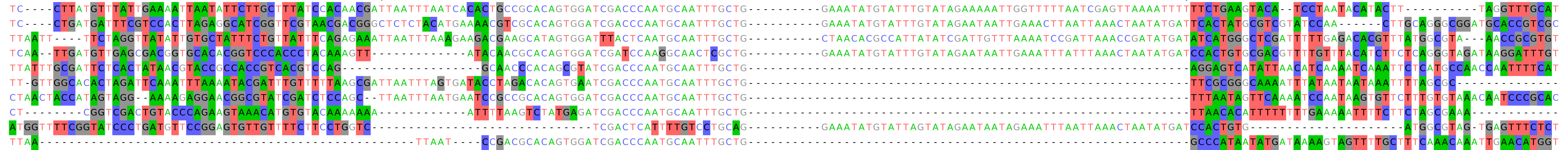


Start Crop Point

End Crop Point

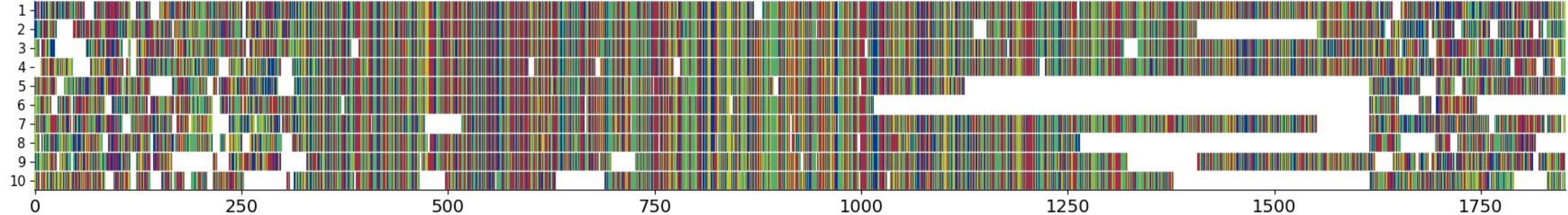
