

Start crop Point



End crop Point



MSA length = 1223



Start crop Point



End crop Point



0

10

0

250

500

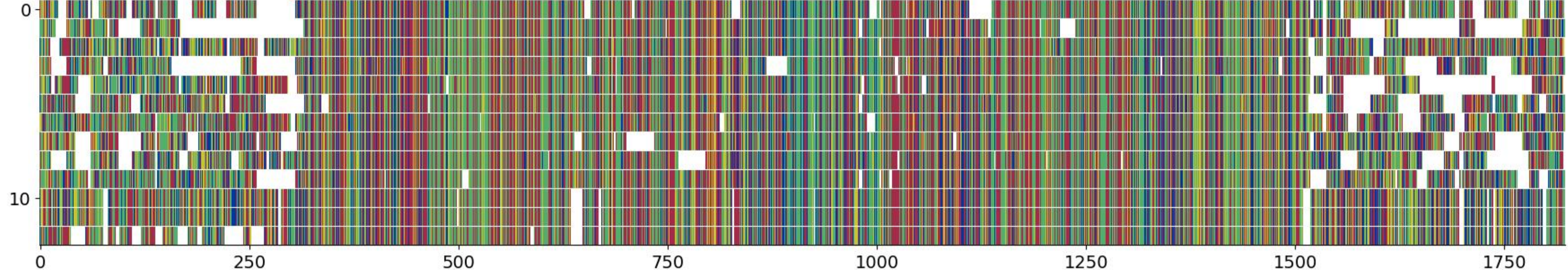
750

1000

