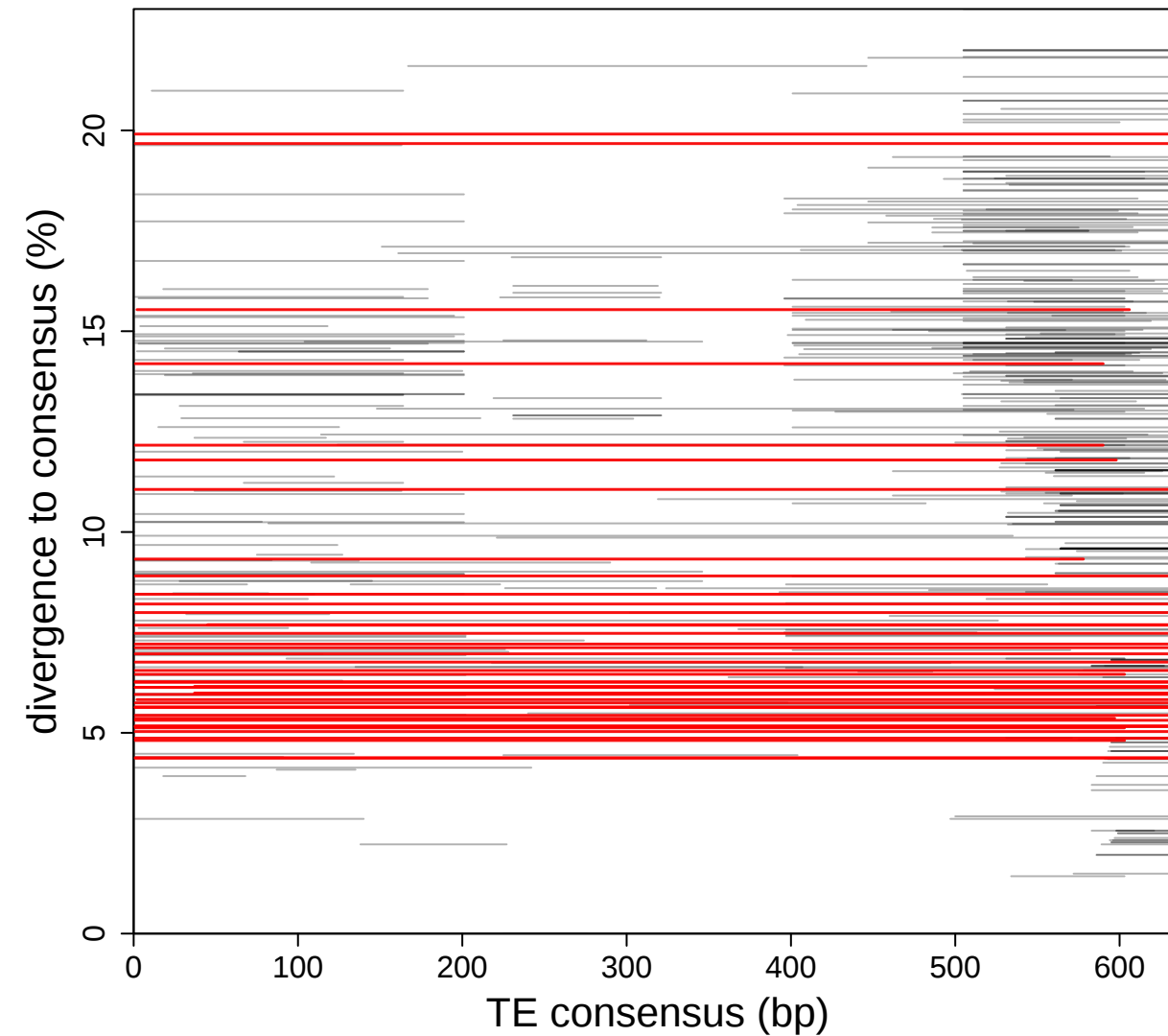


1.fasta.b.bed_uf.bed_g_3.bed_fm_1.bed_0_0_bcIn.fa_aln.fa_cl.f
size: 1004bp; fragments: 1386; full length: 0 (>=903.6bp)