MSA length = 1315



Start crop Point

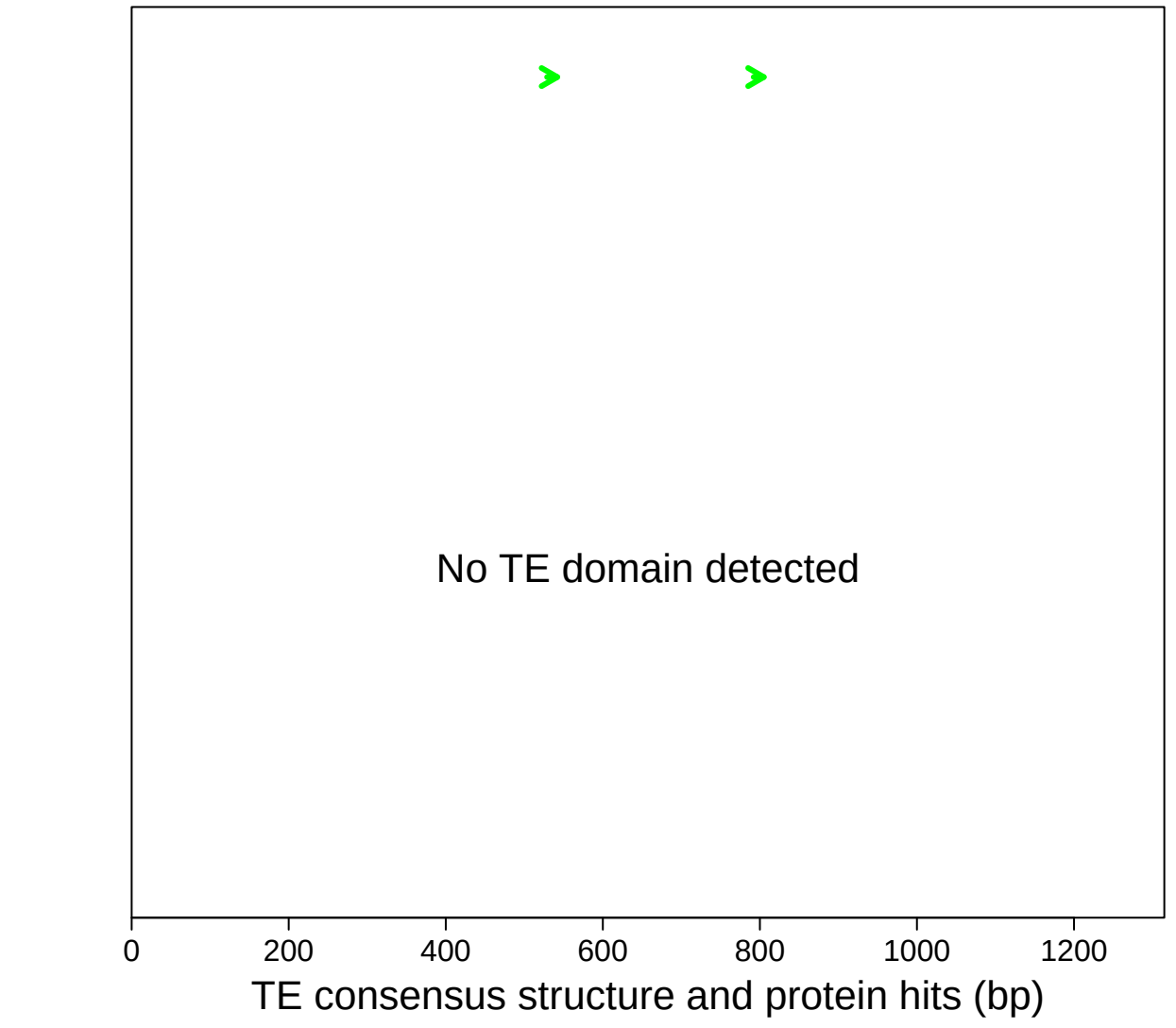
