

Start crop Point



End crop Point



MSA length = 1186



Start crop Point



End crop Point



0

10

0

200

400

600

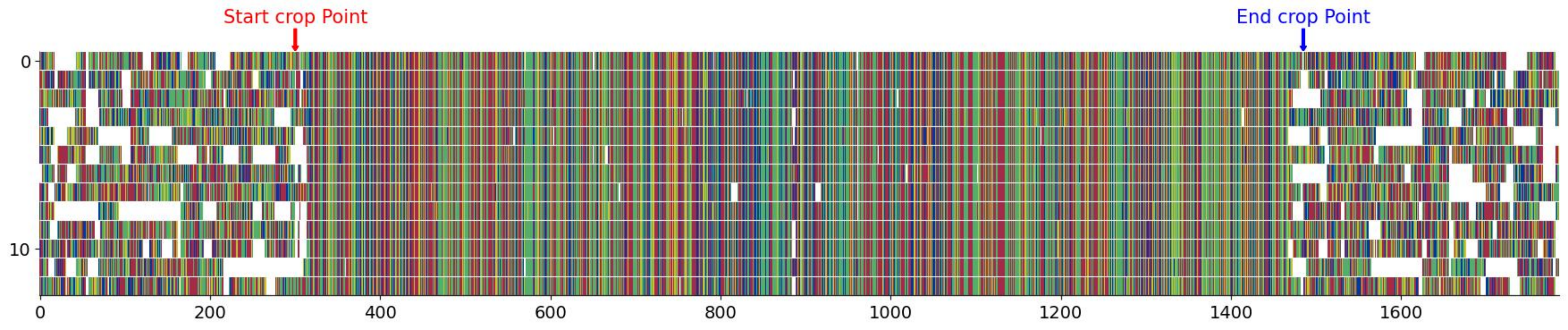
800

1000

1200

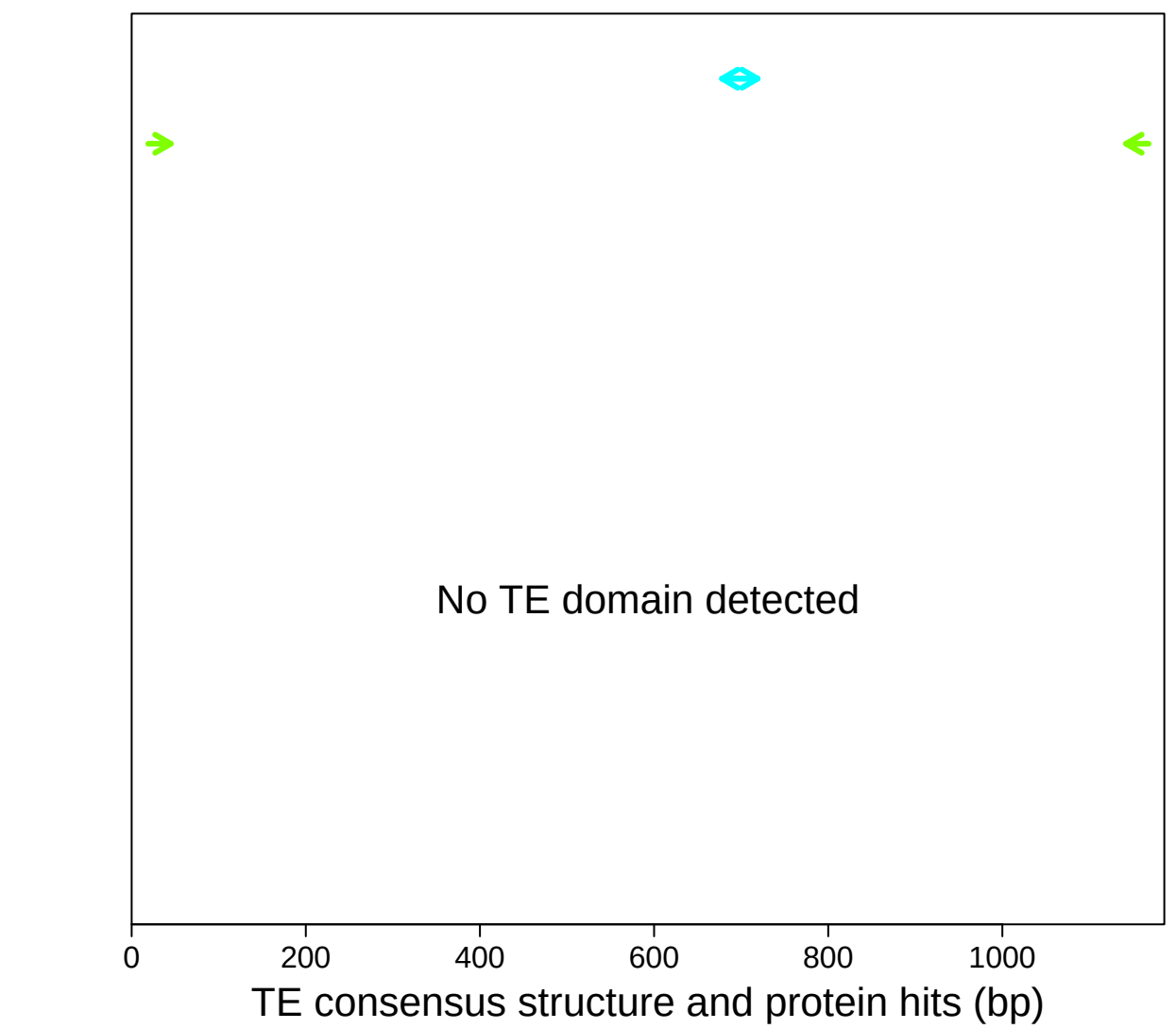