1250

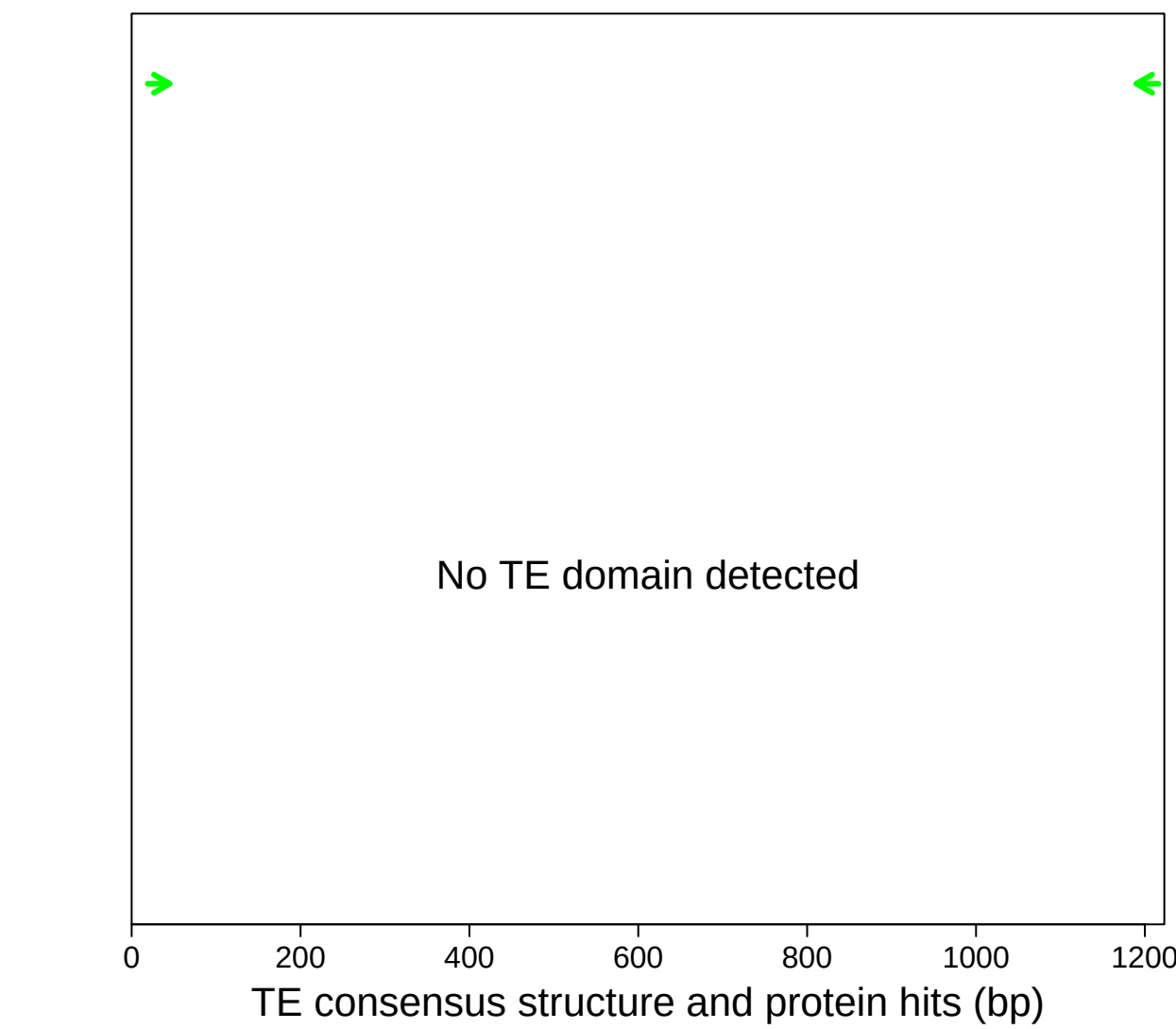
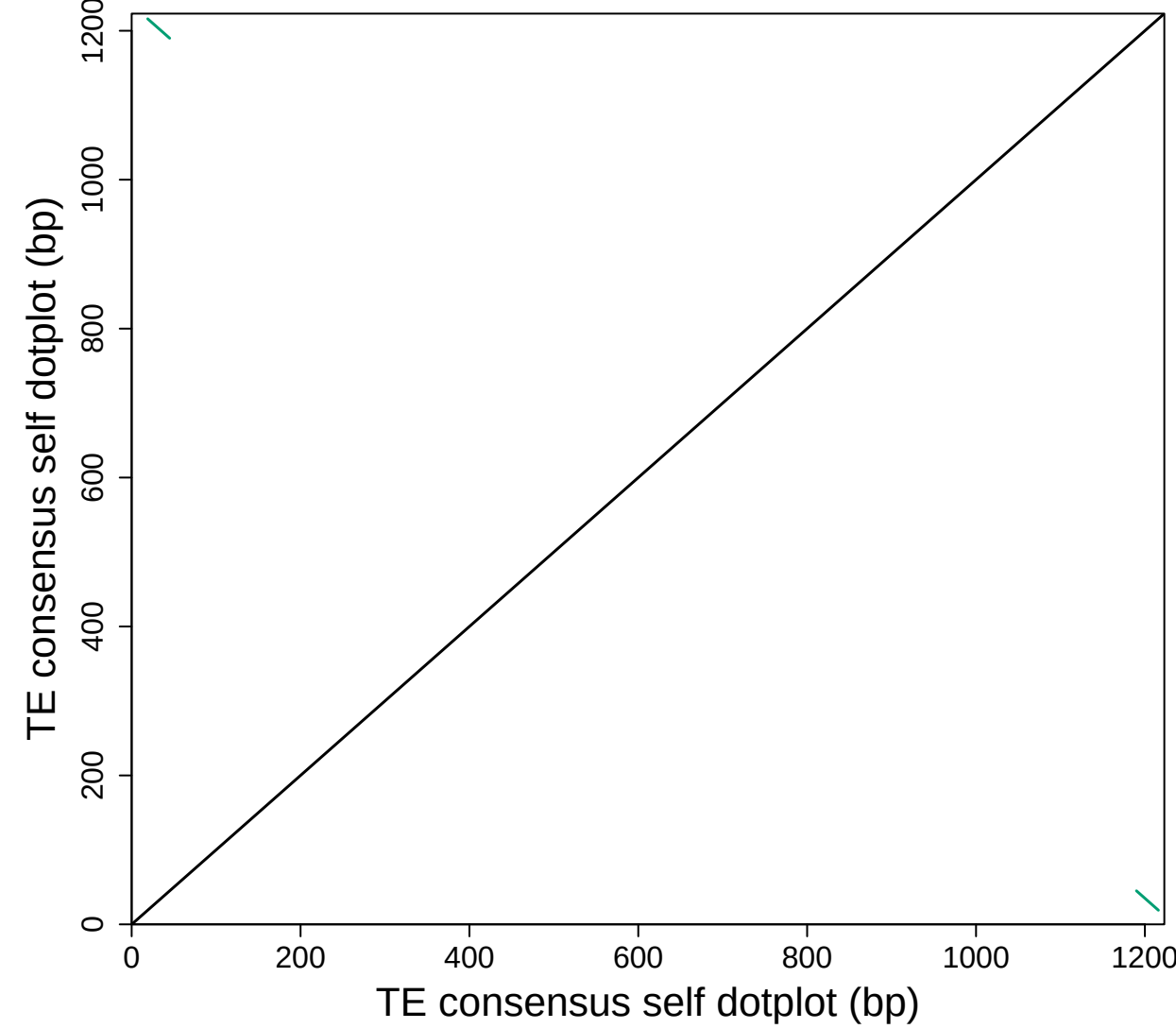
