

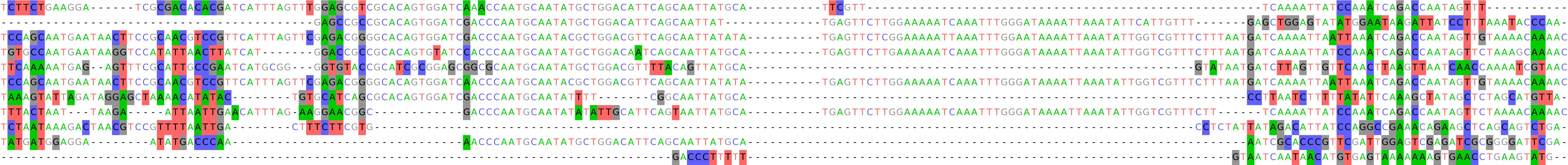
Start crop Point



End crop Point



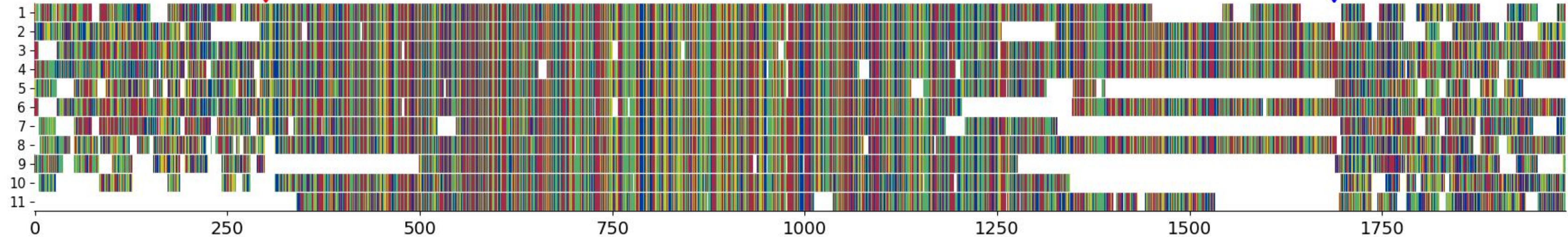