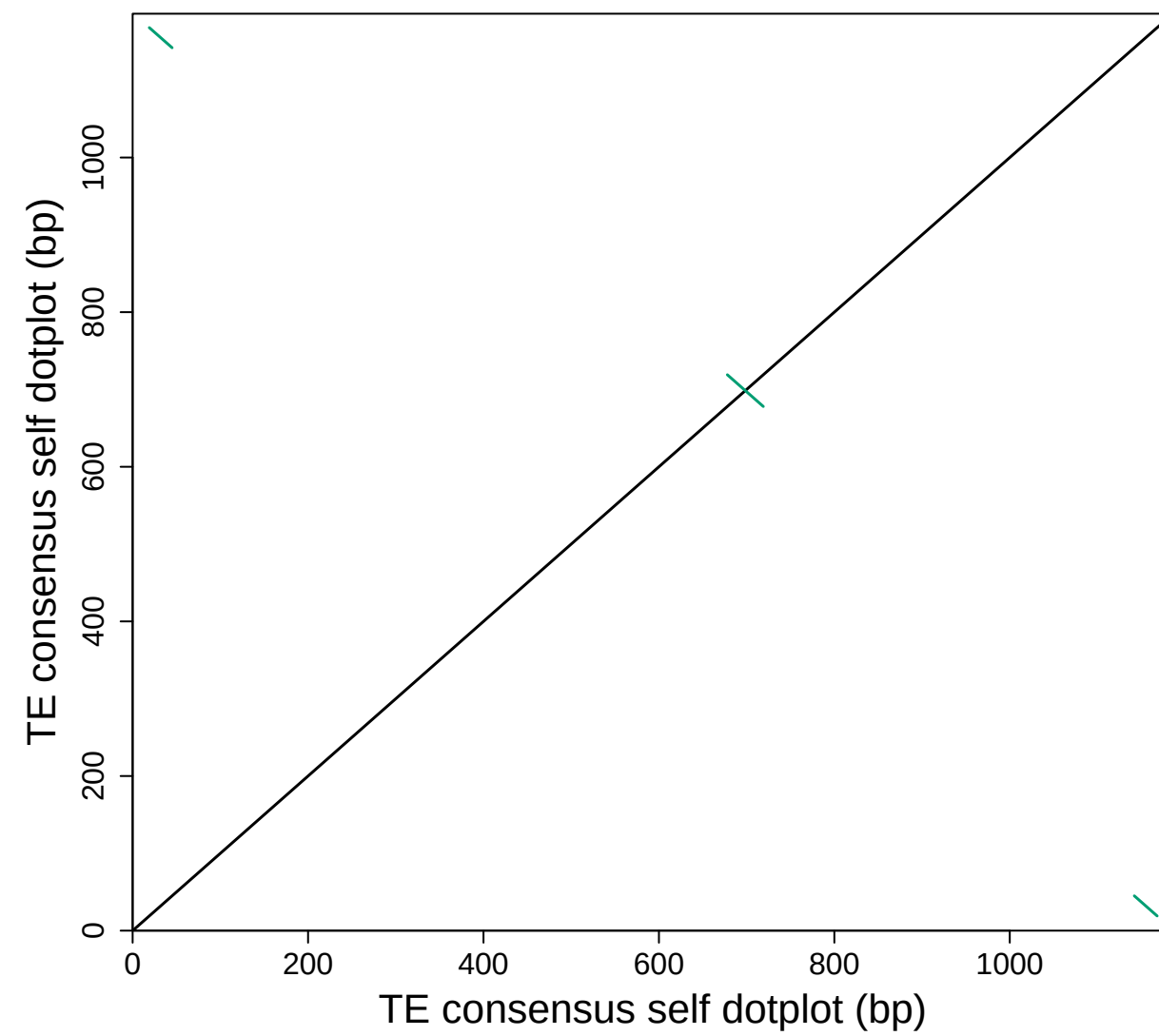
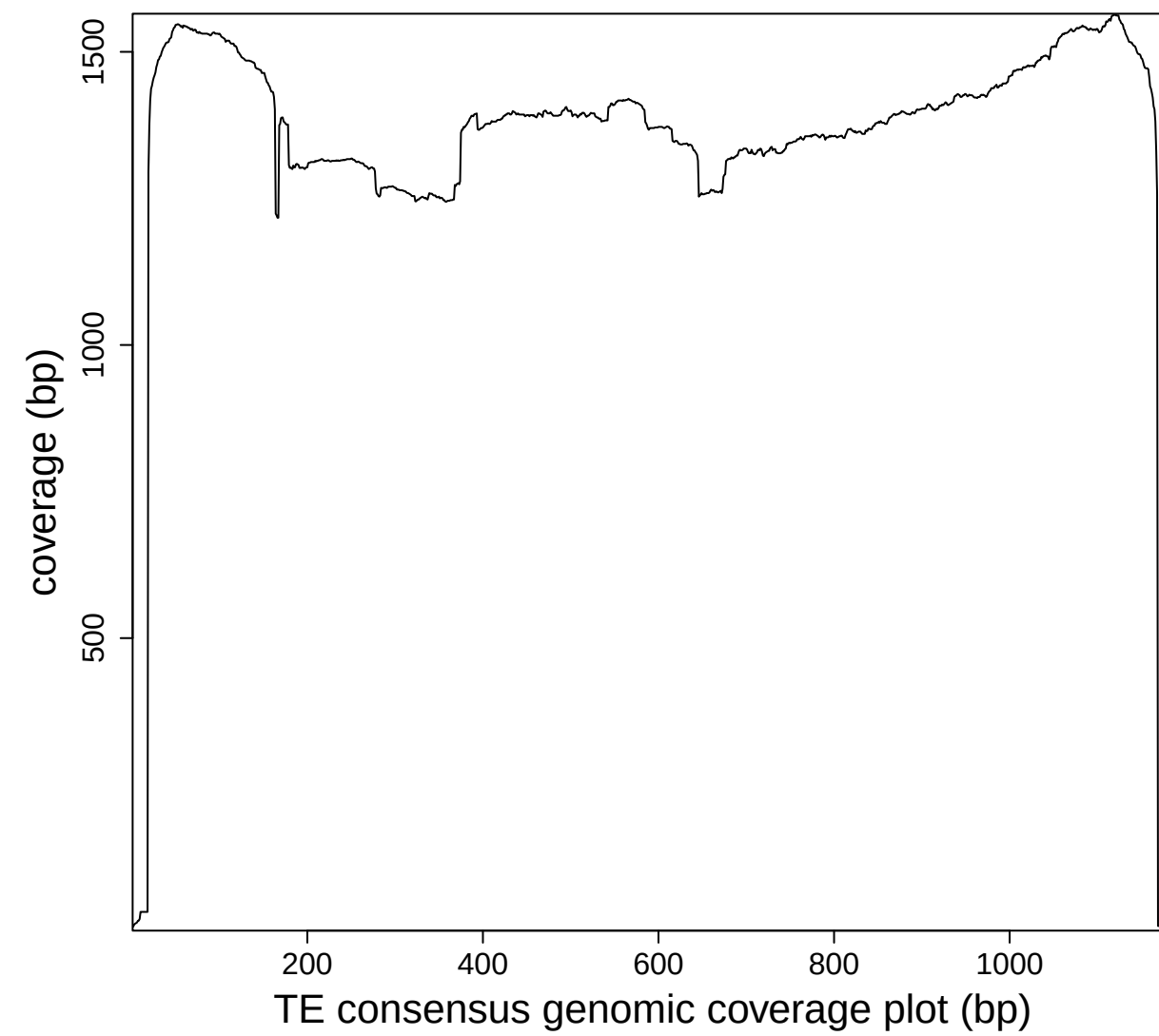
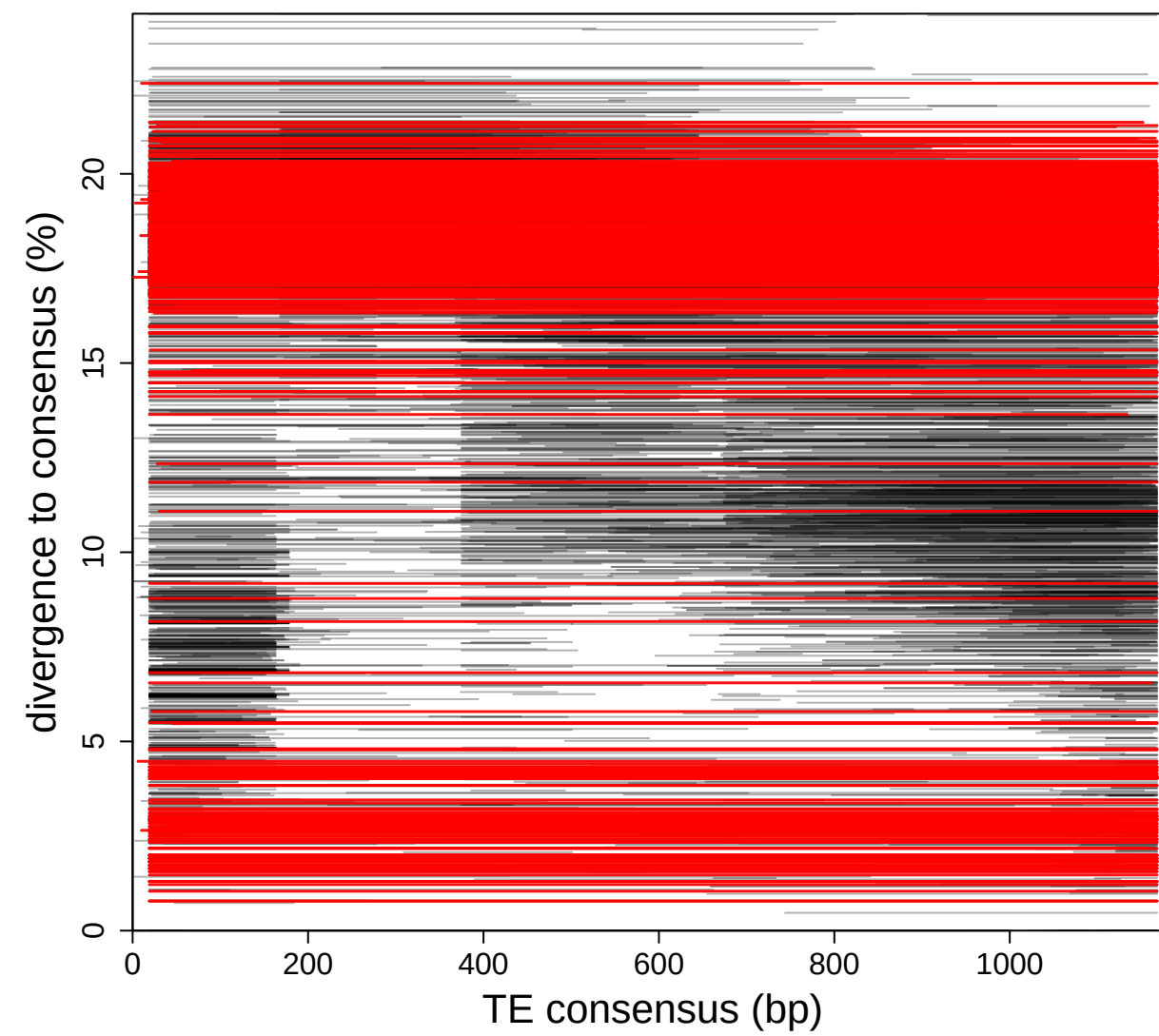
1400