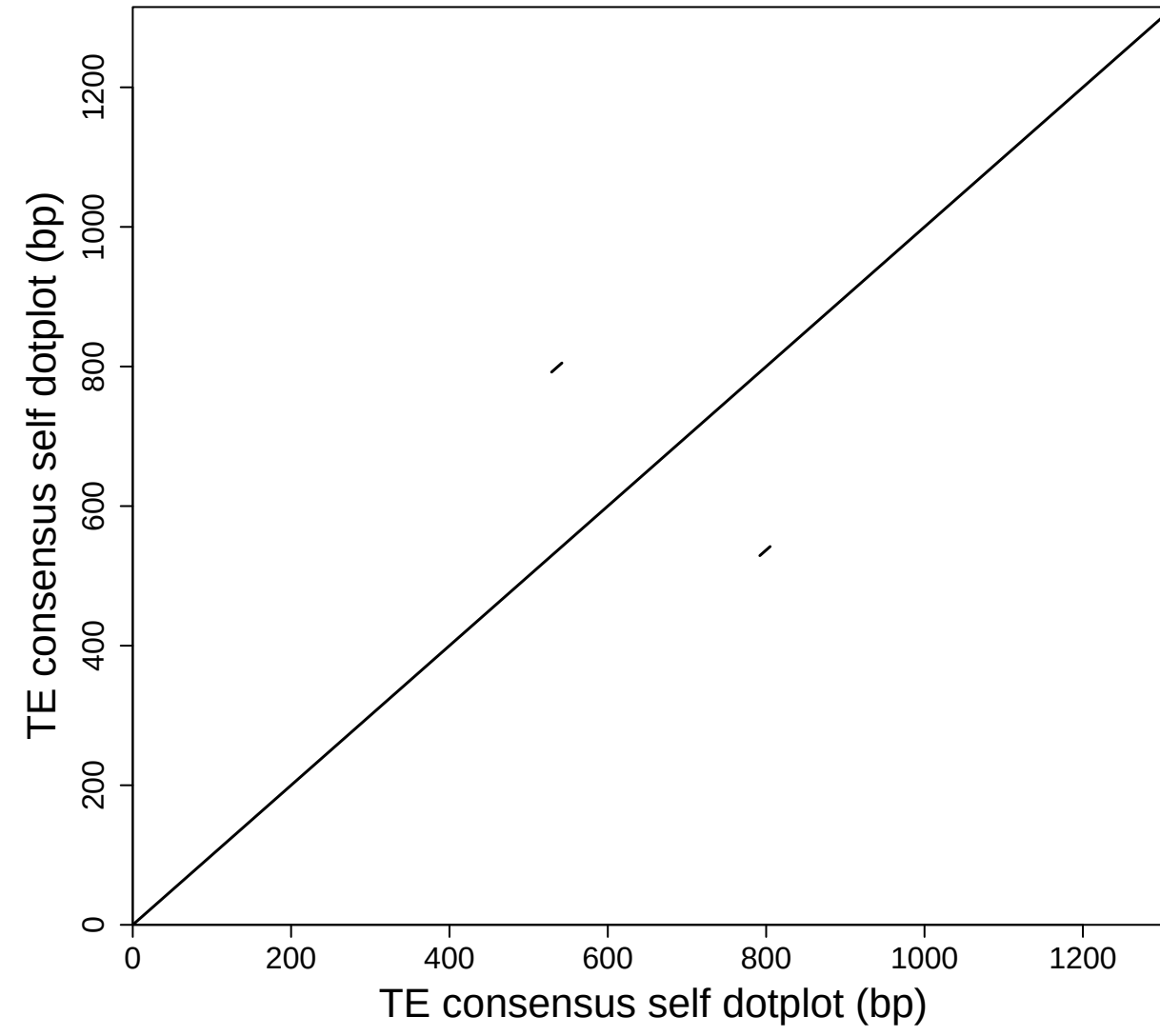
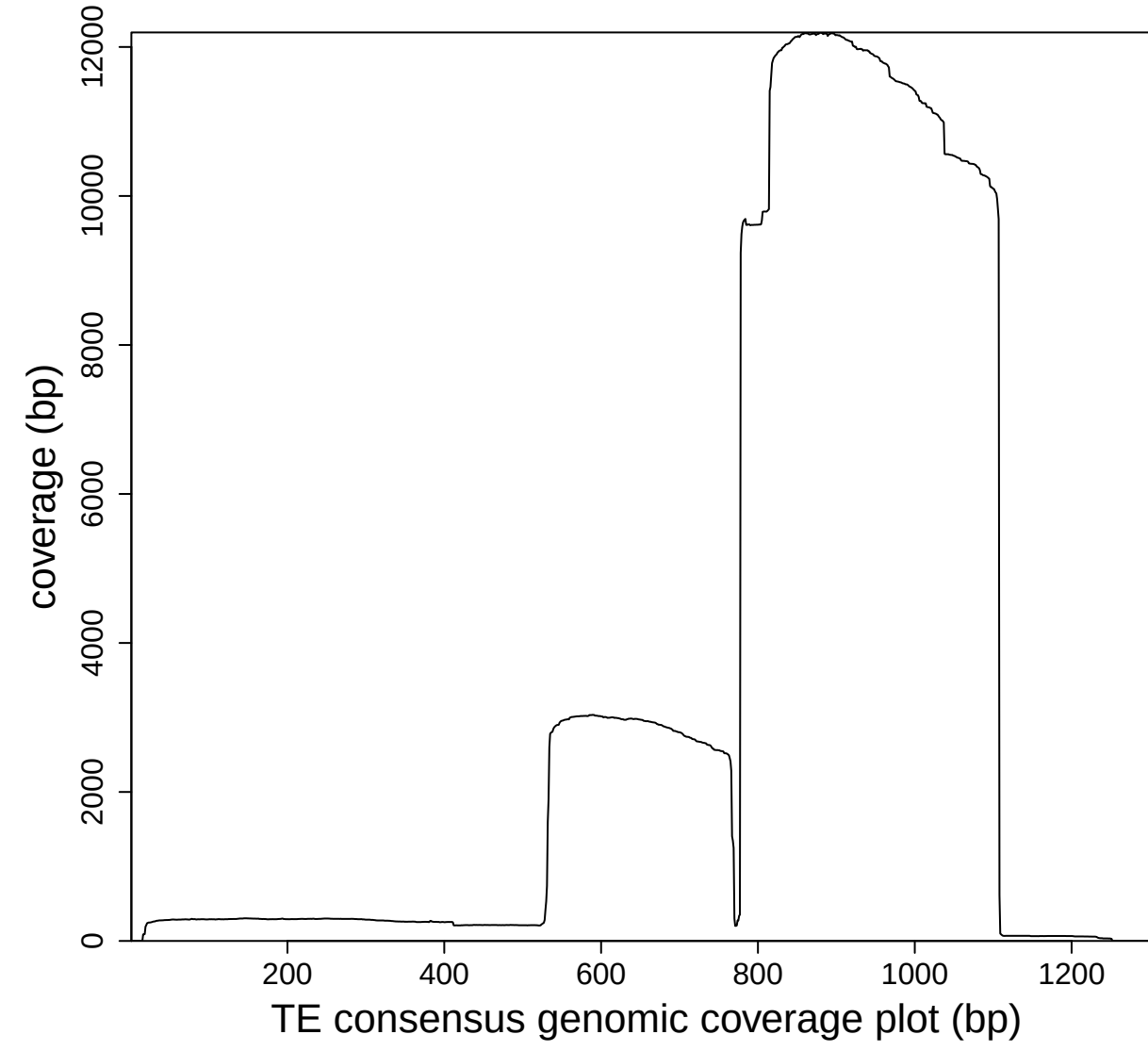
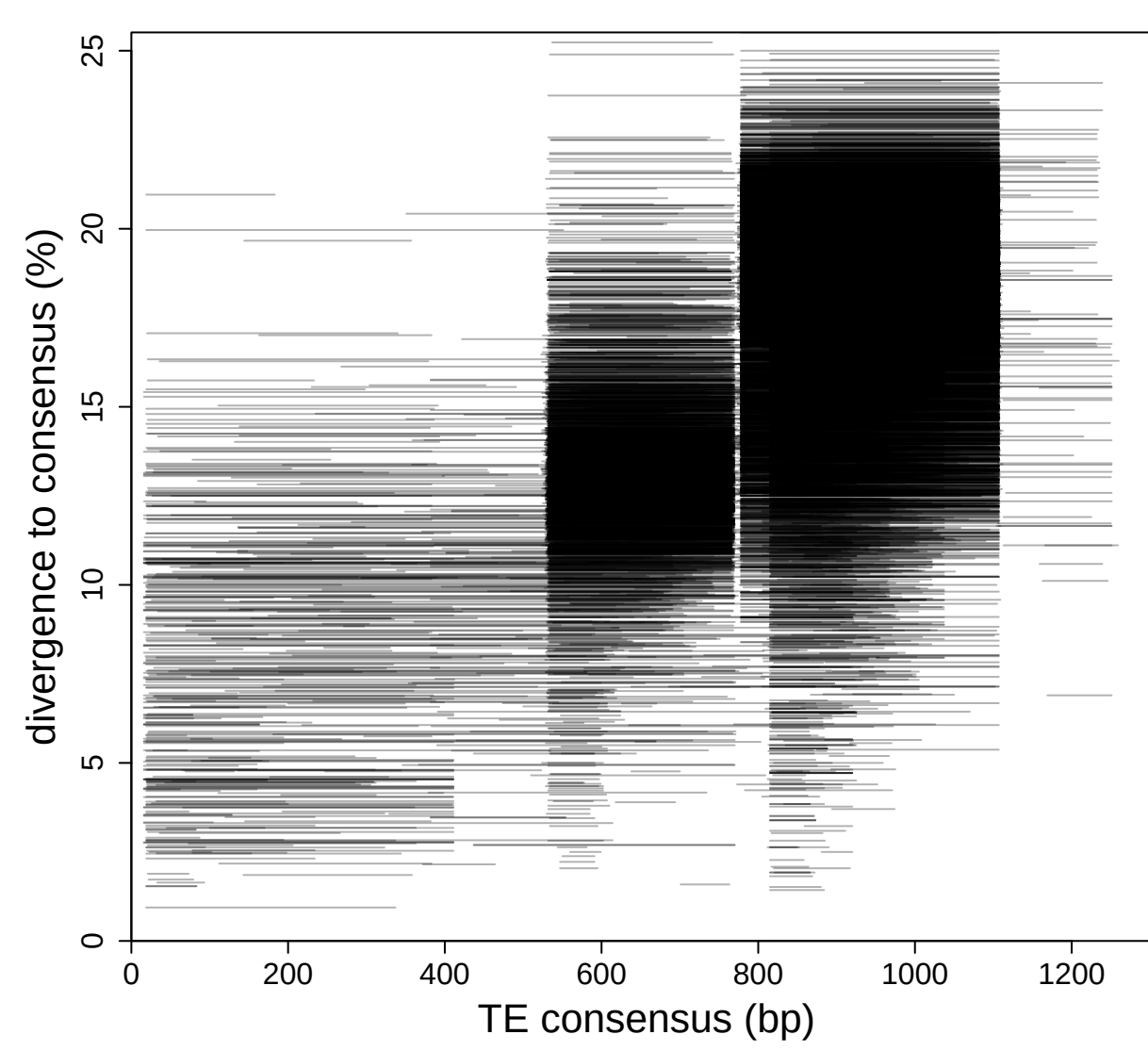