MSA length = 1389



Start crop Point

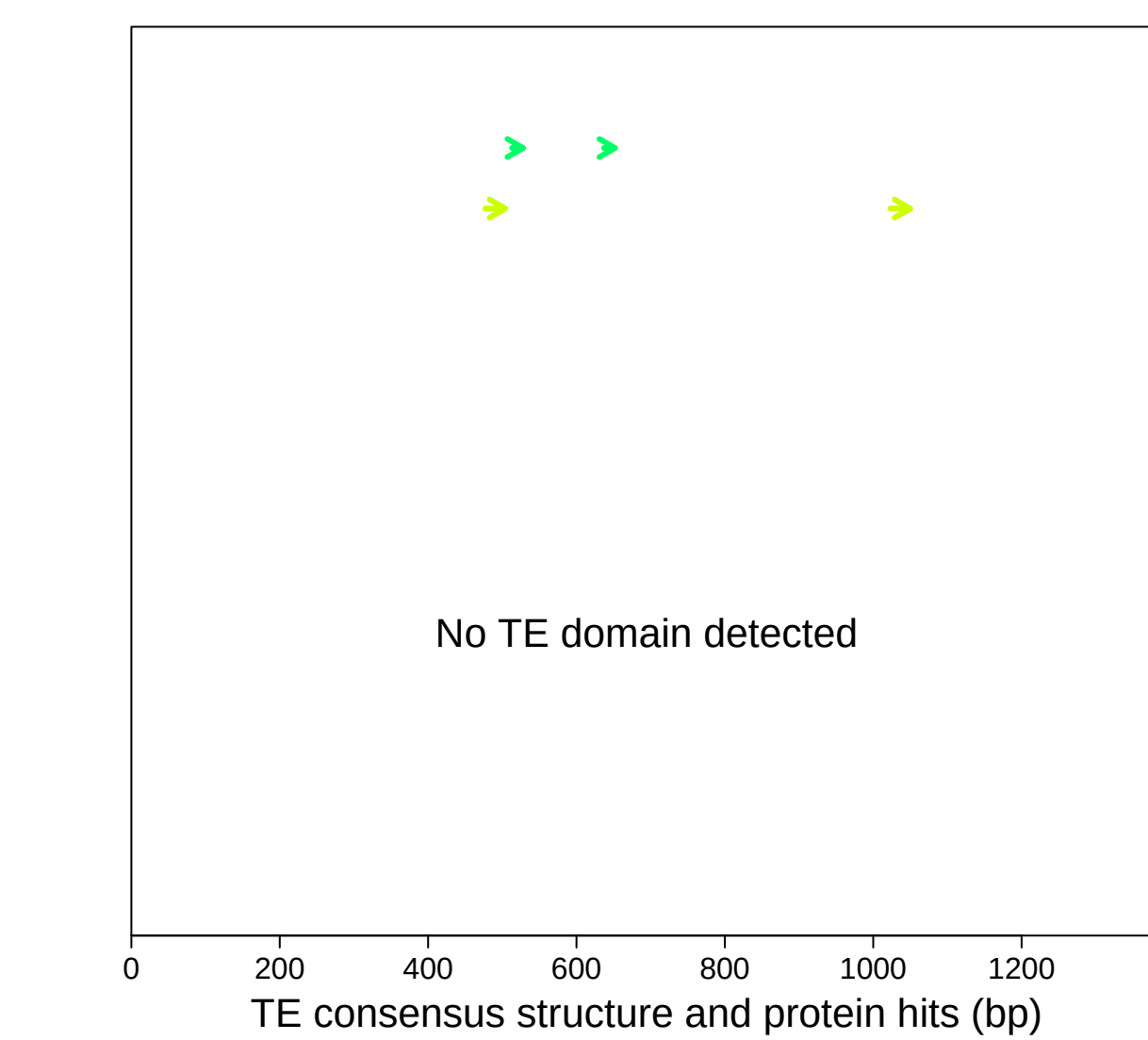