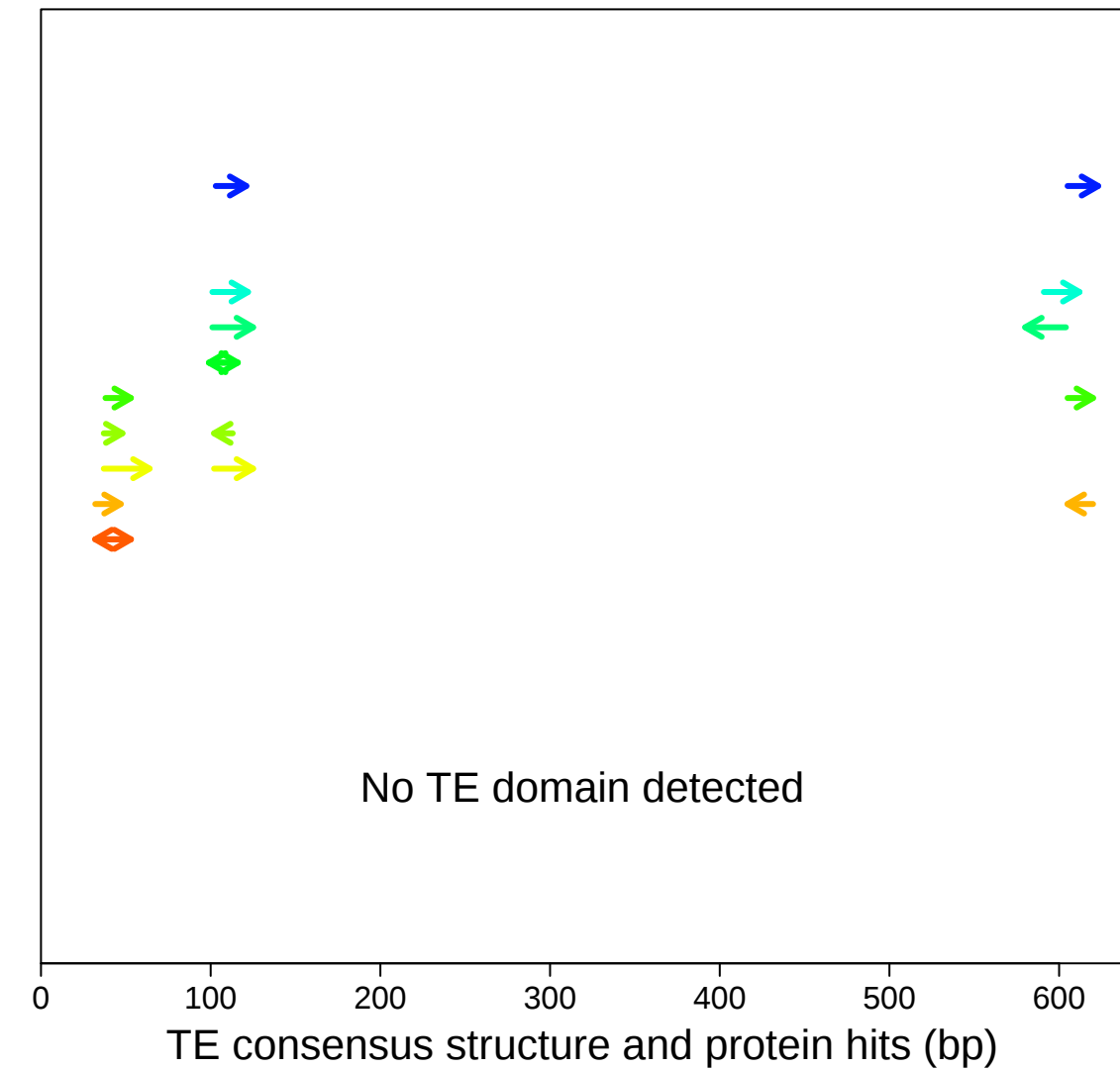
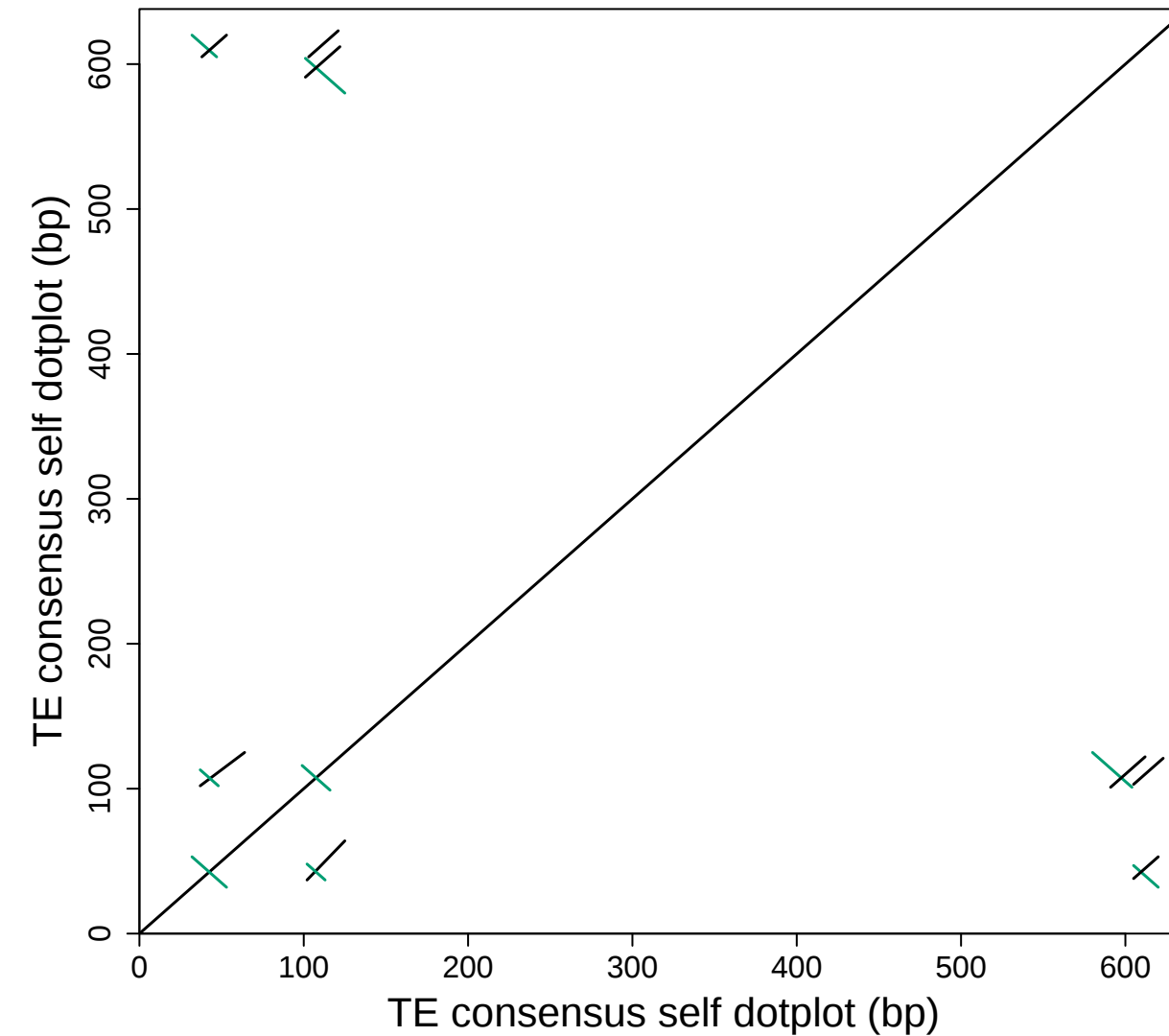
After TEtrimmer Extended plot Blue lines are boundaries