1600



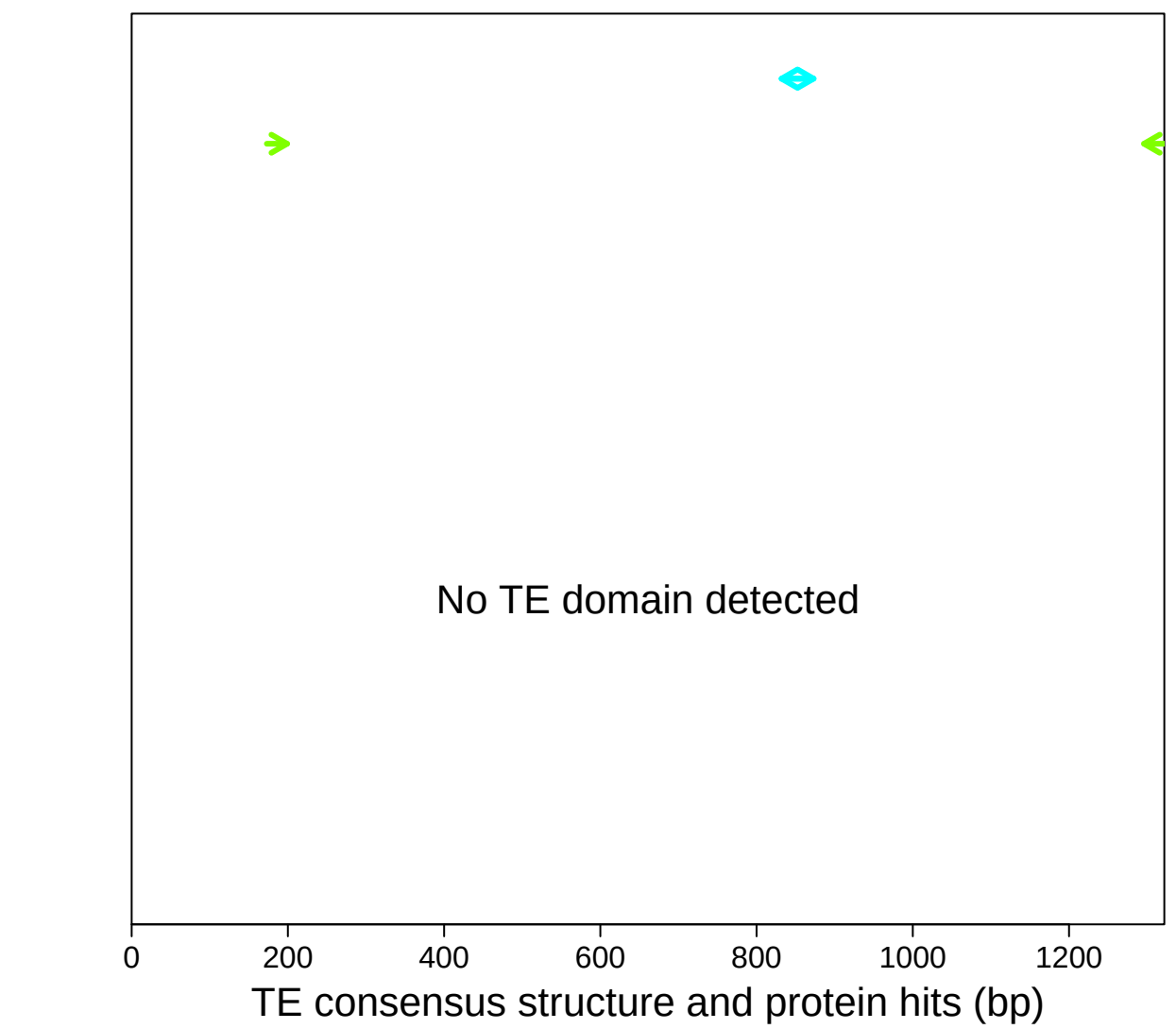
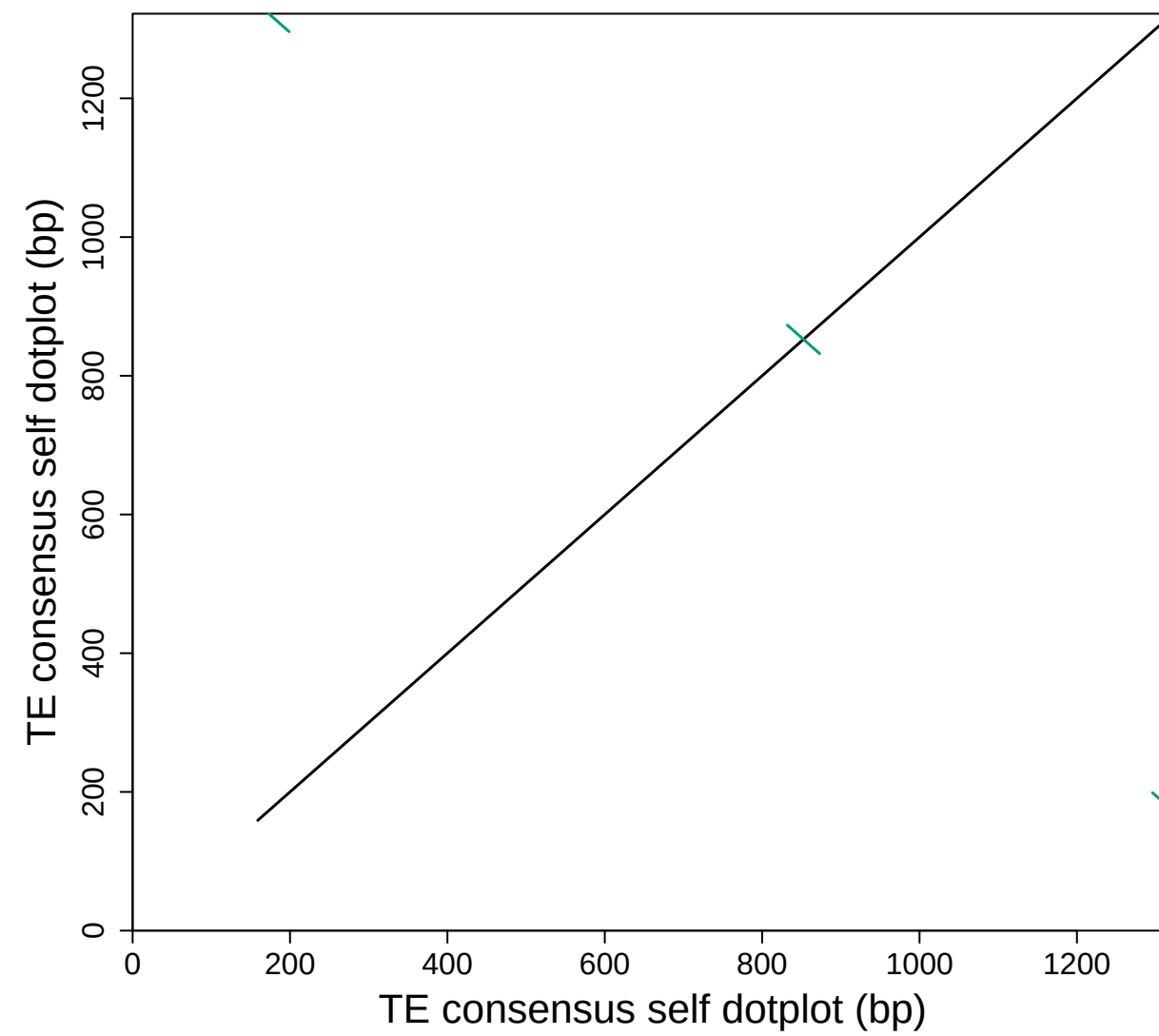
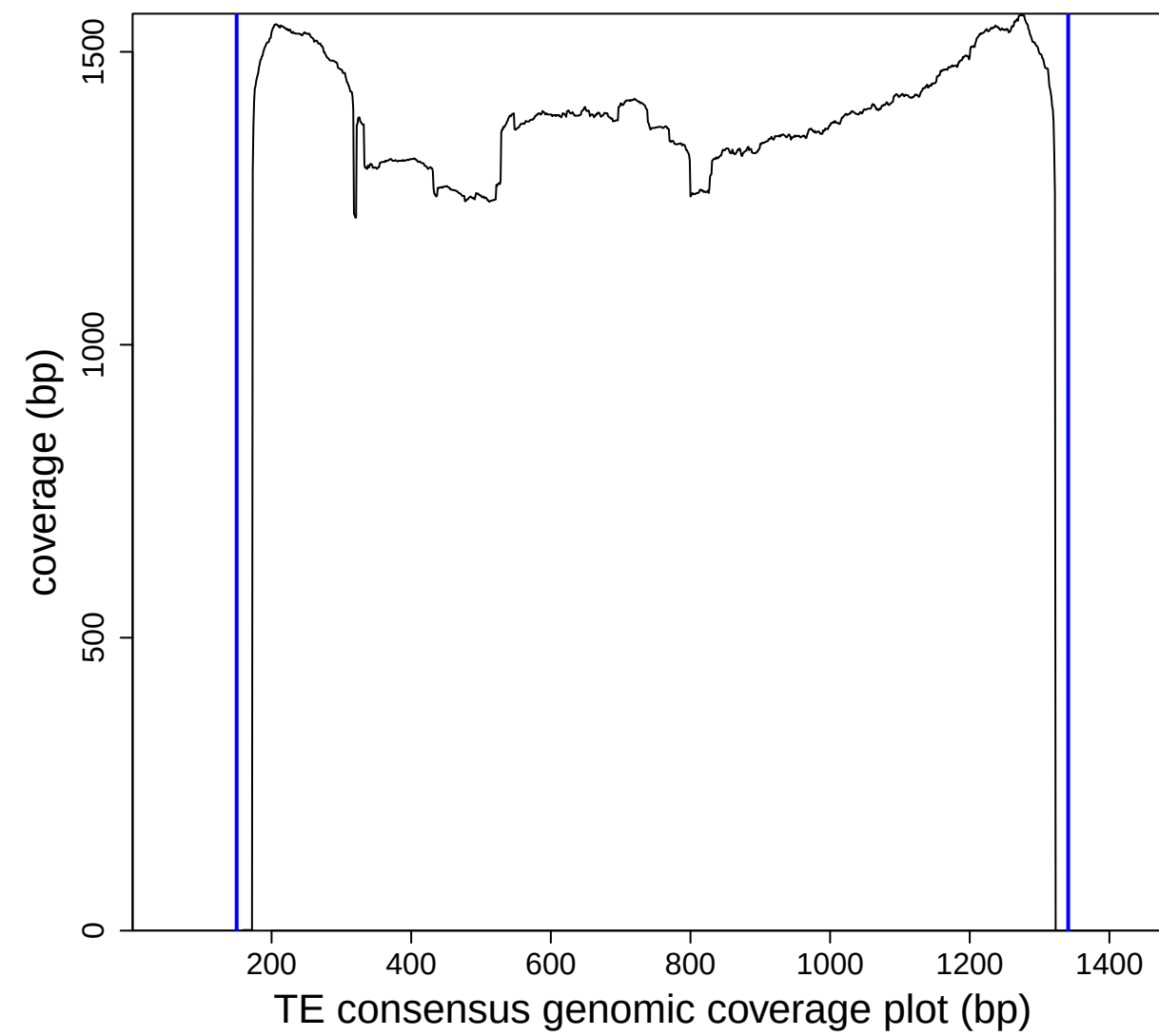