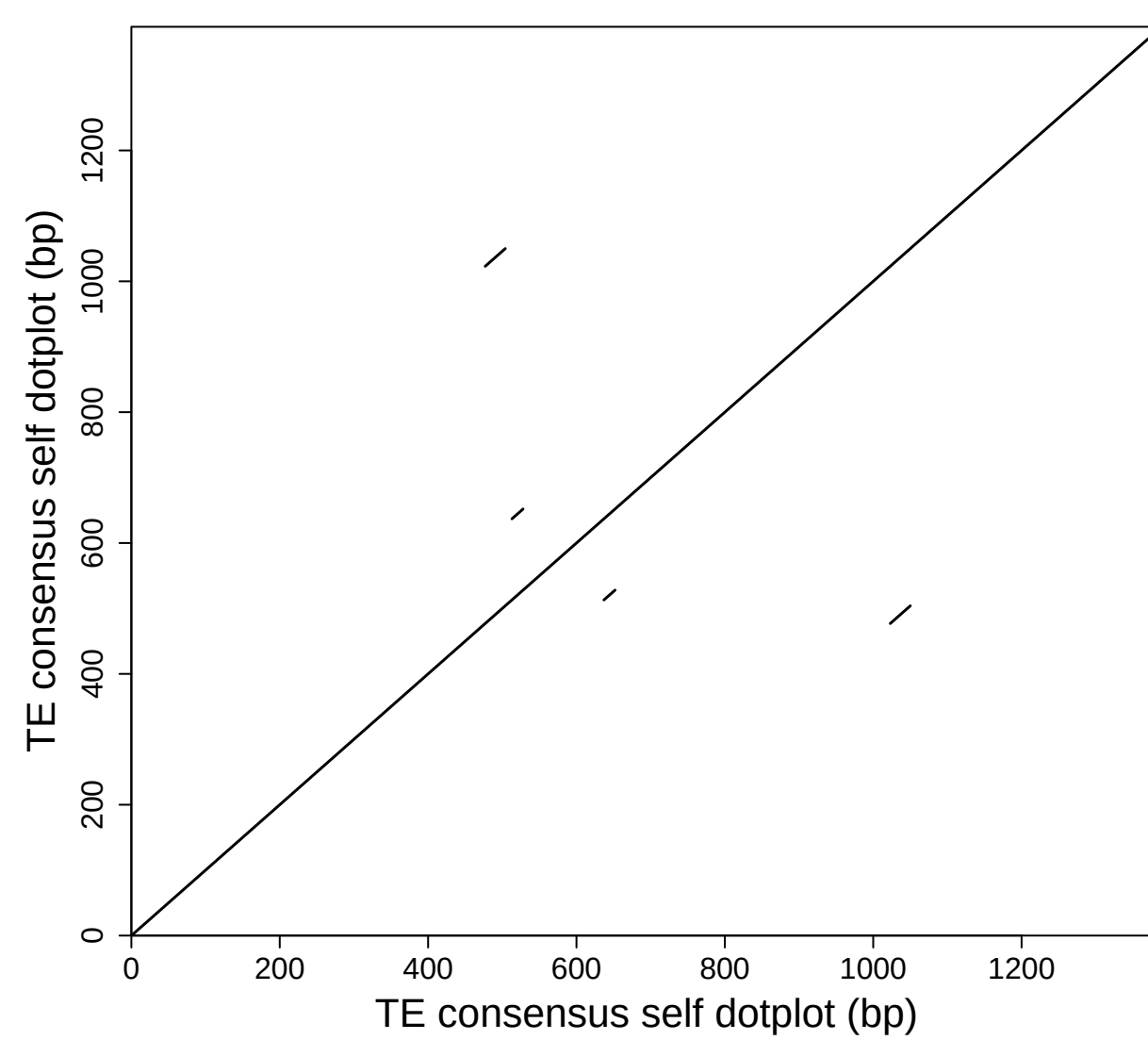
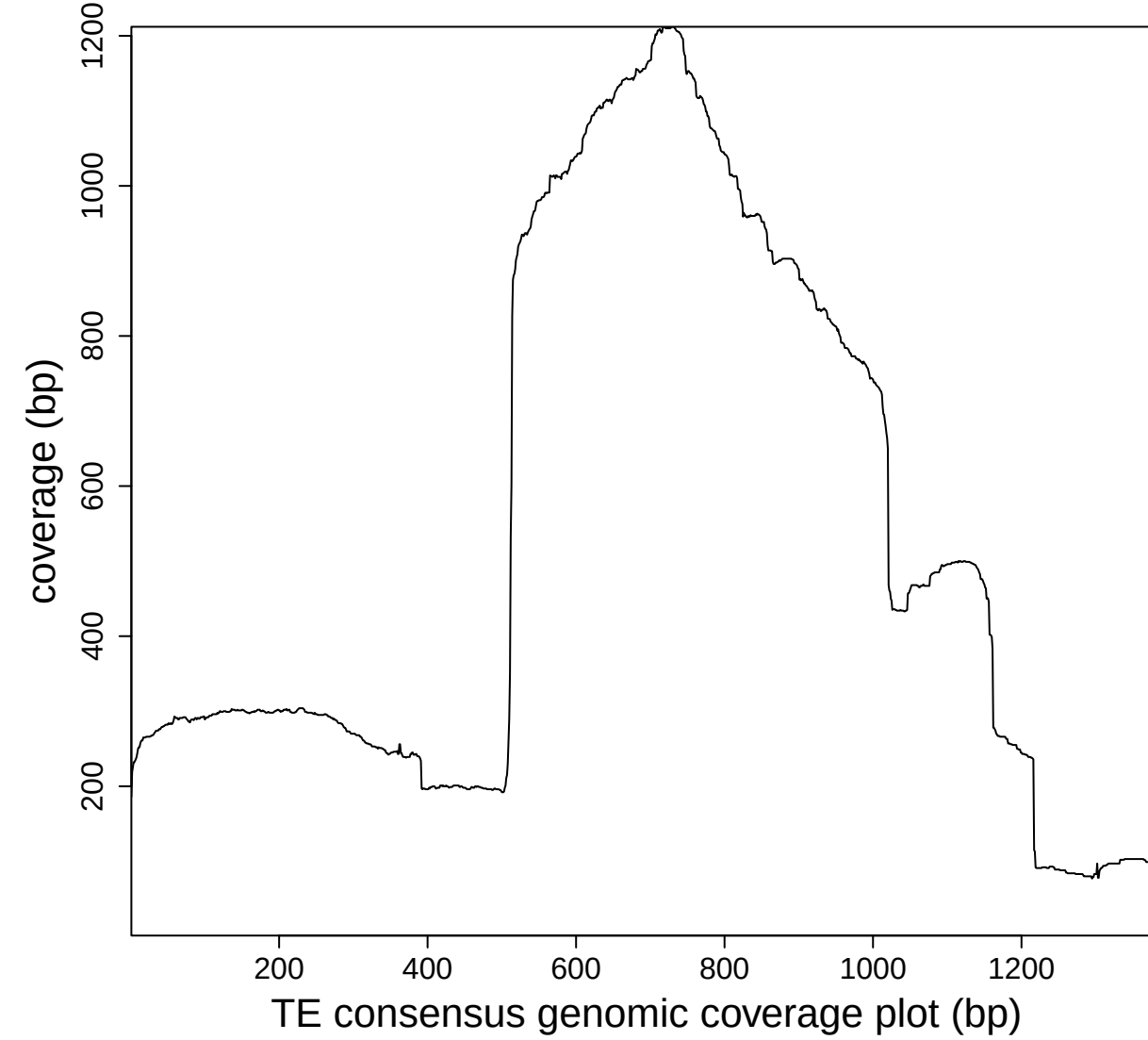
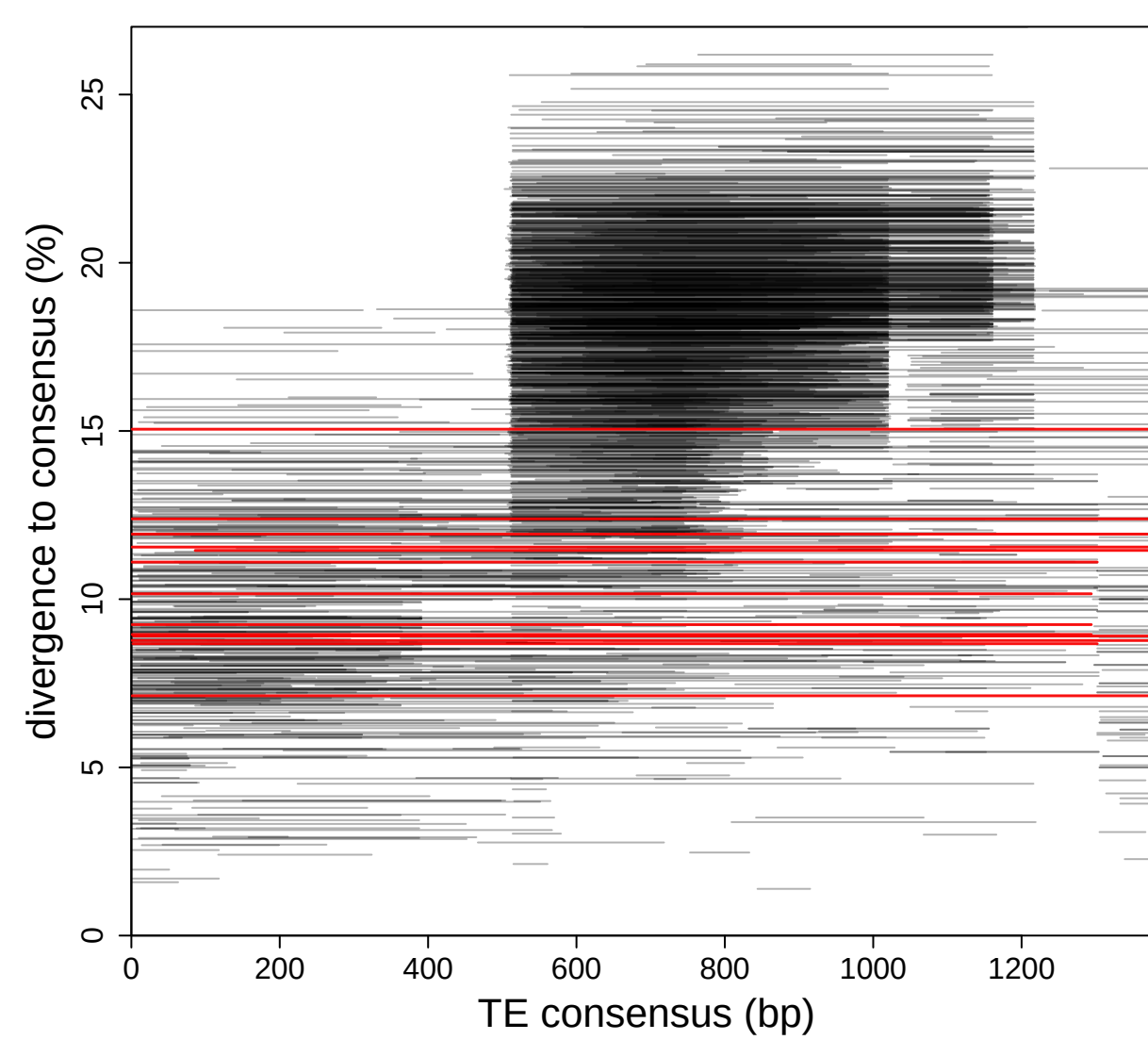