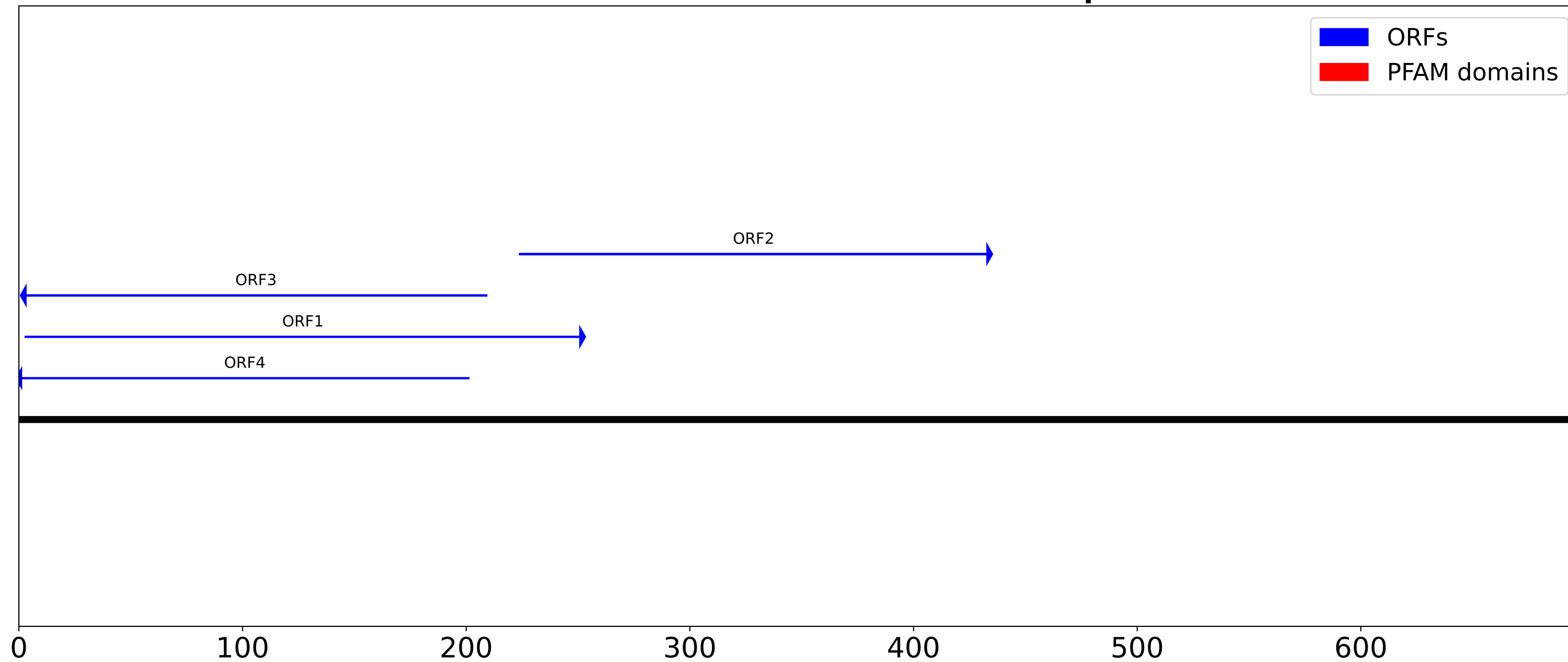
TE: rnd_5_family_2161
size: 638bp; fragments: 454; full length: 45 (≥ 574.2 bp)