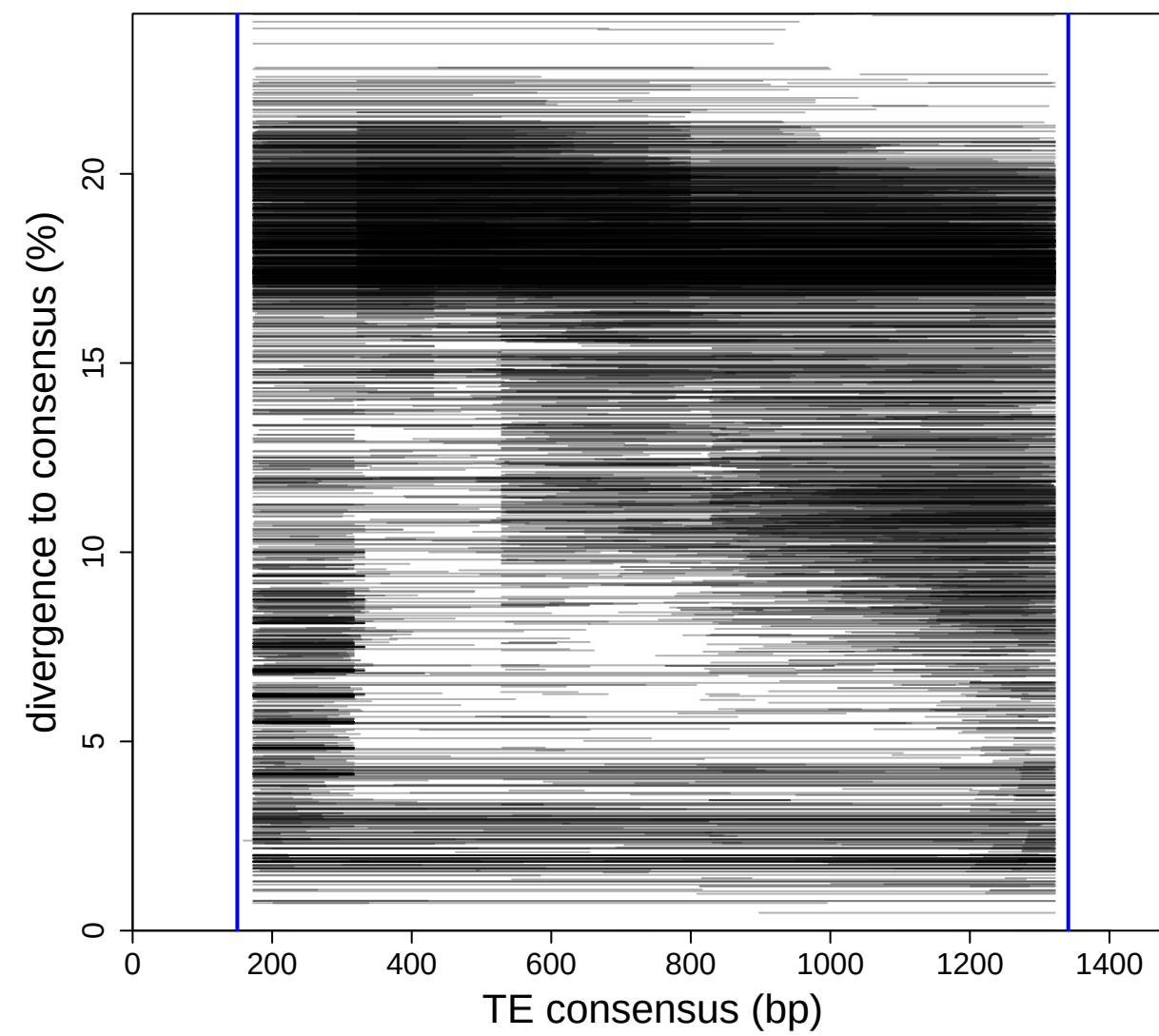
b.bed uf.bed g 2.bed fm 1.bed 0 0 bcln.fa aln.fa cl.fa gs.fa
size: 1186bp; fragments: 3041; full length: 562 (≥ 1067.4 bp)