End crop Point



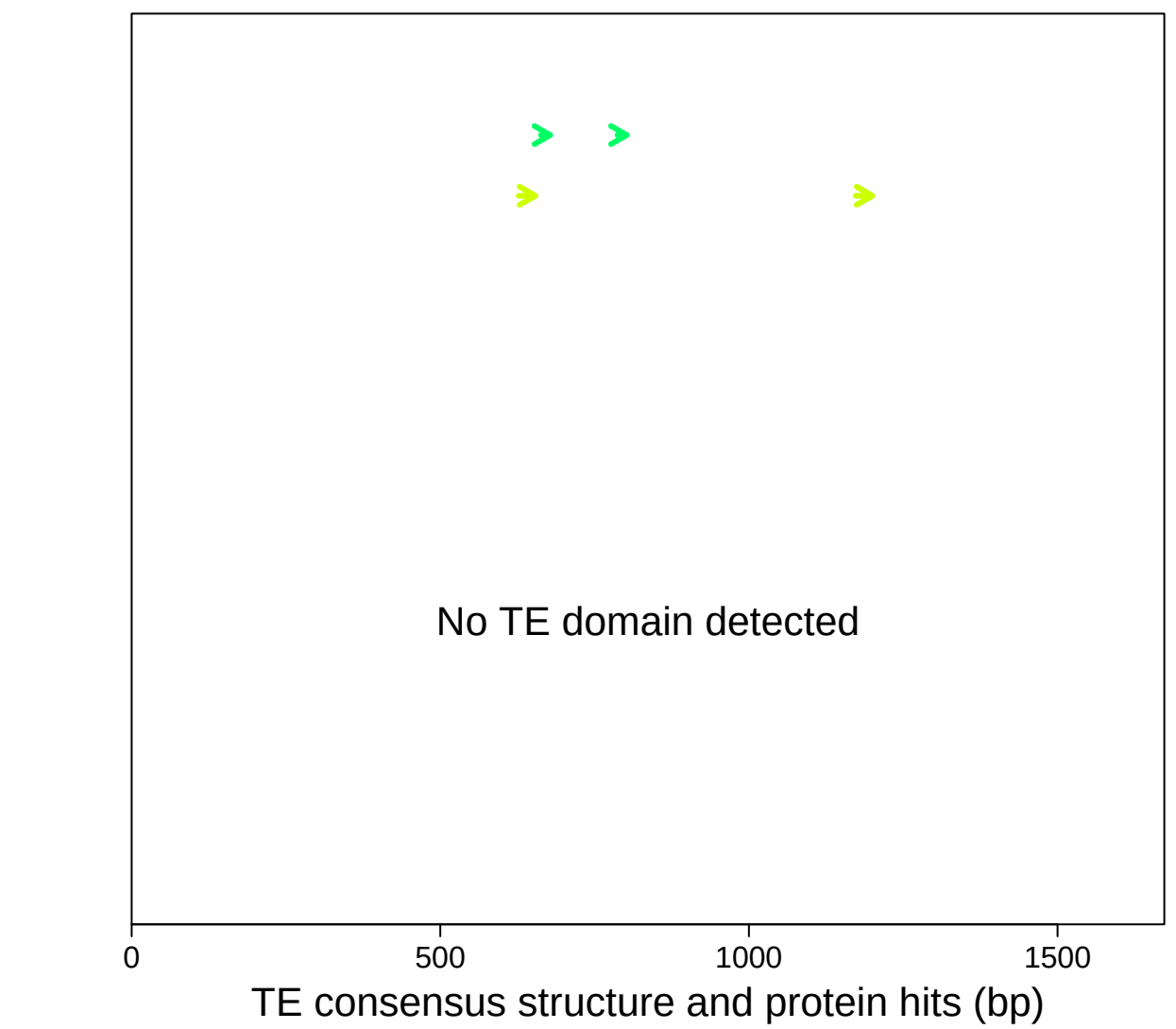
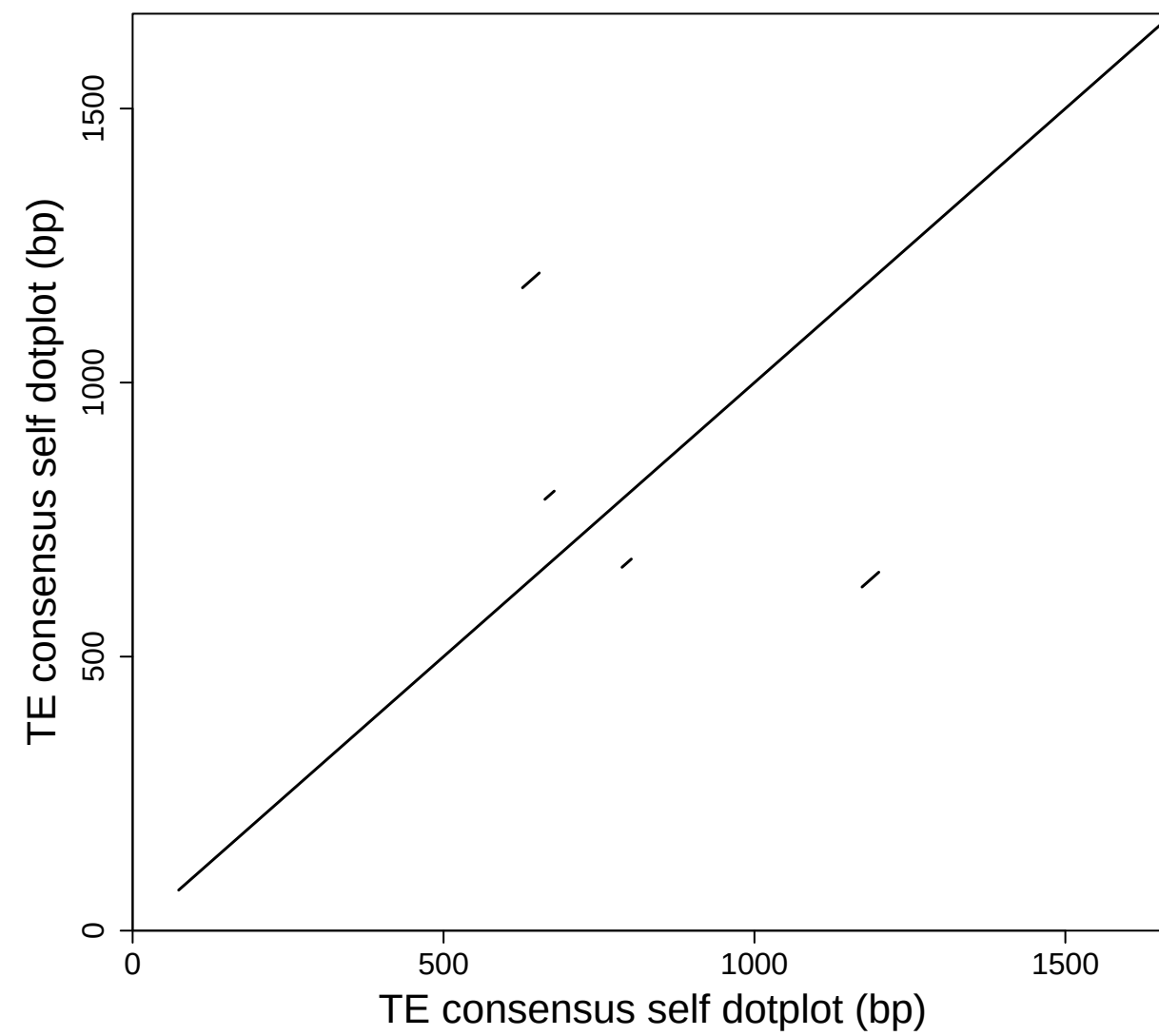
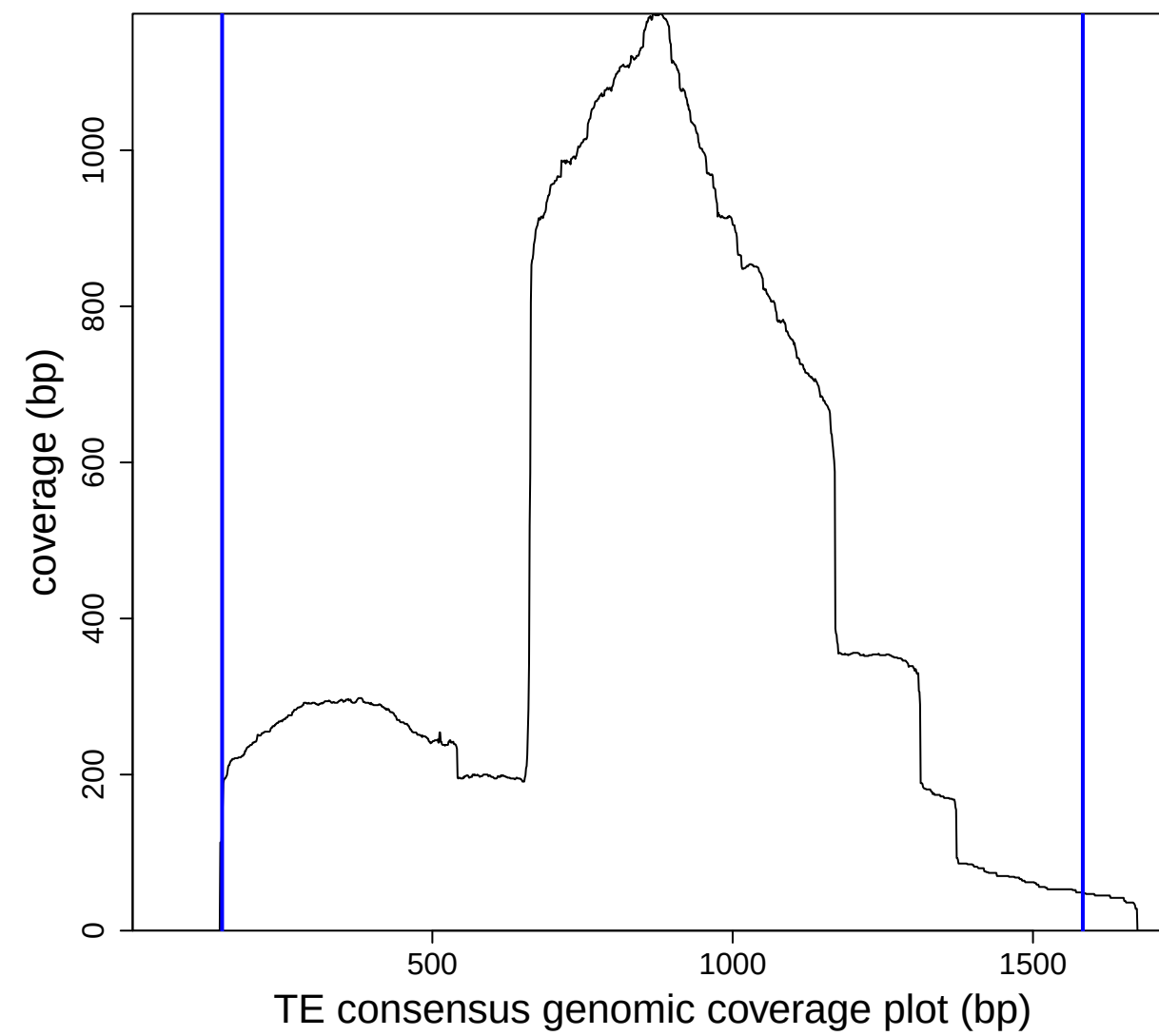