End crop Point



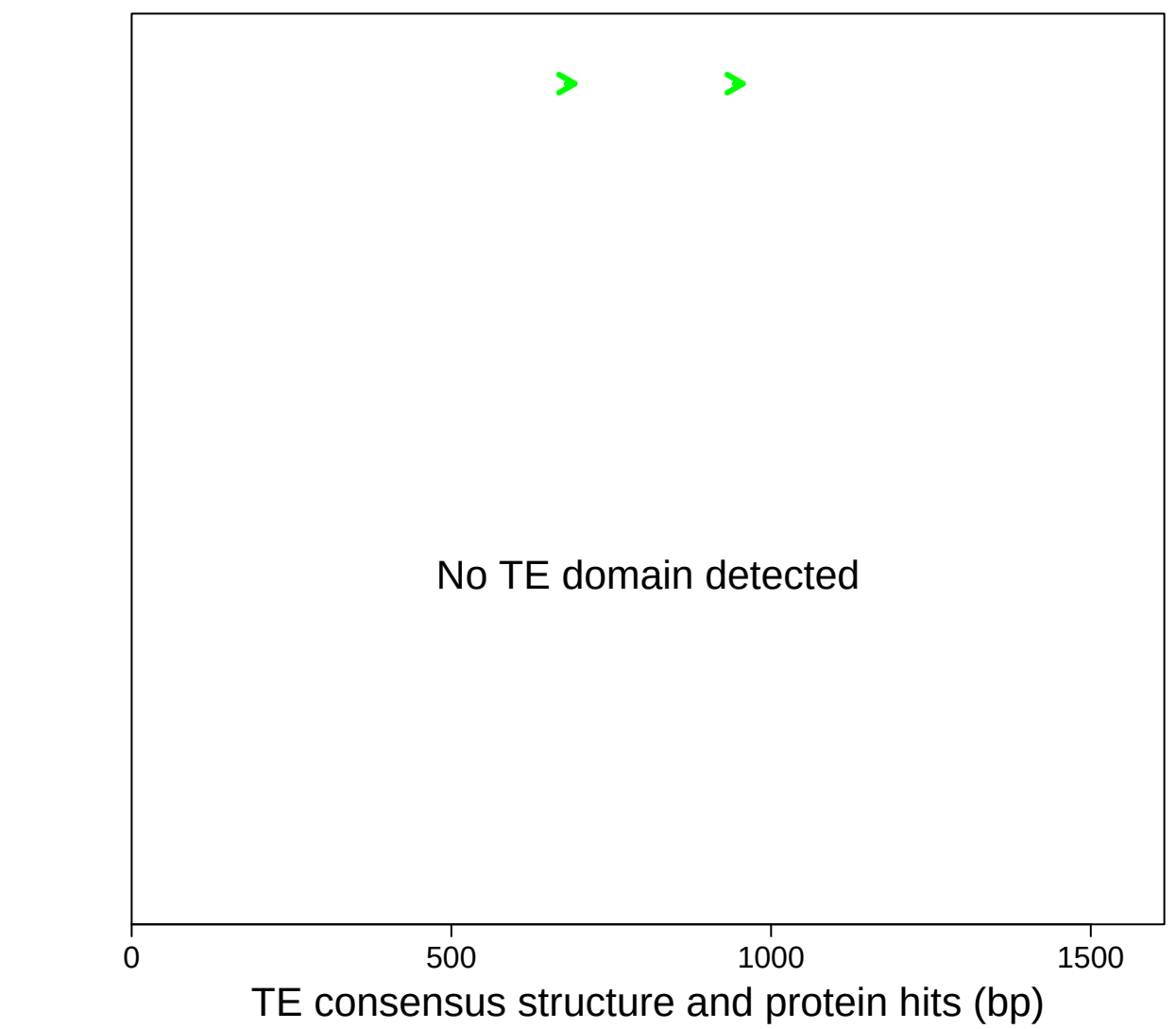
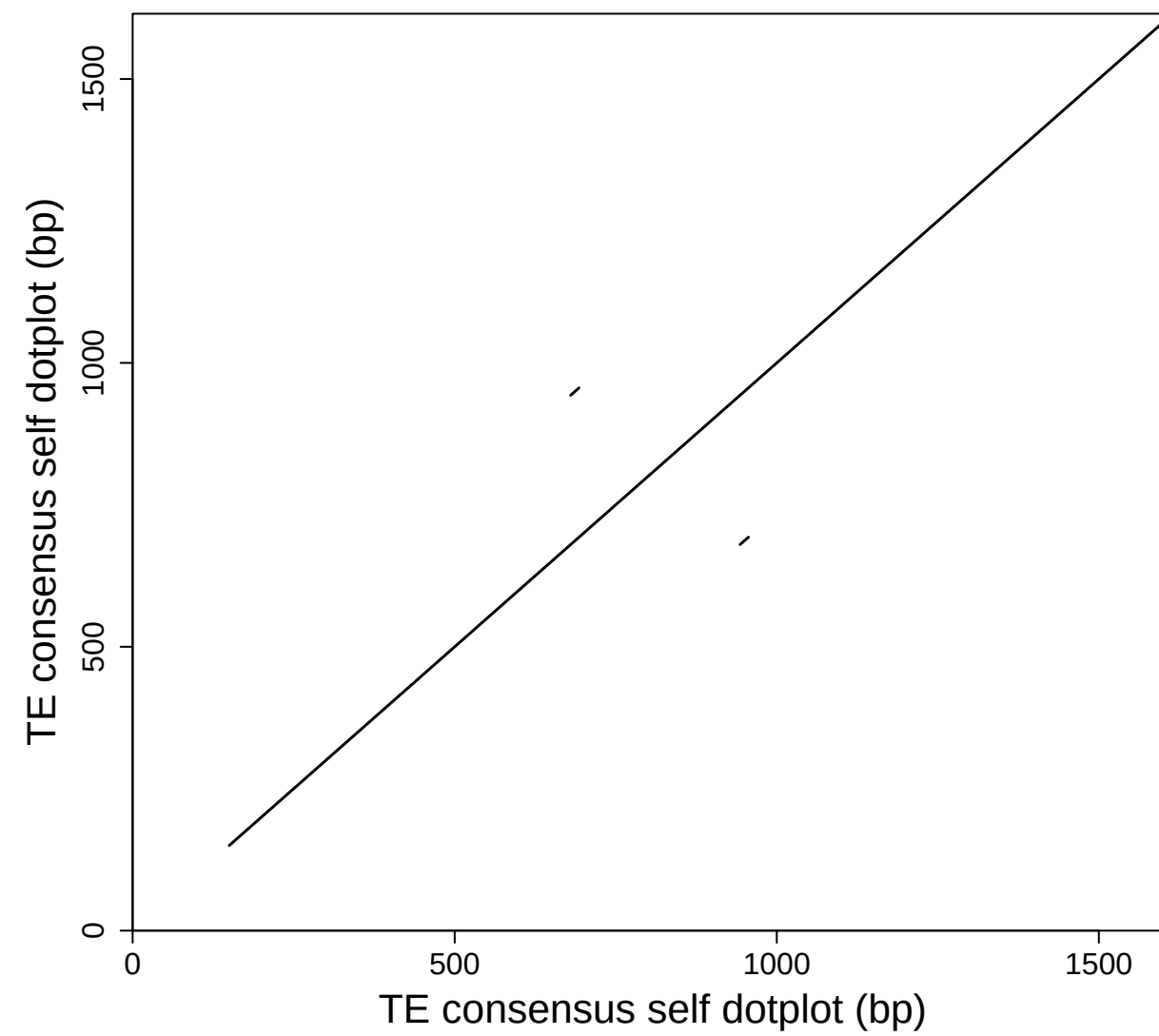
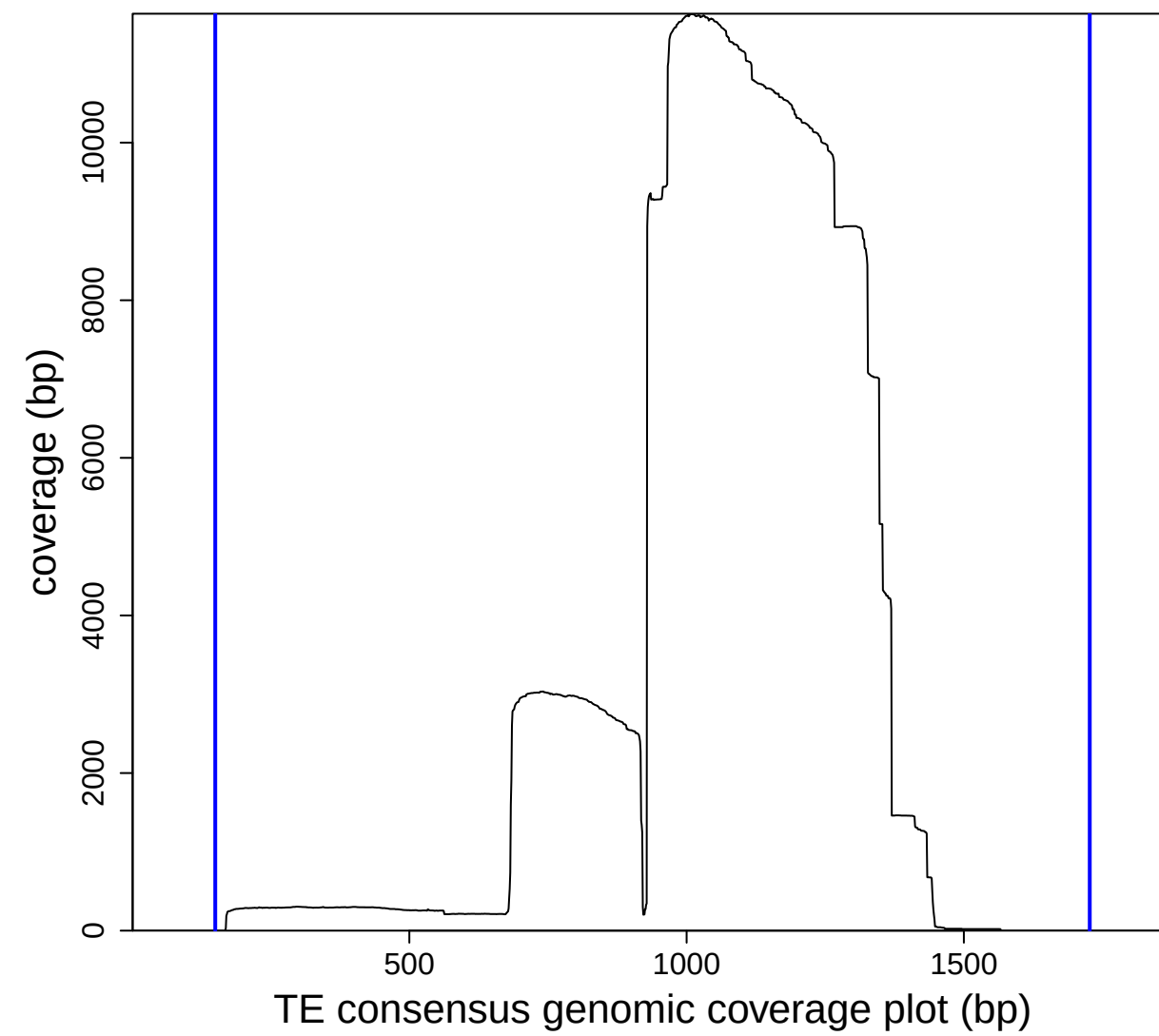