After TEtrimmer 1186 bp



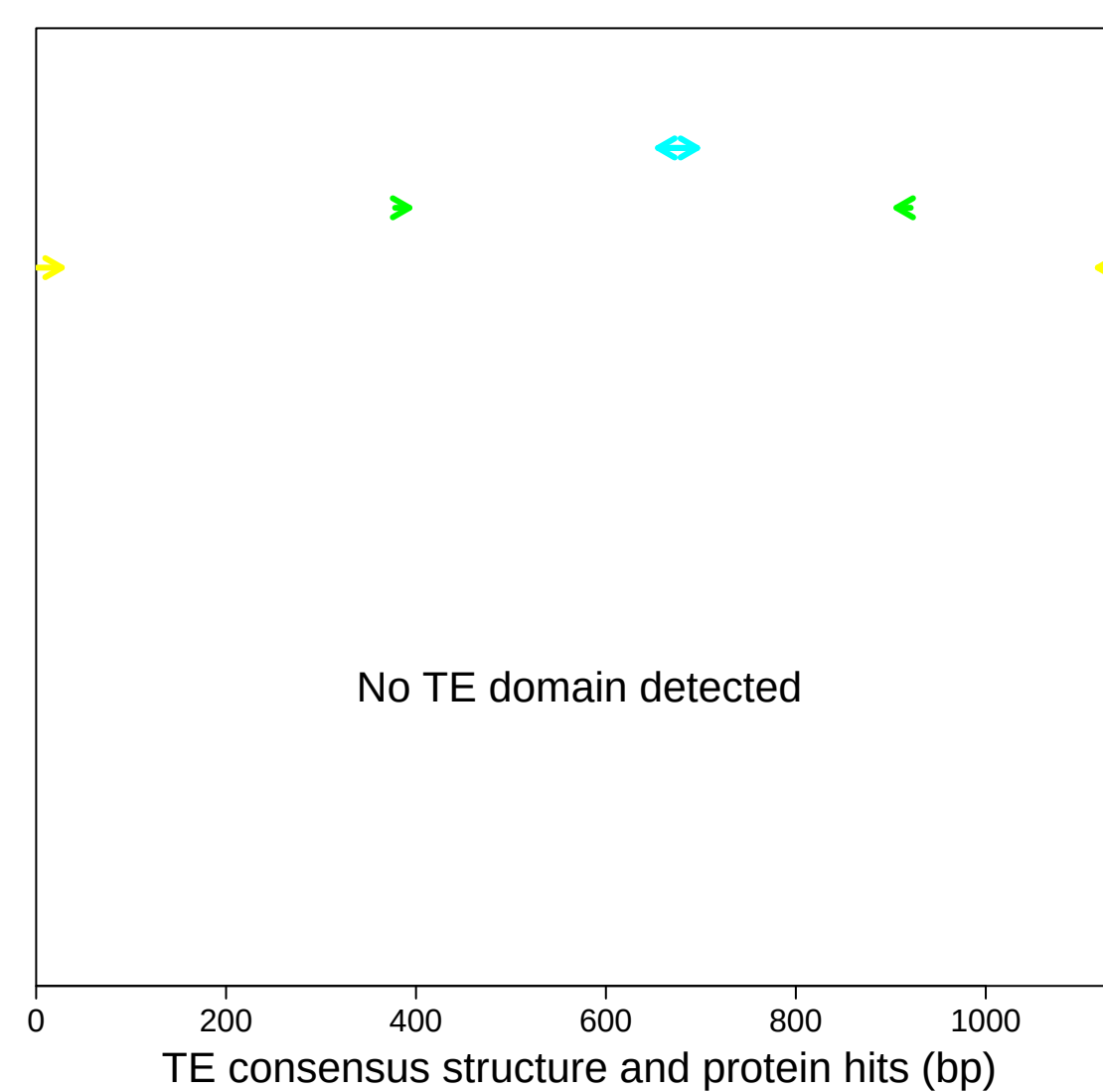
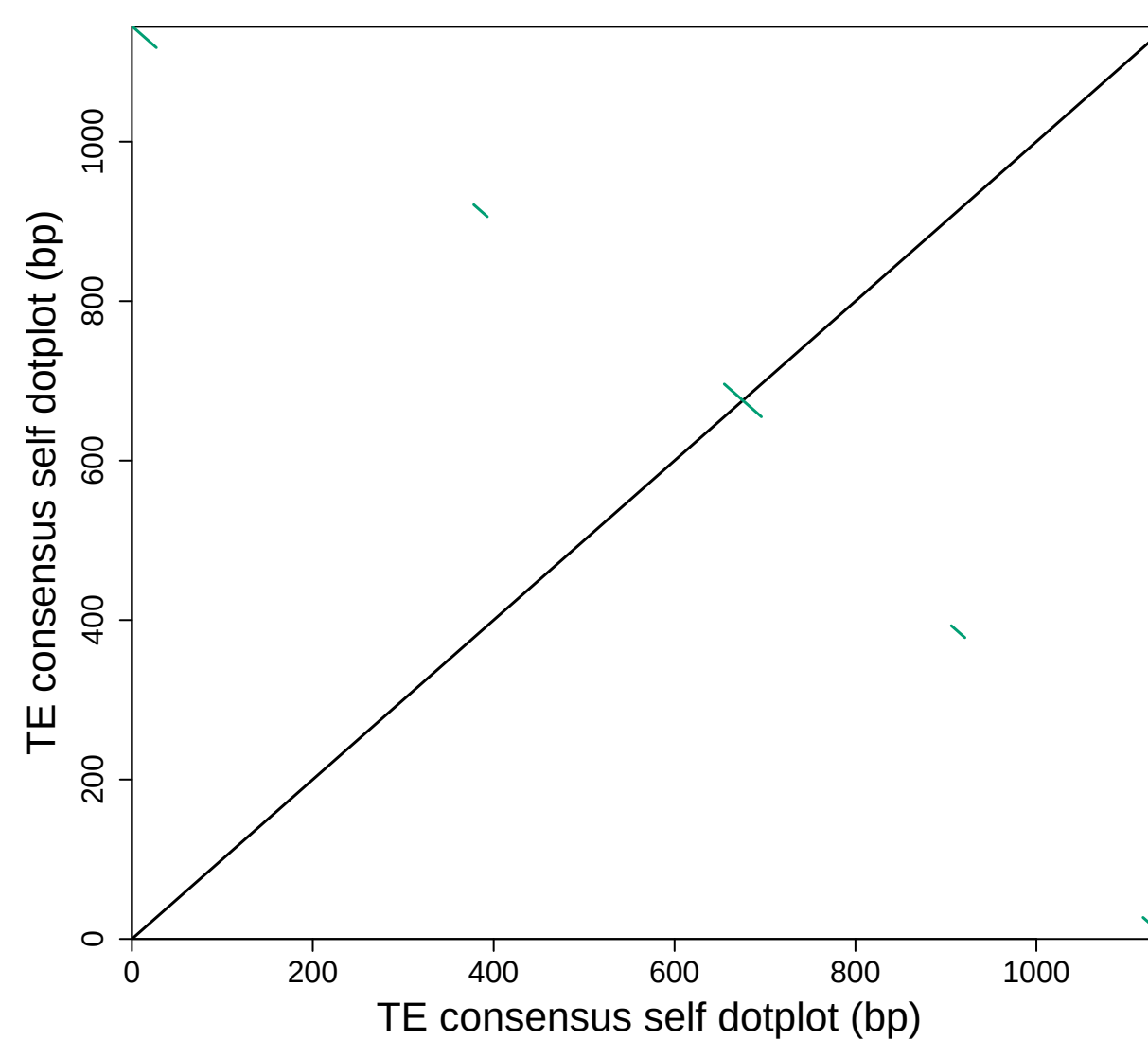
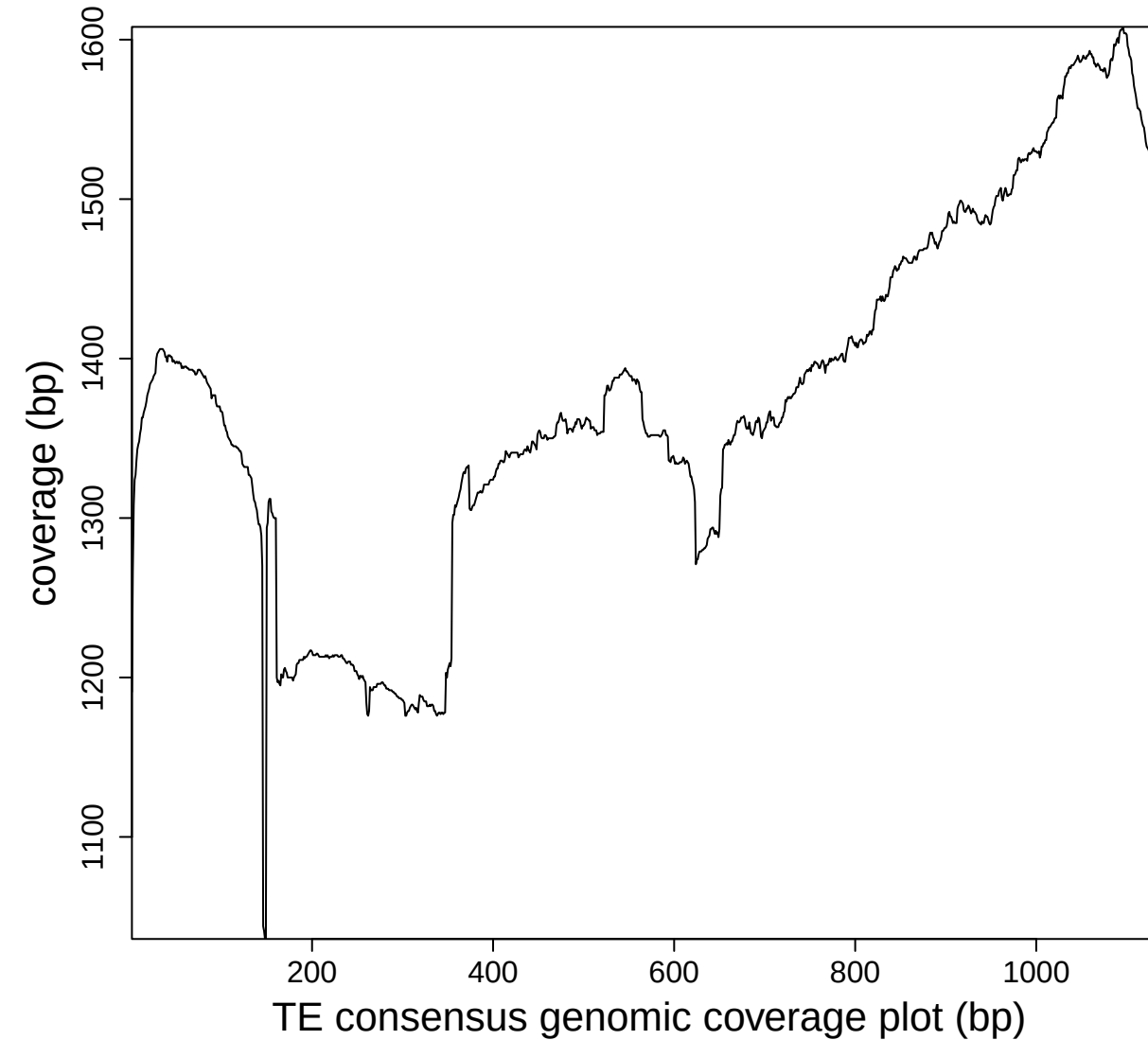