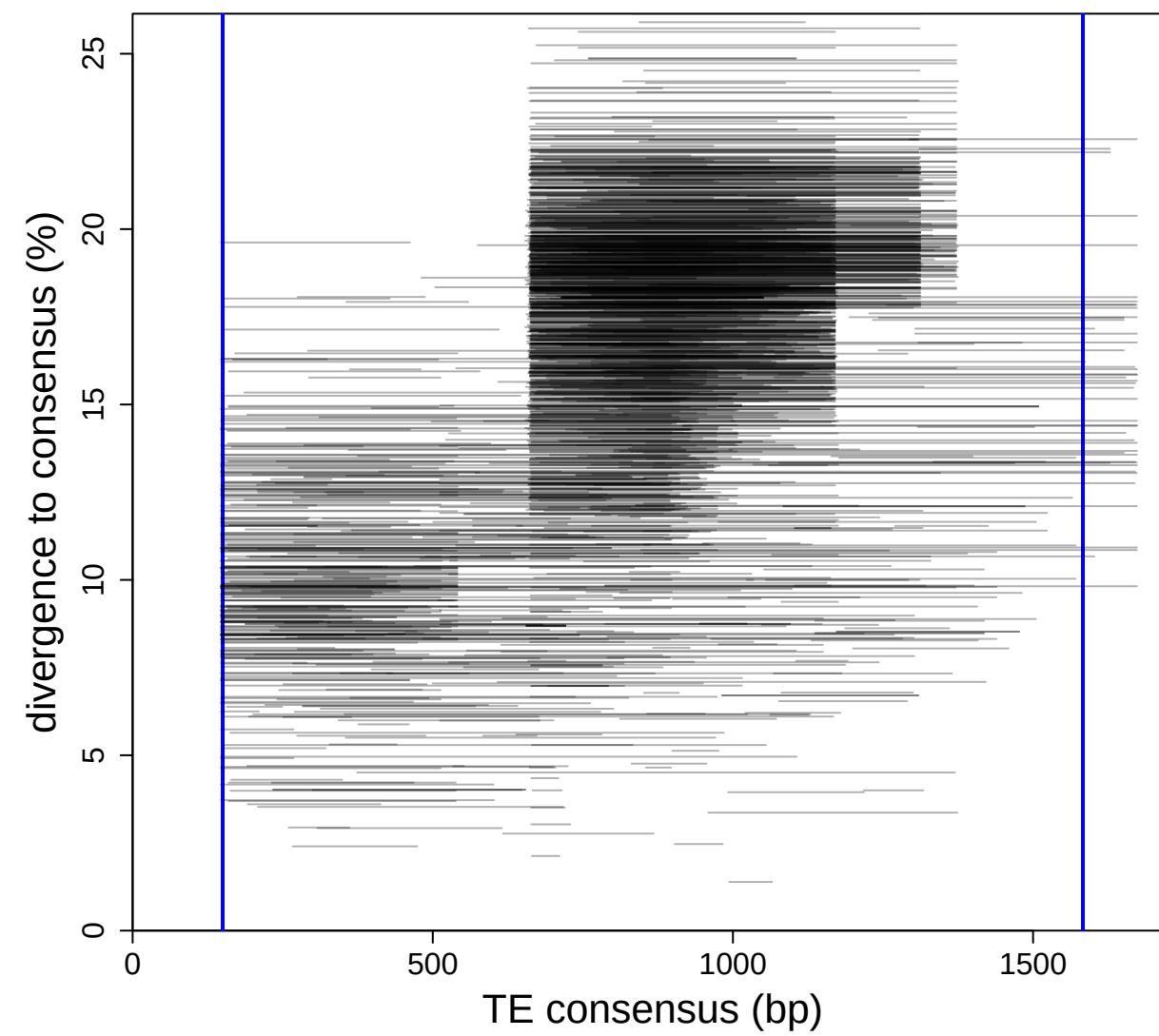
a.b.bed_uf.bed_g_2.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.f
size: 1389bp; fragments: 1822; full length: 13 (>=1250.1bp)