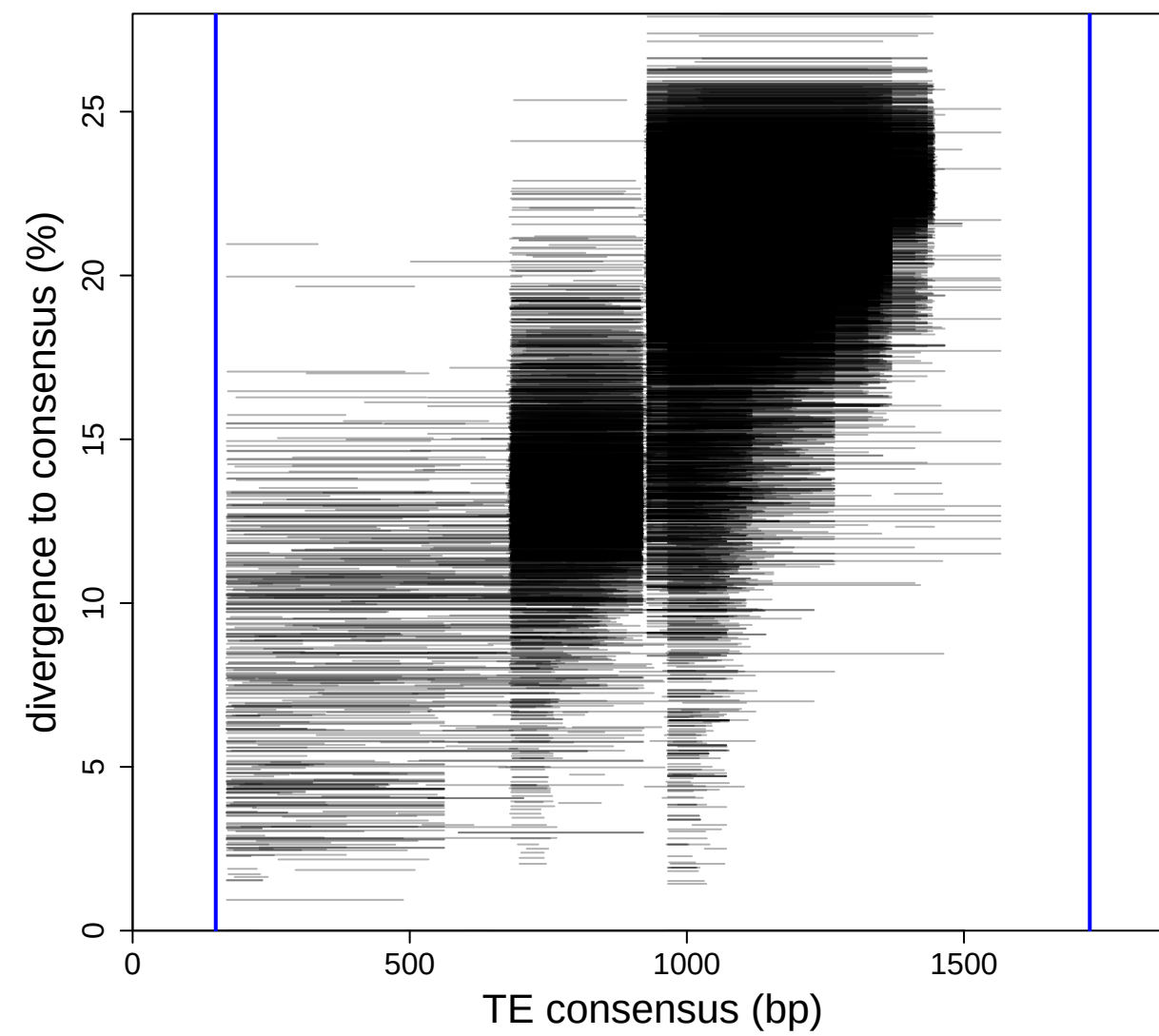
a.b.bed_uf.bed_g_3.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.f
size: 1315bp; fragments: 16014; full length: 0 (>=1183.5bp)