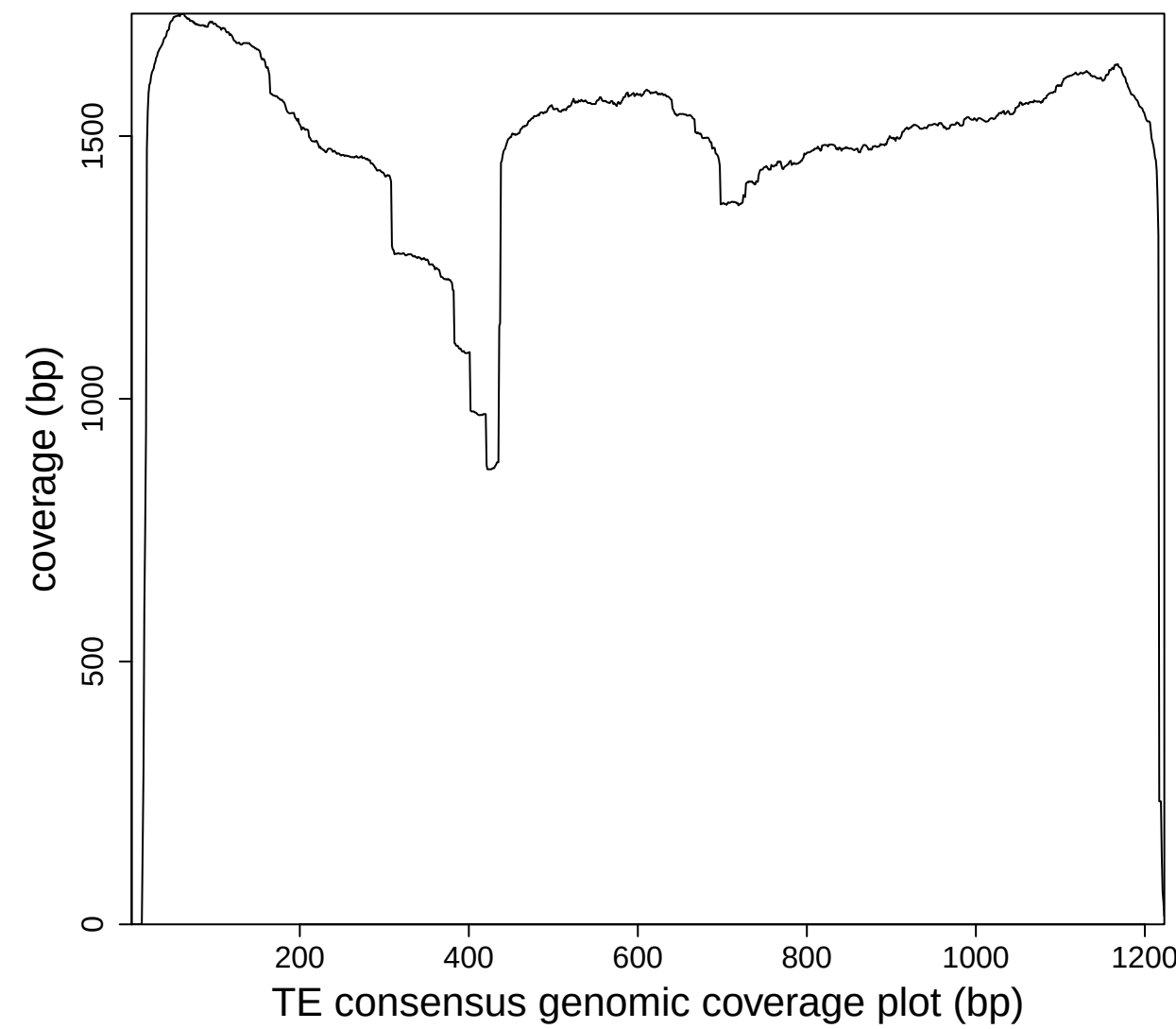
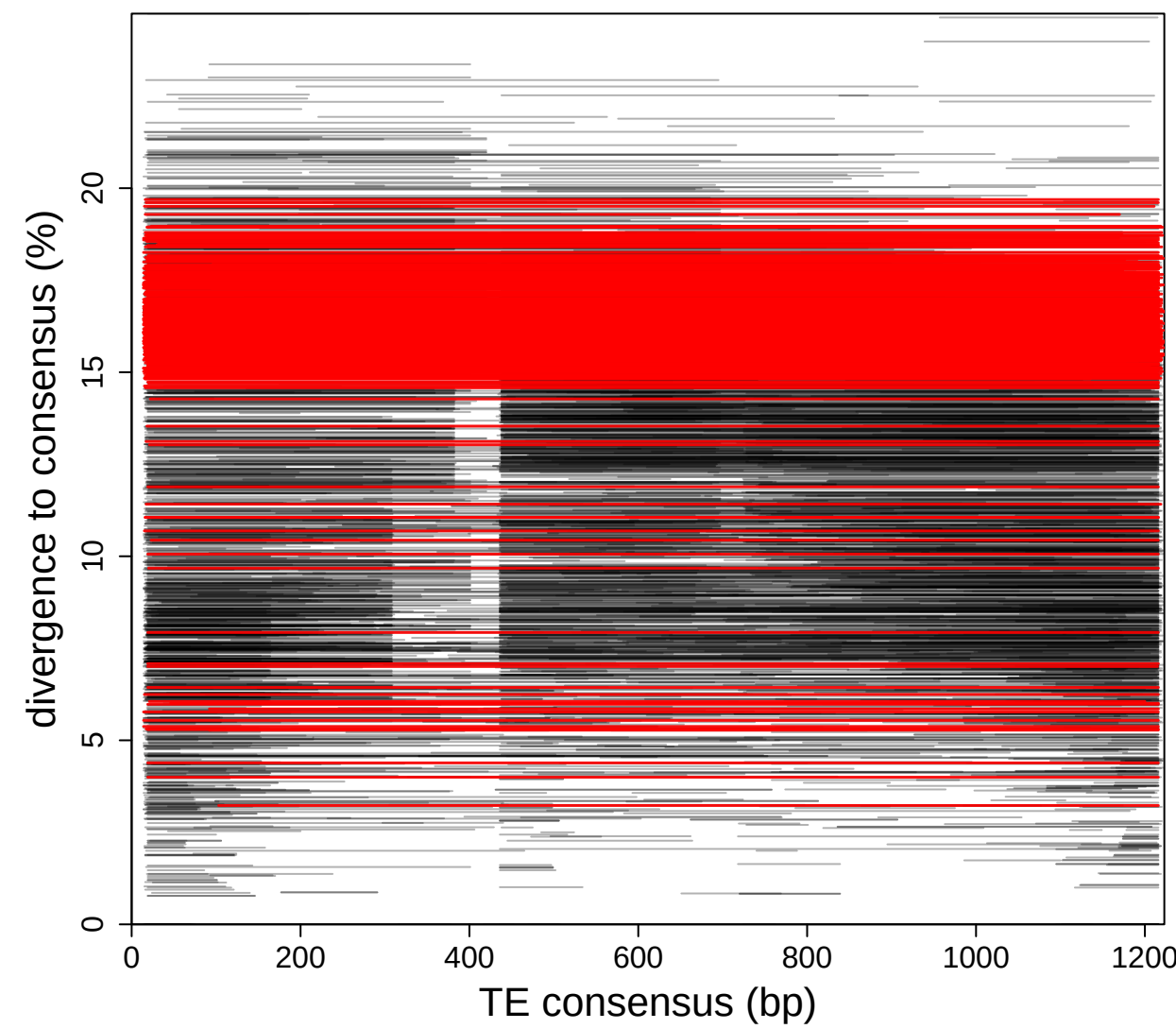
1500