After TEtrimmer 1389 bp



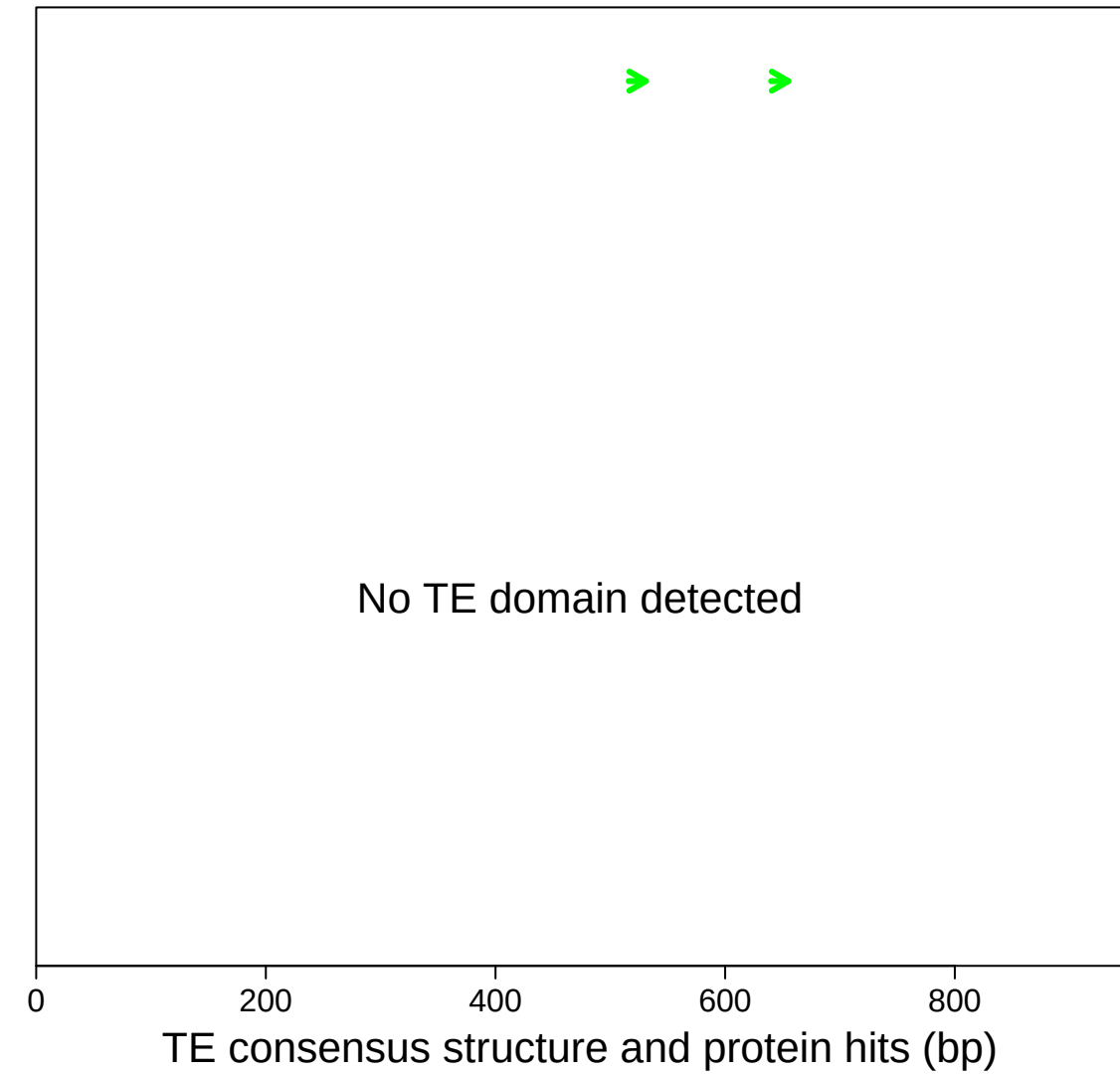
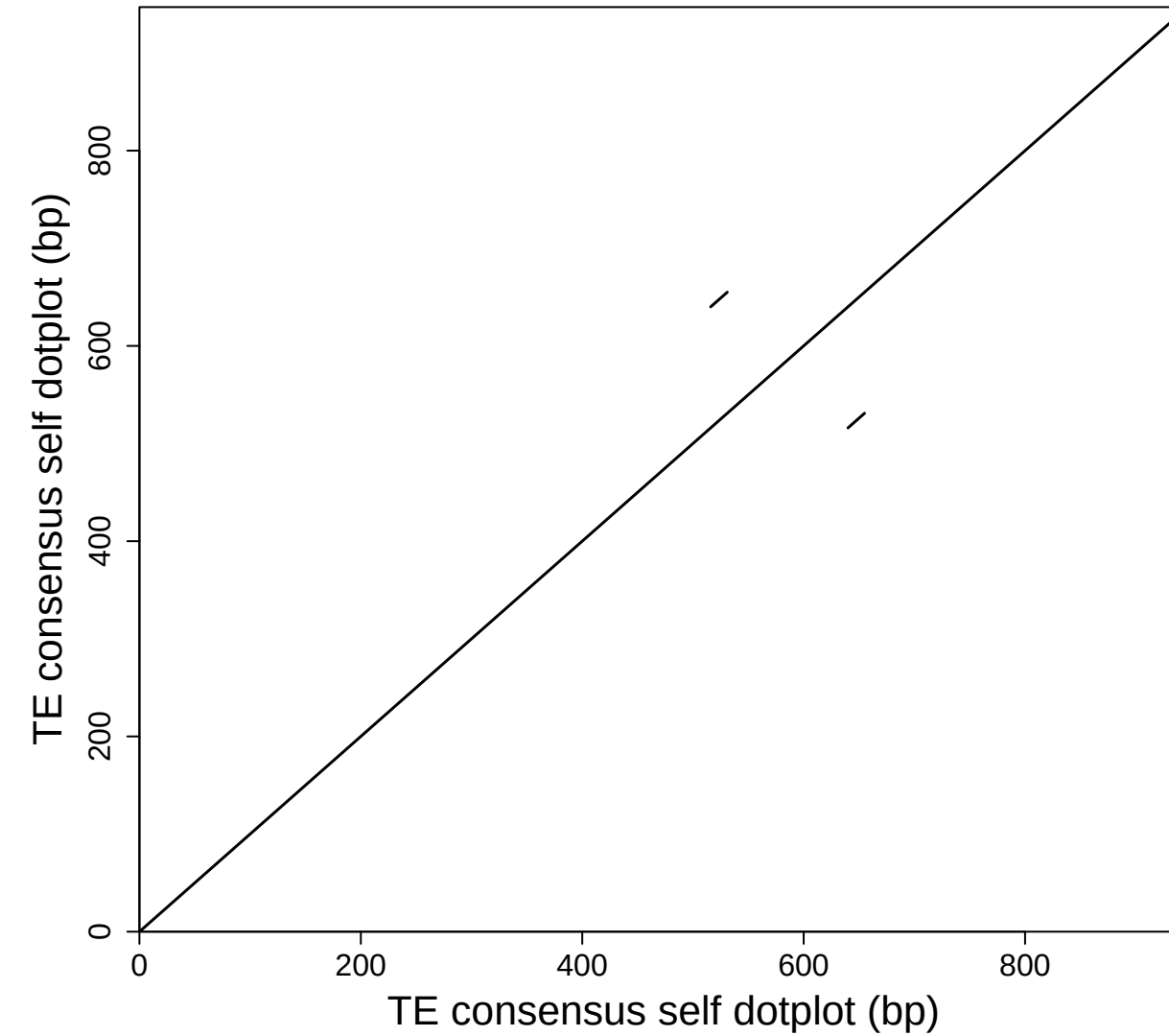
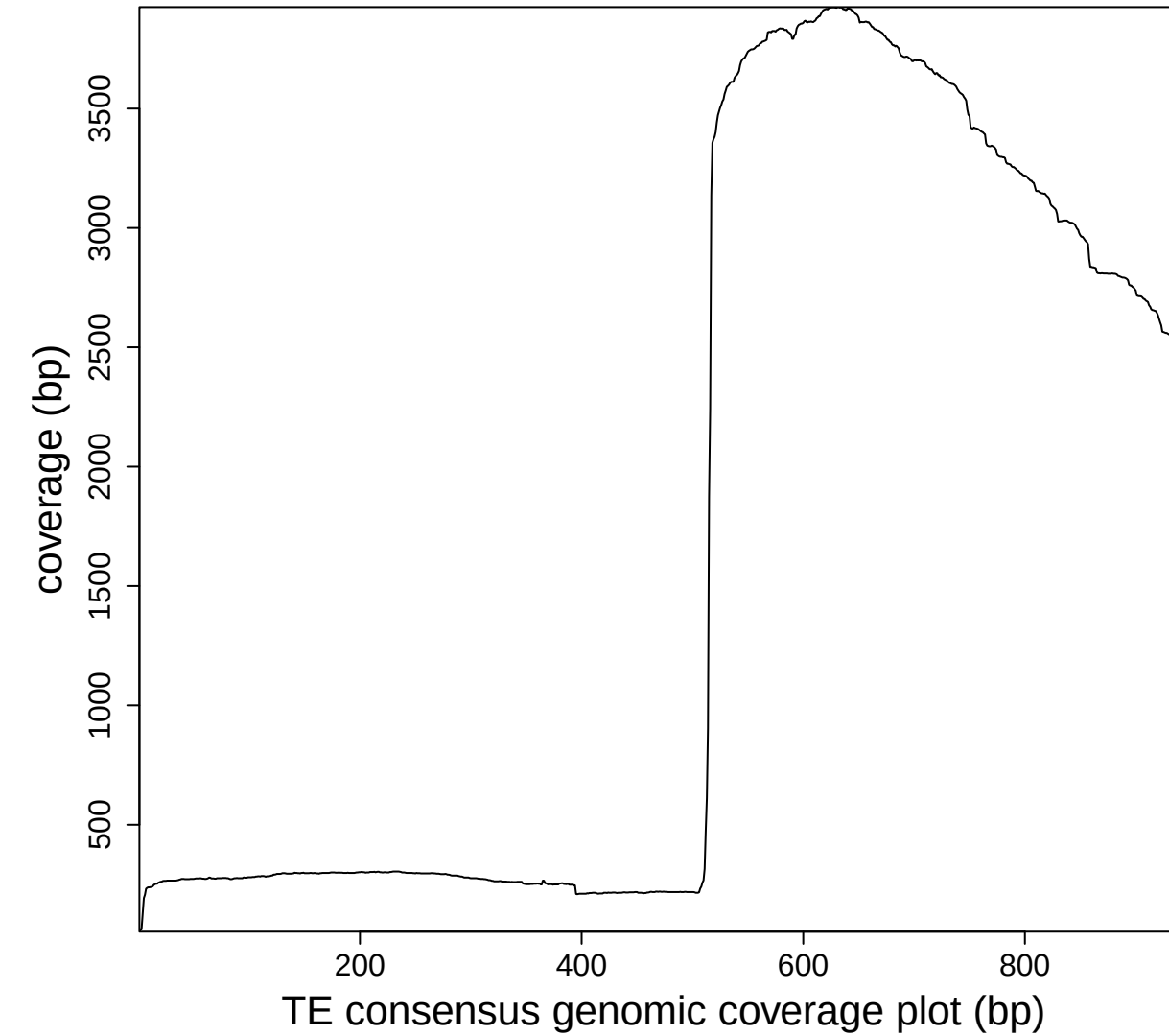