After TEtrimmer 1315 bp



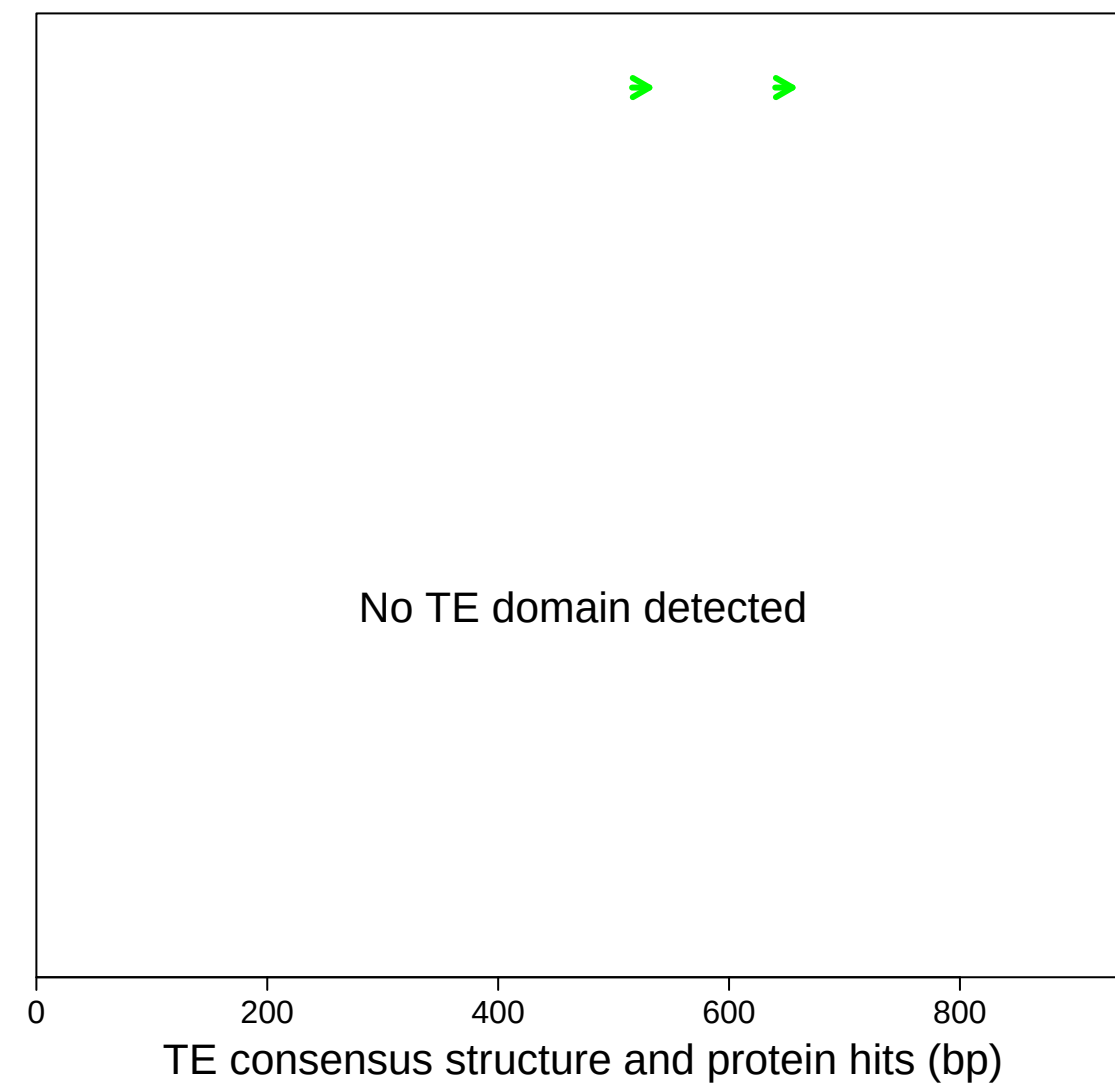
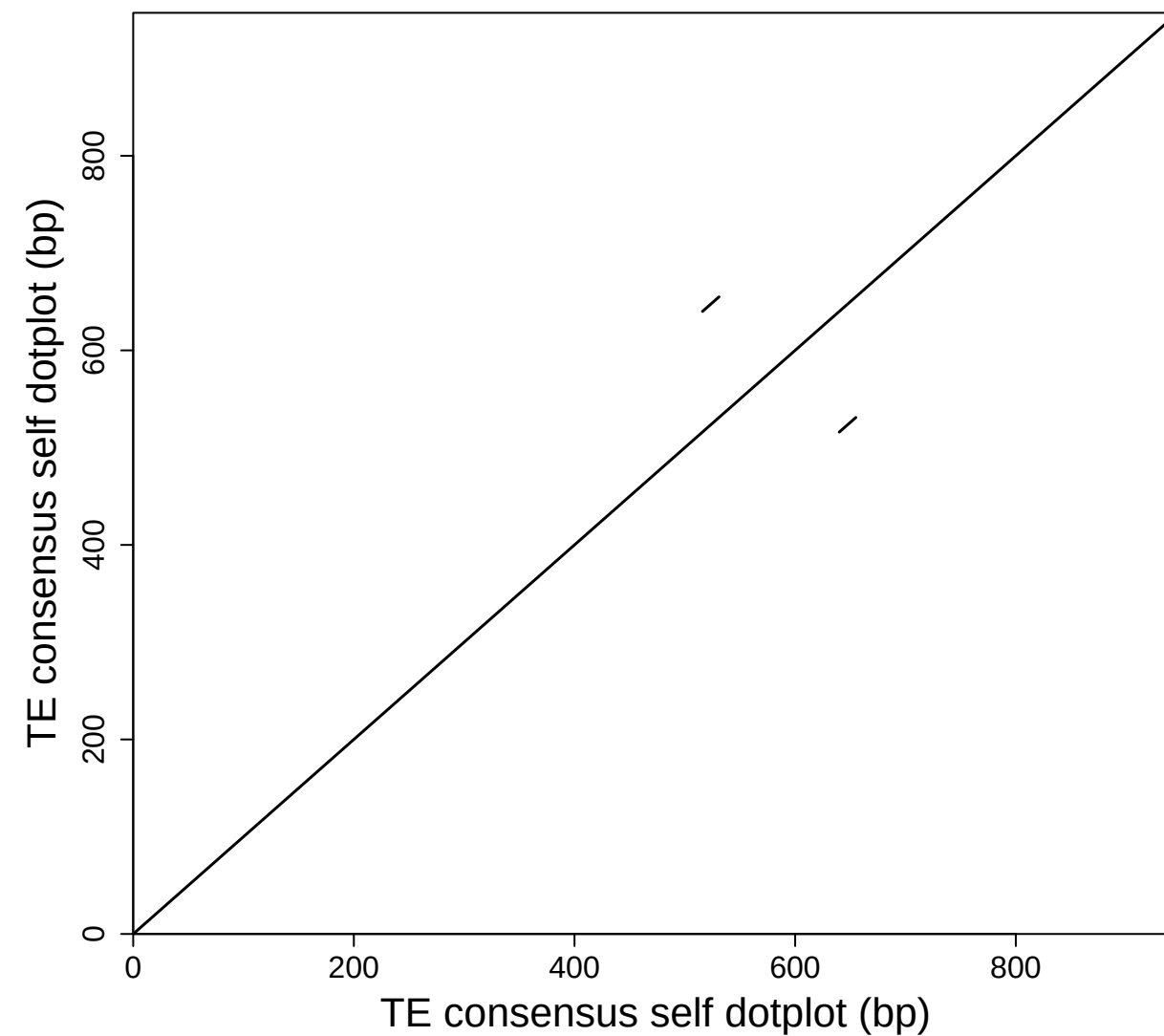
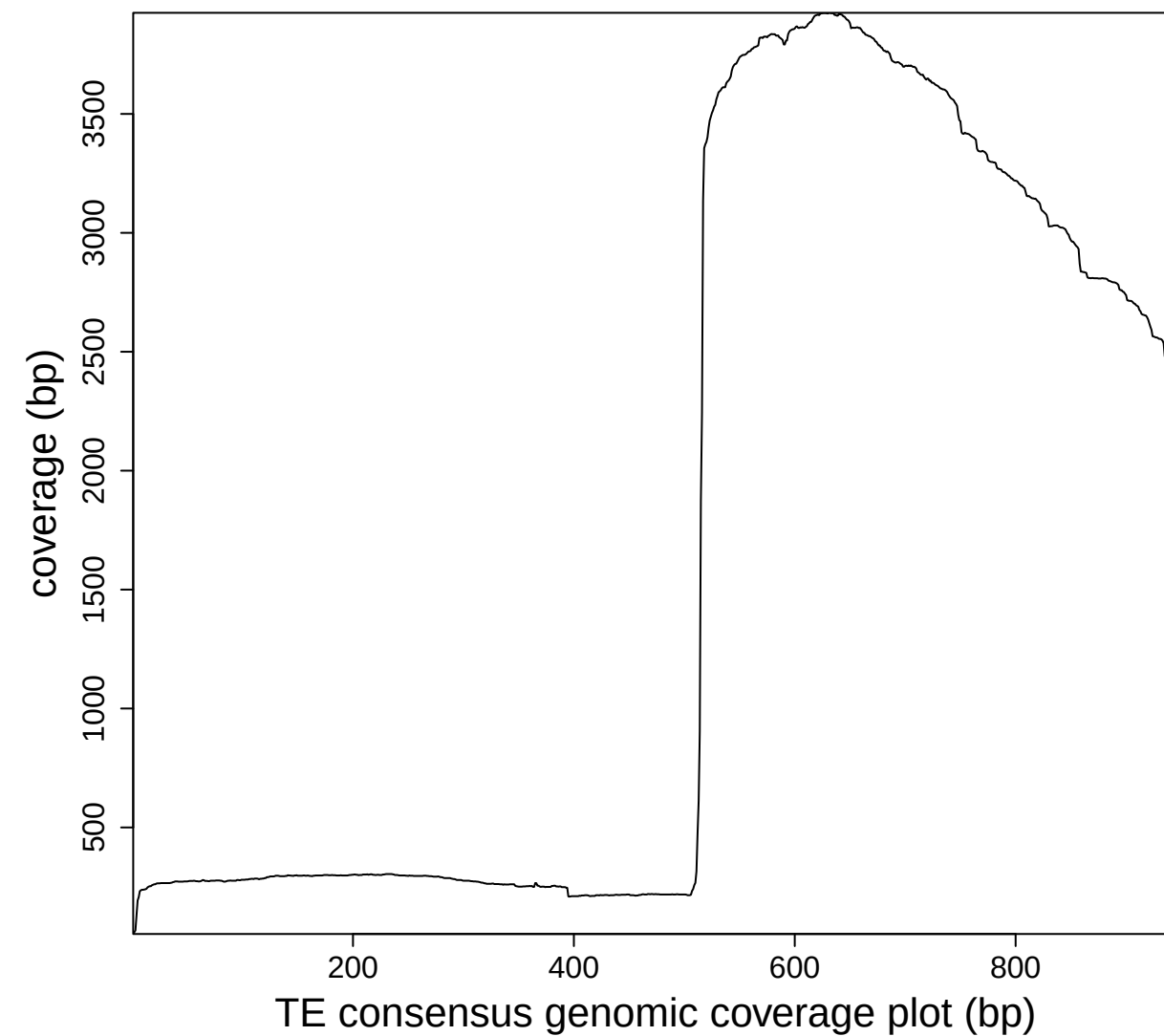