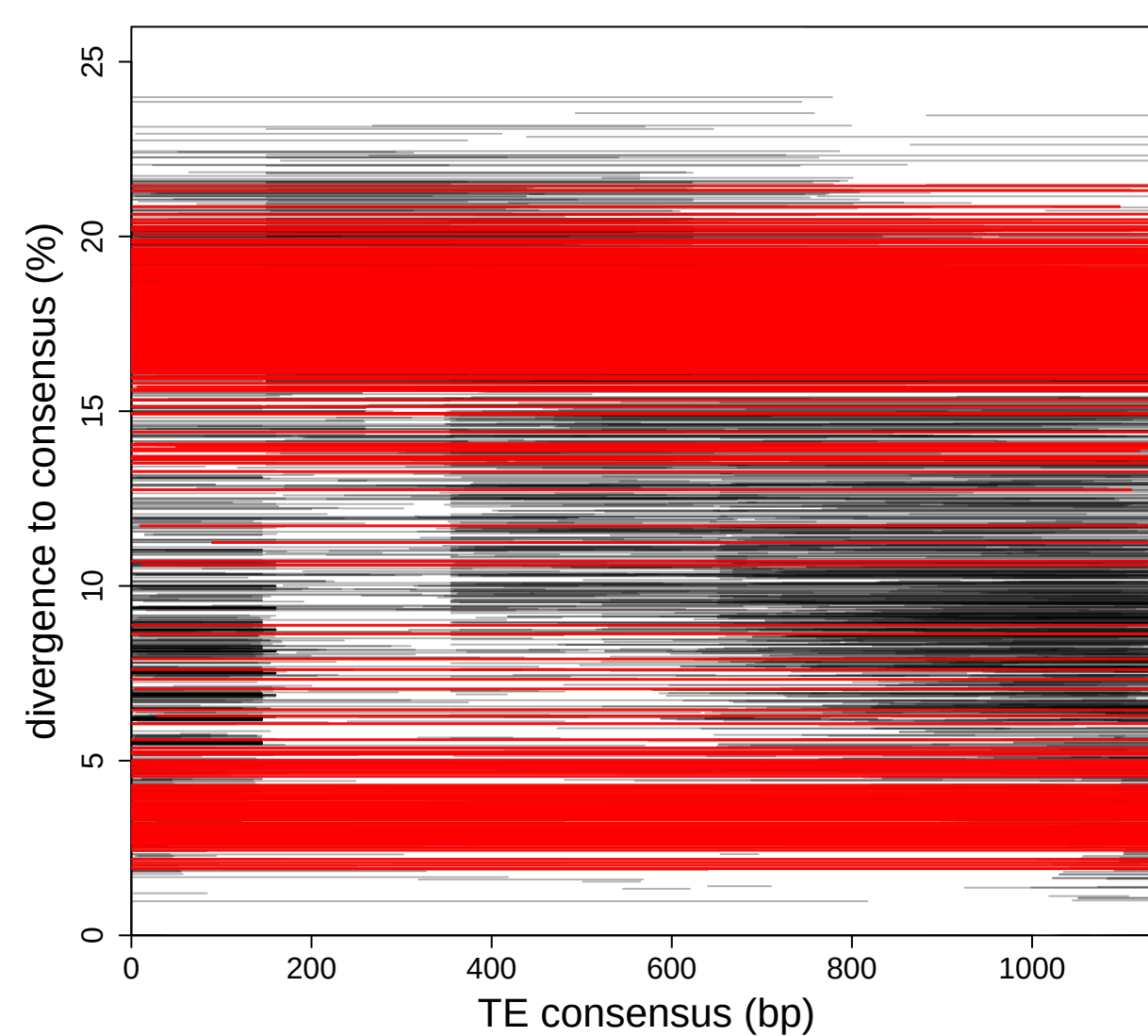
2.fasta.b.bed_uf.bed_g_2.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa
size: 1491bp; fragments: 3041; full length: 0 (>=1341.9bp)