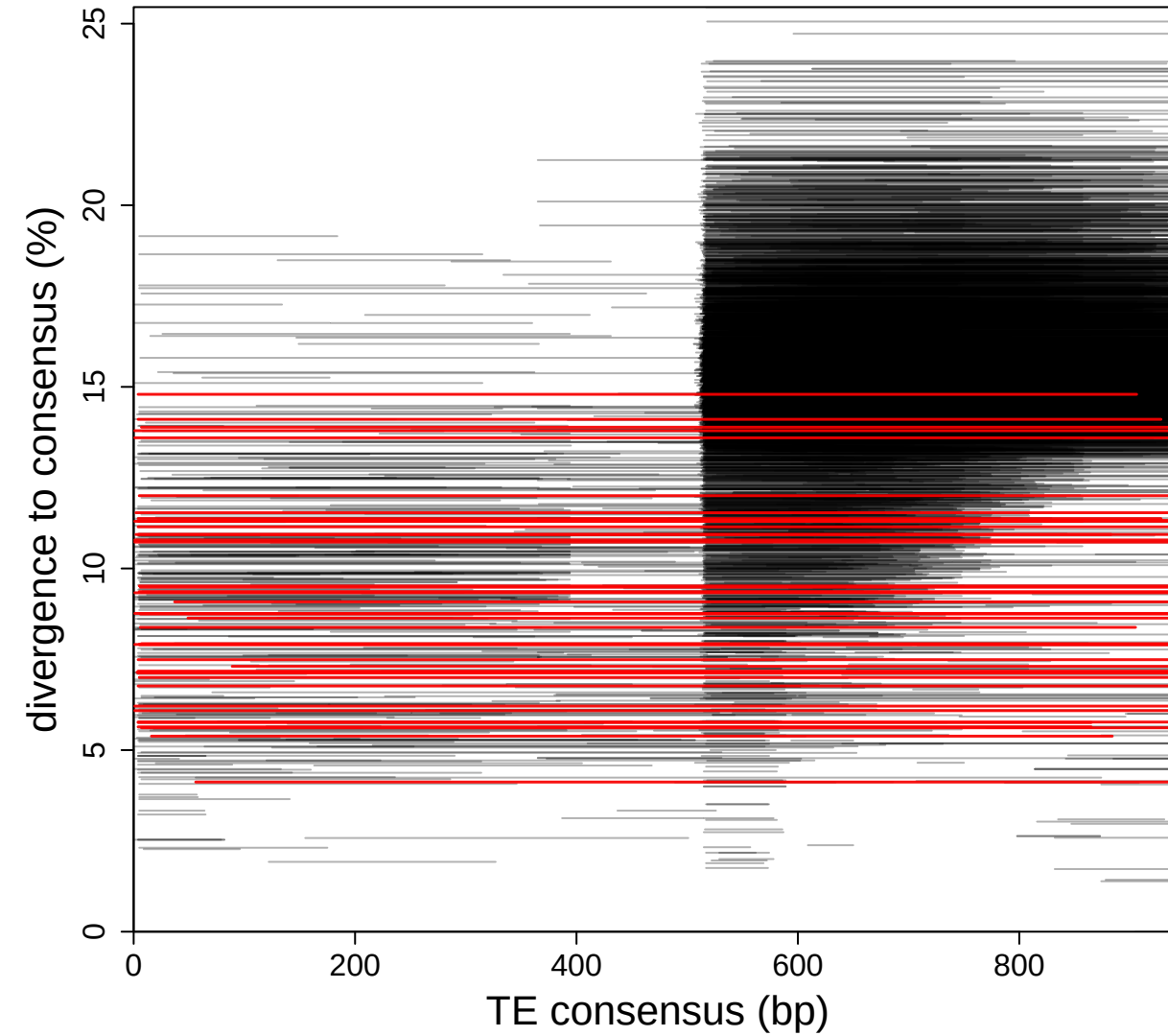
1.fasta.b.bed uf.bed g 2.bed fm 1.bed 0 0 bcln.fa aln.fa cl.f
size: 1733bp; fragments: 1597; full length: 0 (>=1559.7bp)