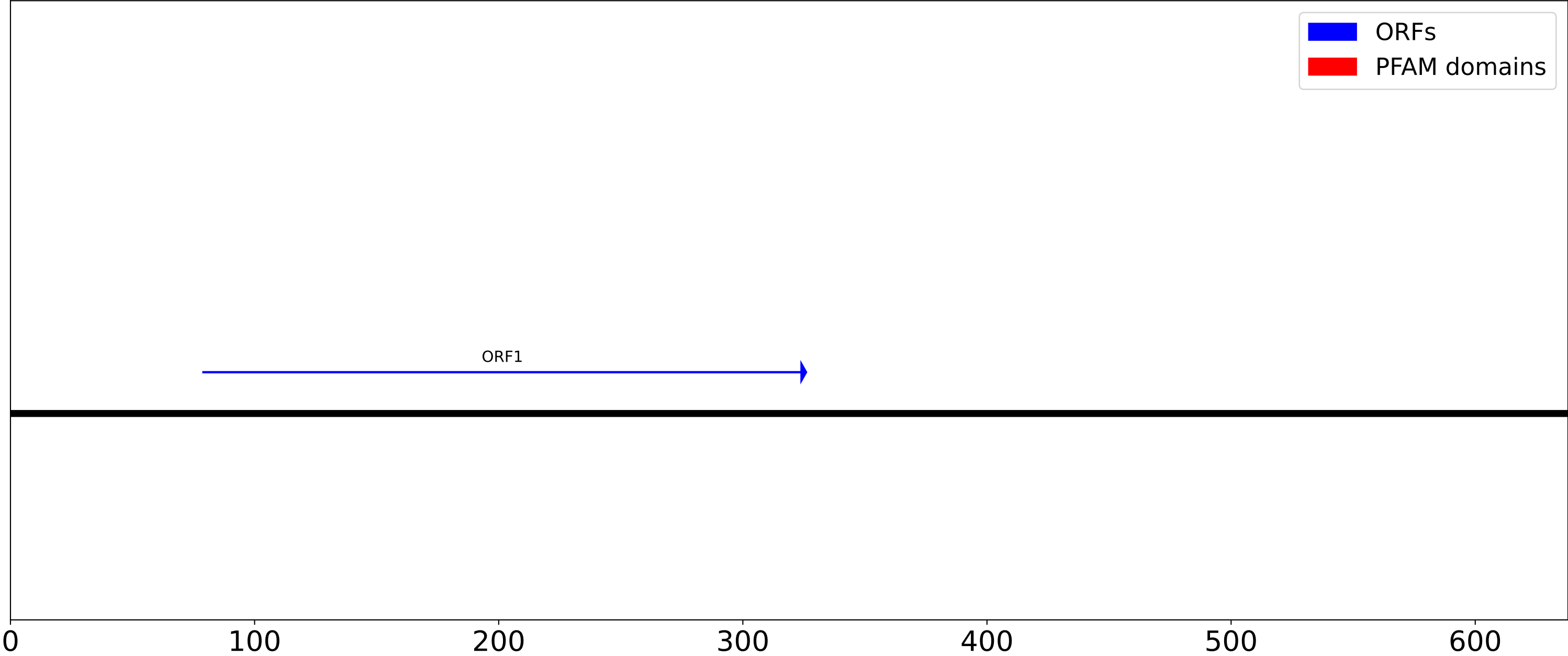
Before TEtrimmer 638 bp



After TEtrimmer ORF and PFAM domain plot



Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