After TEtrimmer Extended plot Blue lines are boundaries

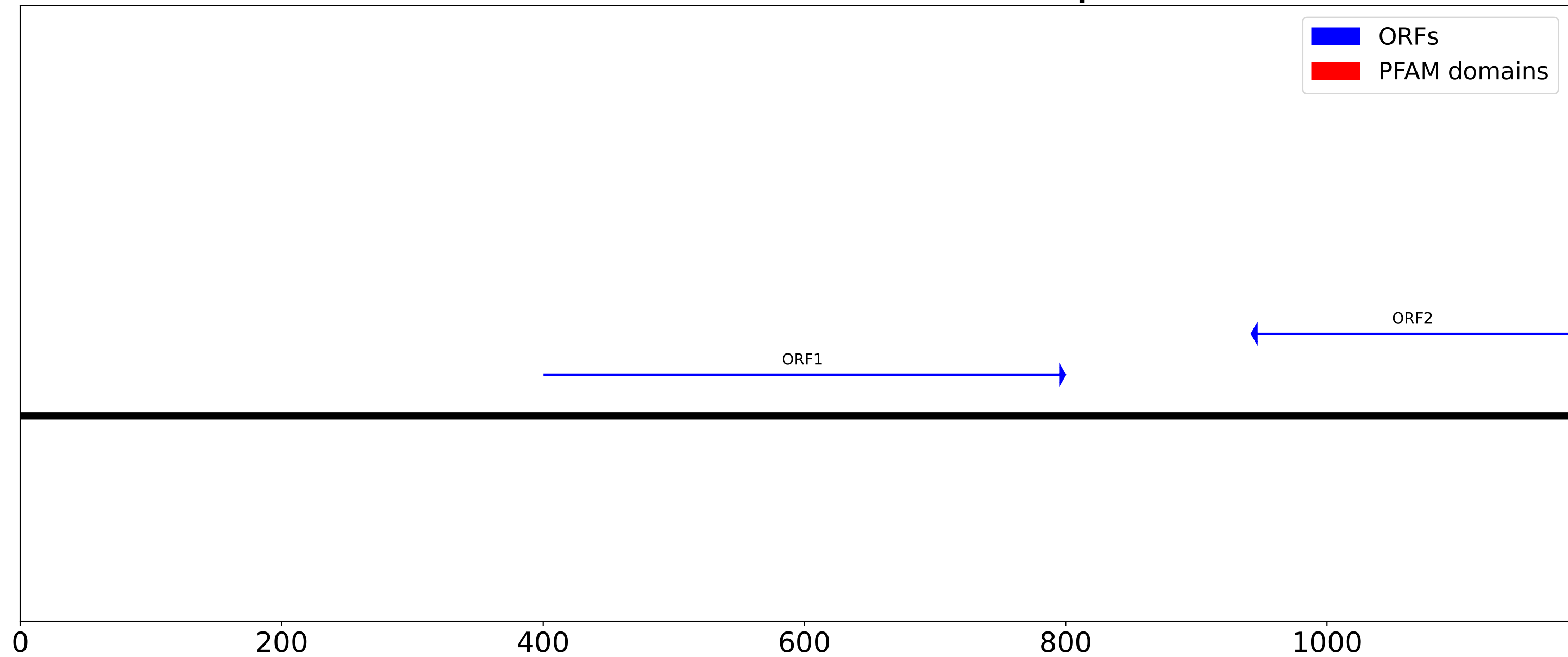


TE: rnd_1_family_92
size: 1144bp; fragments: 3085; full length: 480 (≥ 1029.6 bp)

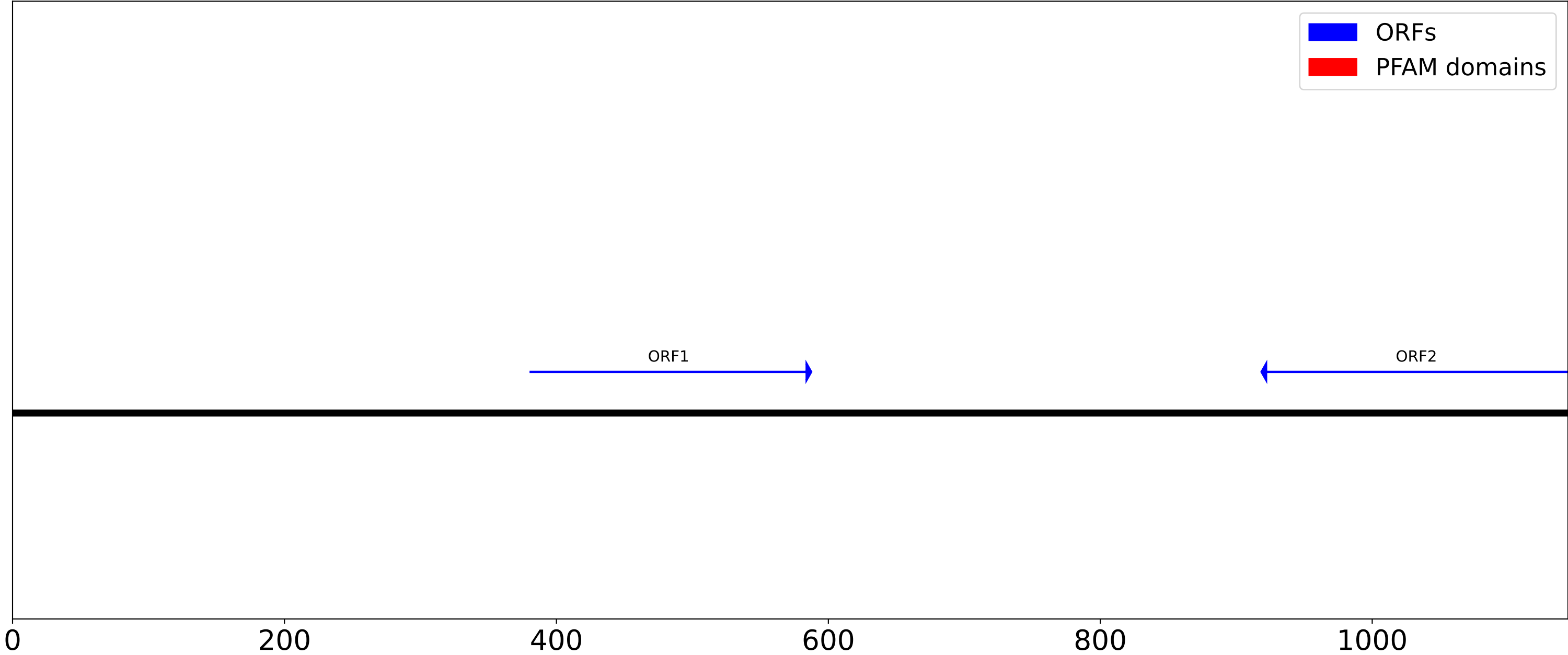
Before TEtrimmer 1144 bp



After TEtrimmer ORF and PFAM domain plot



Before TEtrimmer ORF and PFAM domain plot



Dotplot

(windowsize = 25, threshold = 50.00 09/10/25)