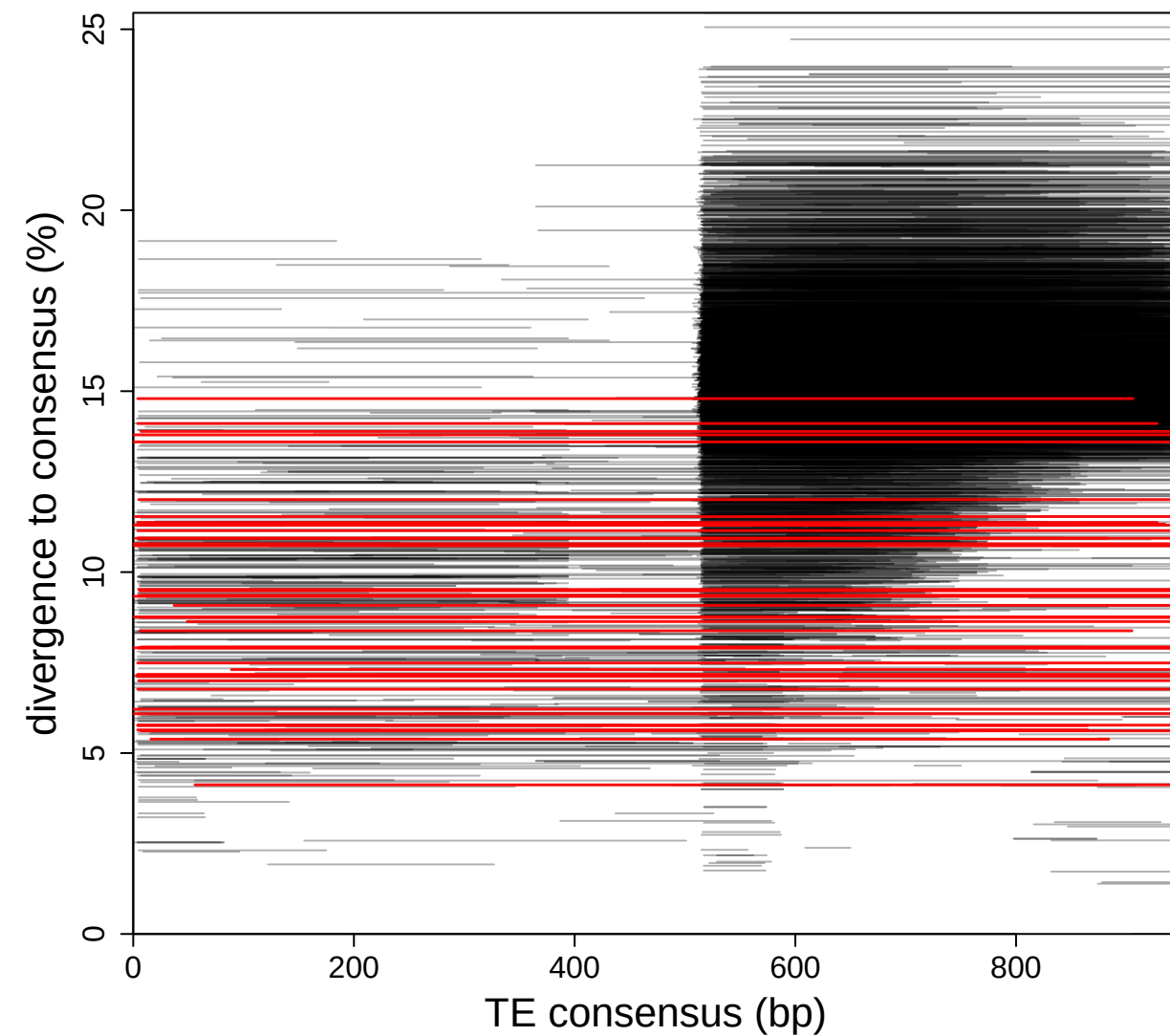
1.fasta.b.bed uf.bed g_3.bed fm_1.bed 0 0 bcln.fa aln.fa cl.f
size: 1877bp; fragments: 15294; full length: 0 (>=1689.3bp)