1750



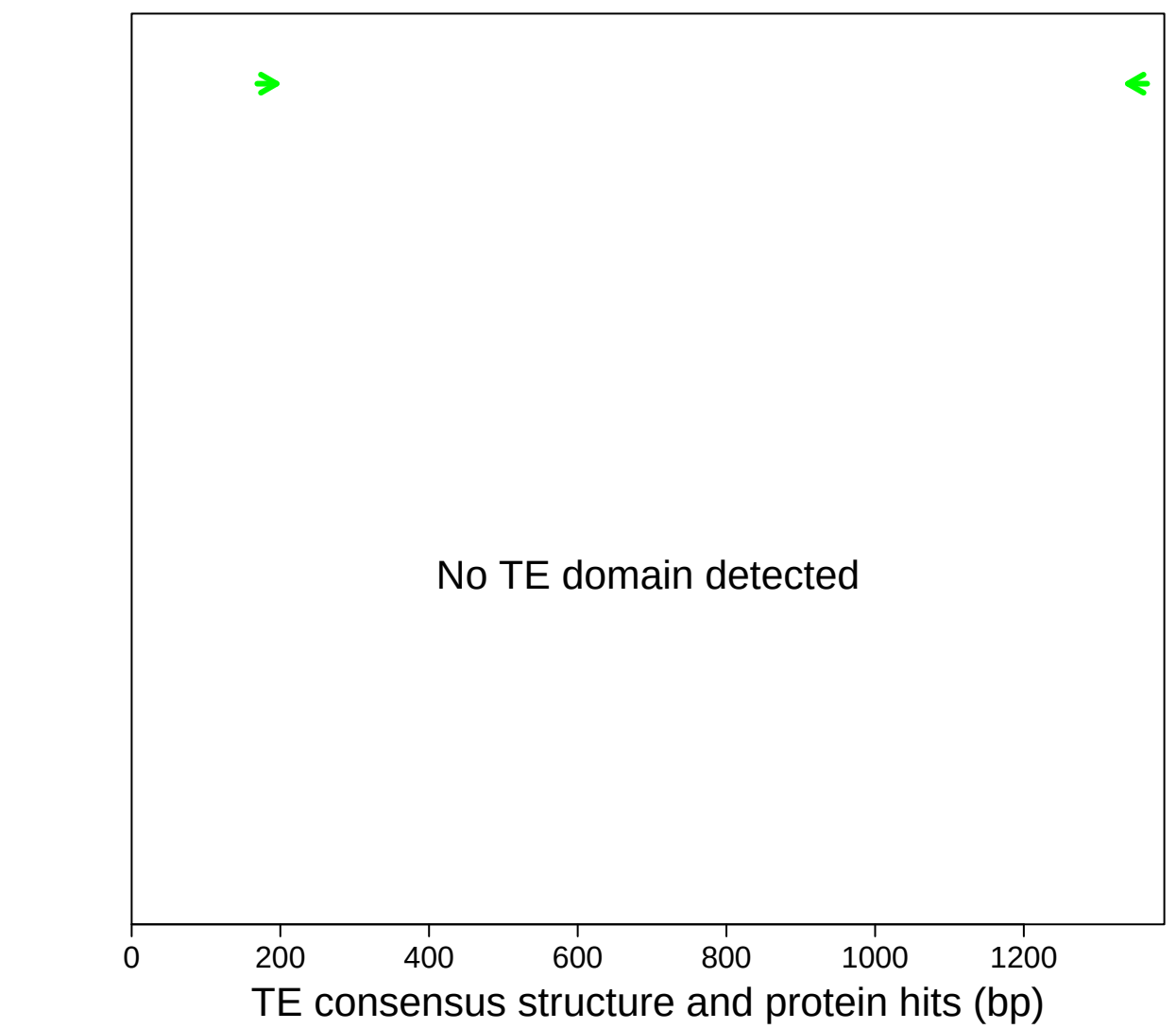