After TEtrimmer Extended plot Blue lines are boundaries

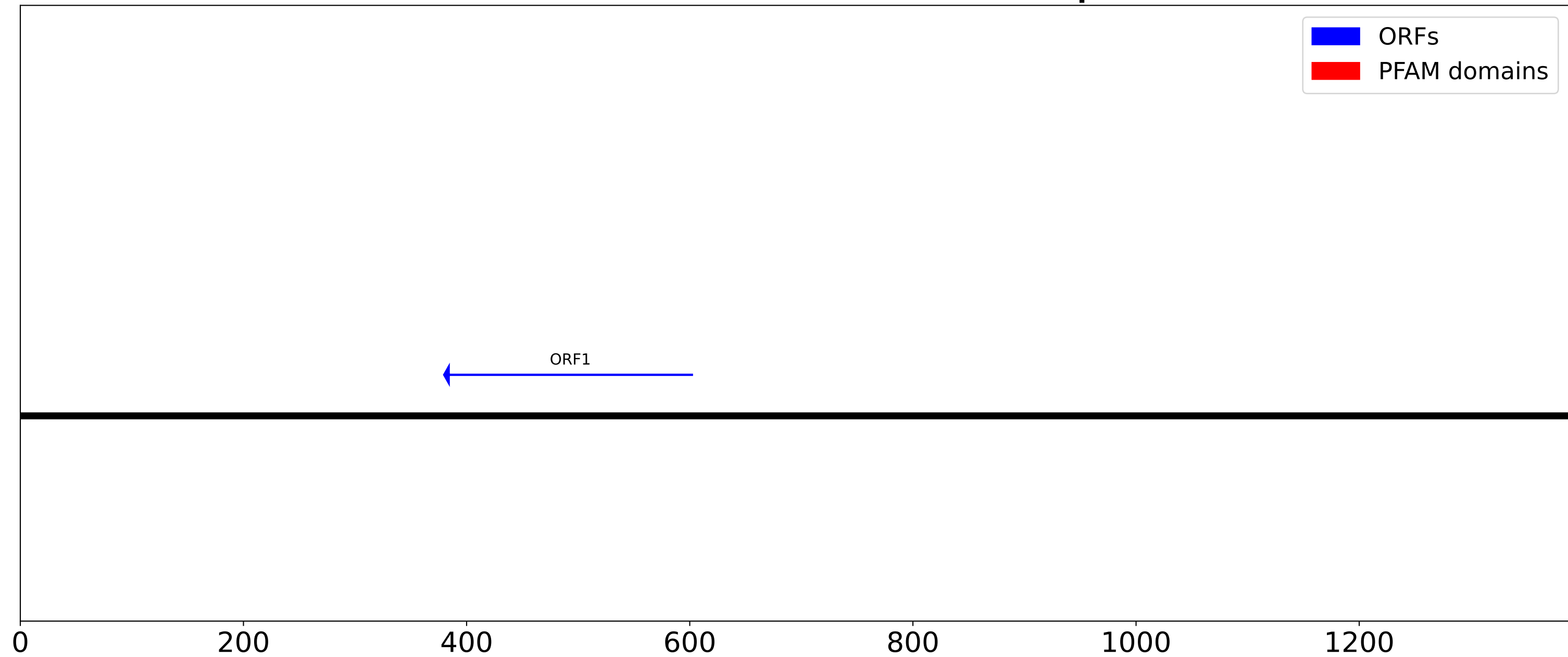


TE: rnd_5_family_4651
size: 947bp; fragments: 4551; full length: 44 (≥ 852.3 bp)

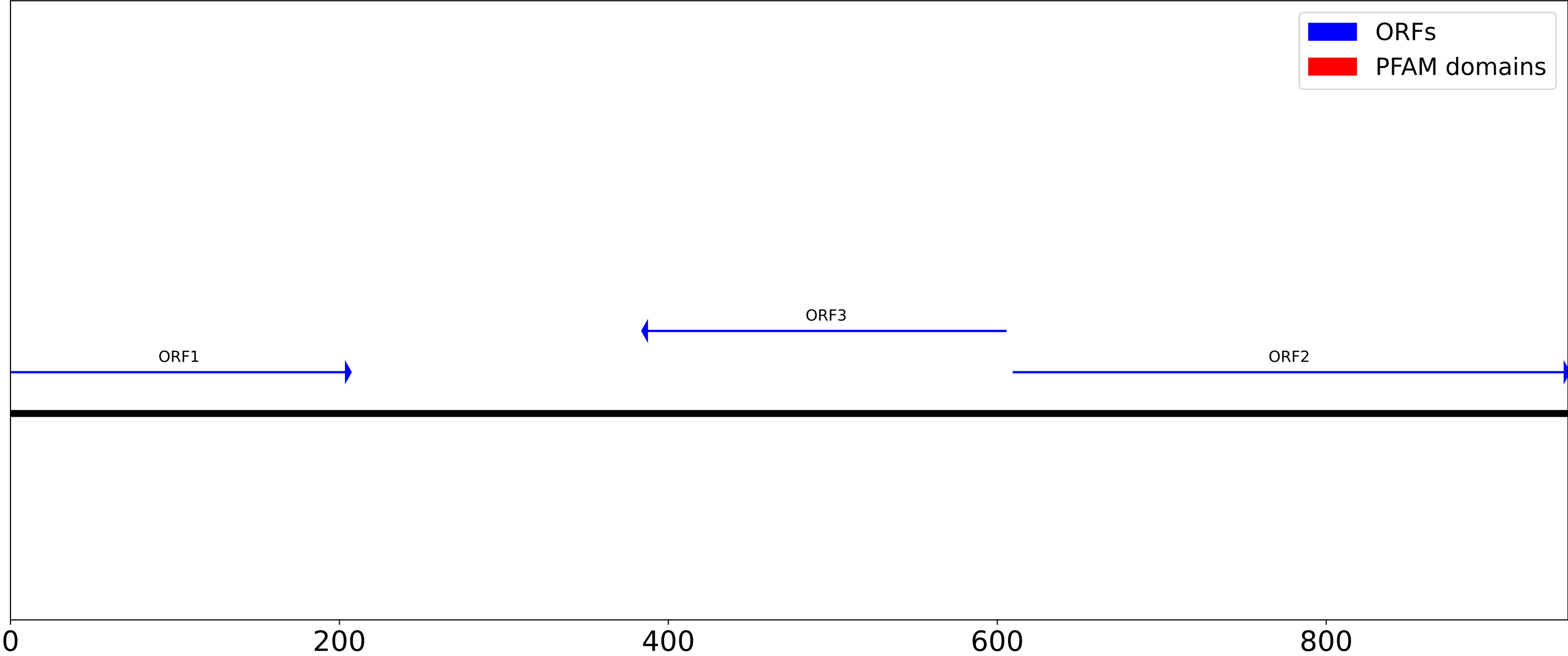
Before TEtrimmer 947 bp



After TEtrimmer ORF and PFAM domain plot



Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