After TEtrimmer Extended plot Blue lines are boundaries

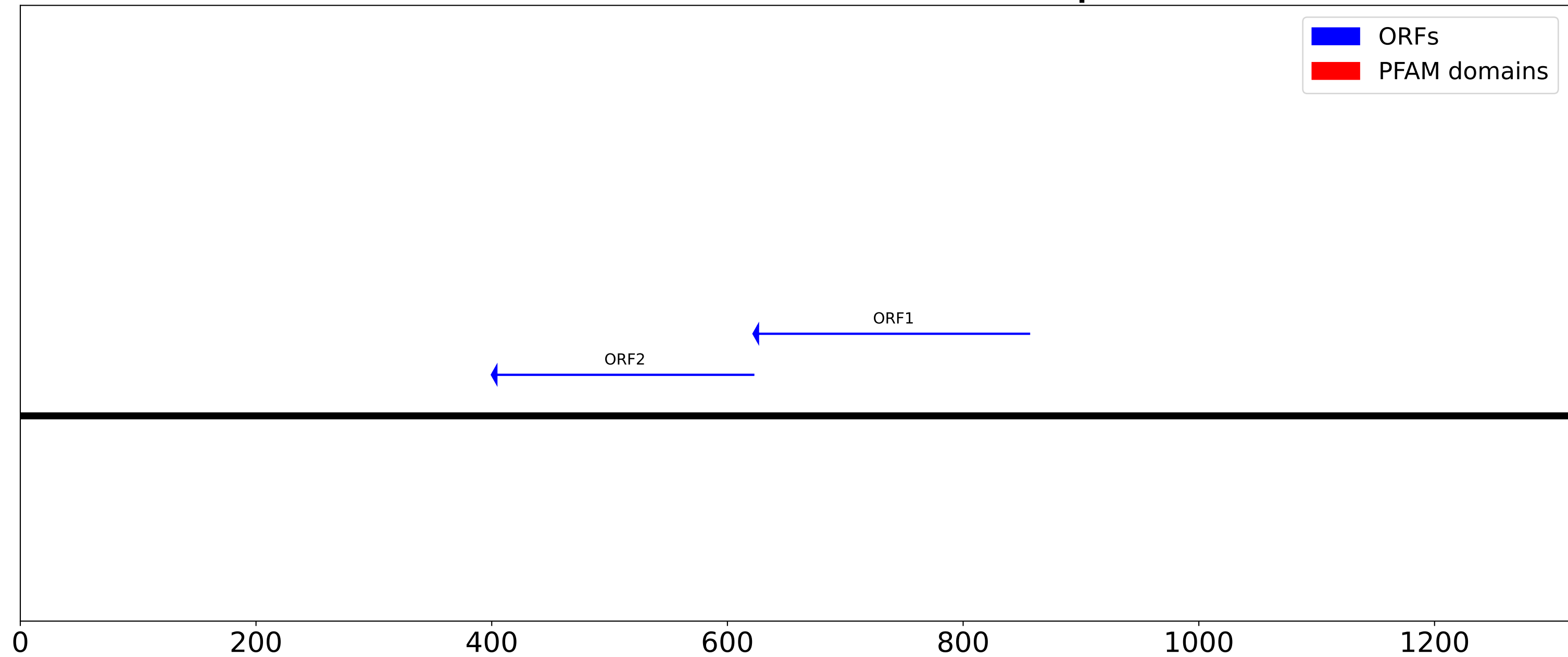


TE: rnd_5_family_4651
size: 947bp; fragments: 4551; full length: 44 (≥ 852.3 bp)

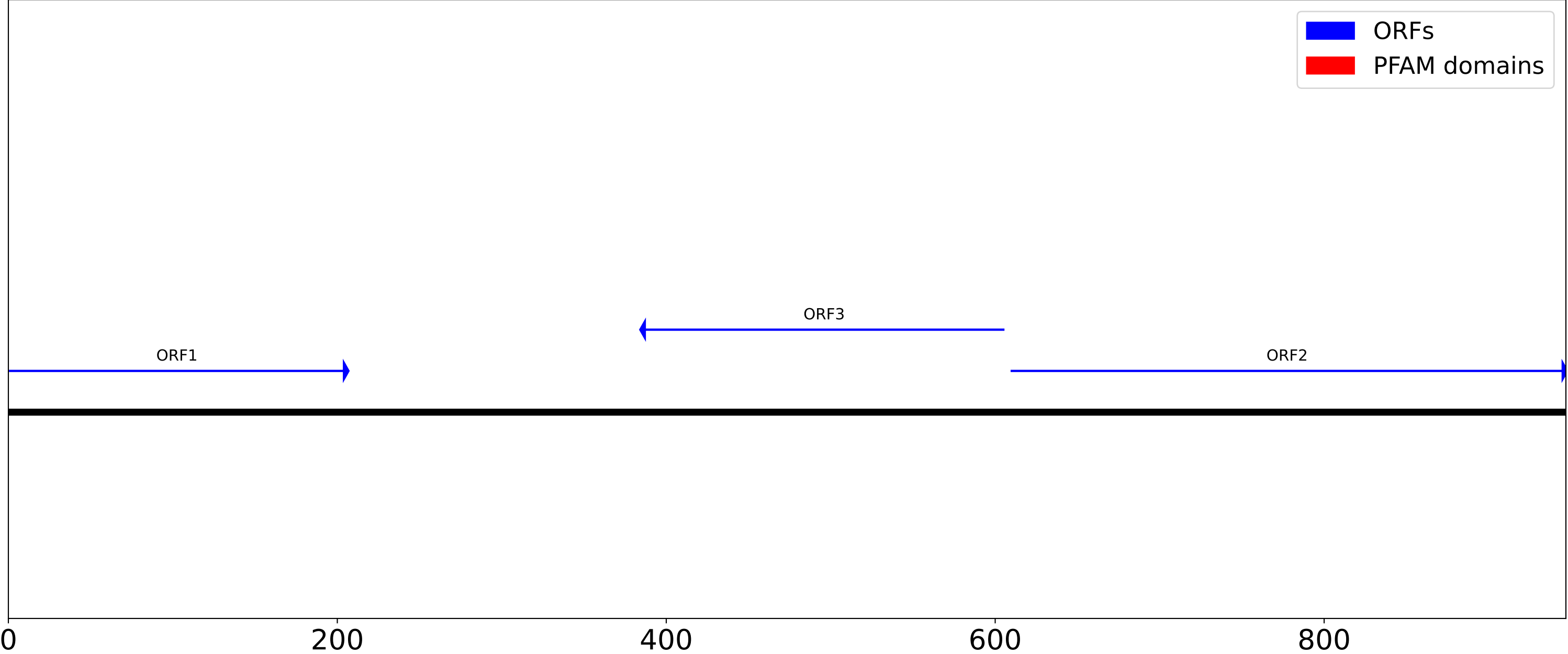
Before TEtrimmer 947 bp



After TEtrimmer ORF and PFAM domain plot



Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