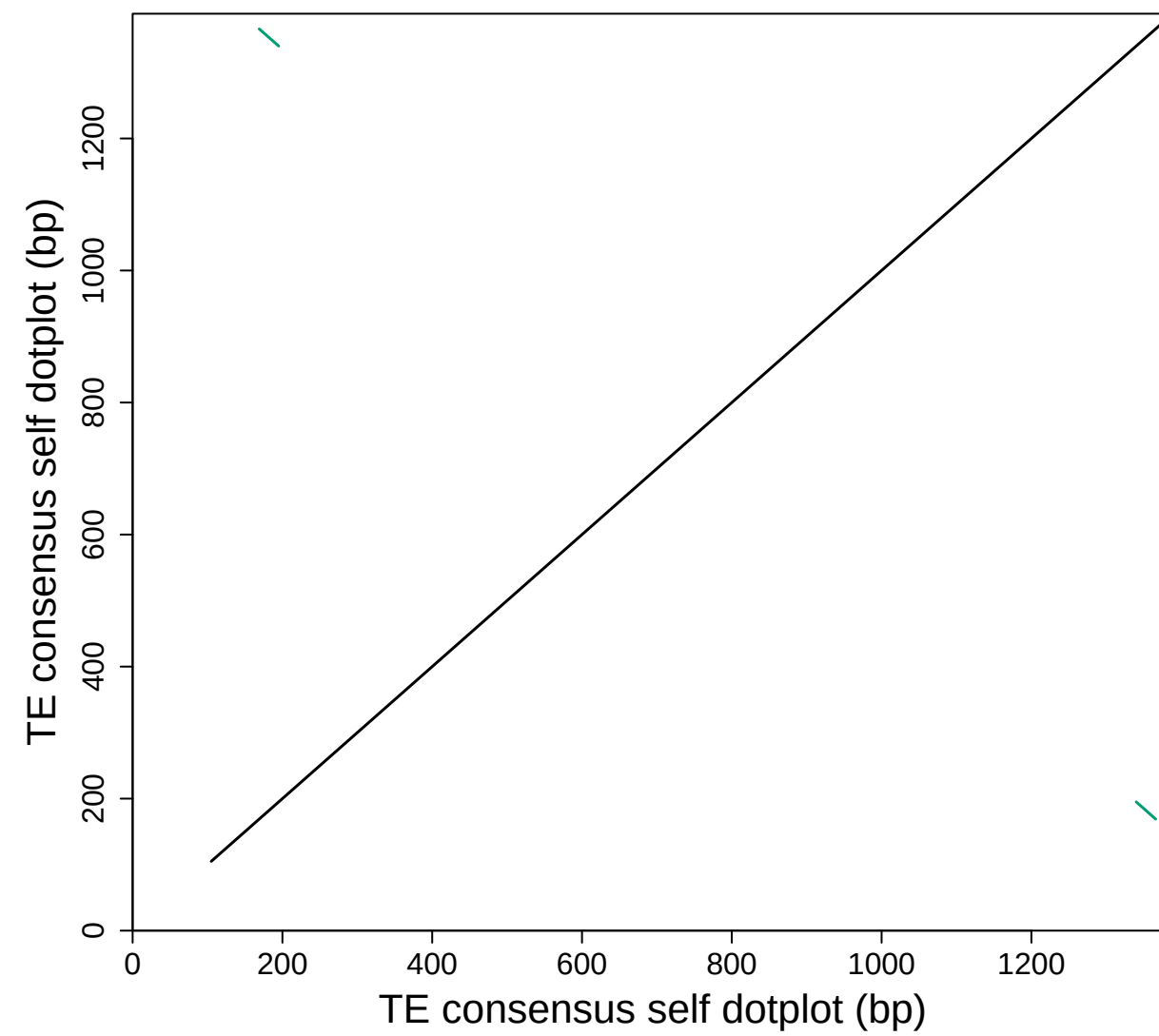
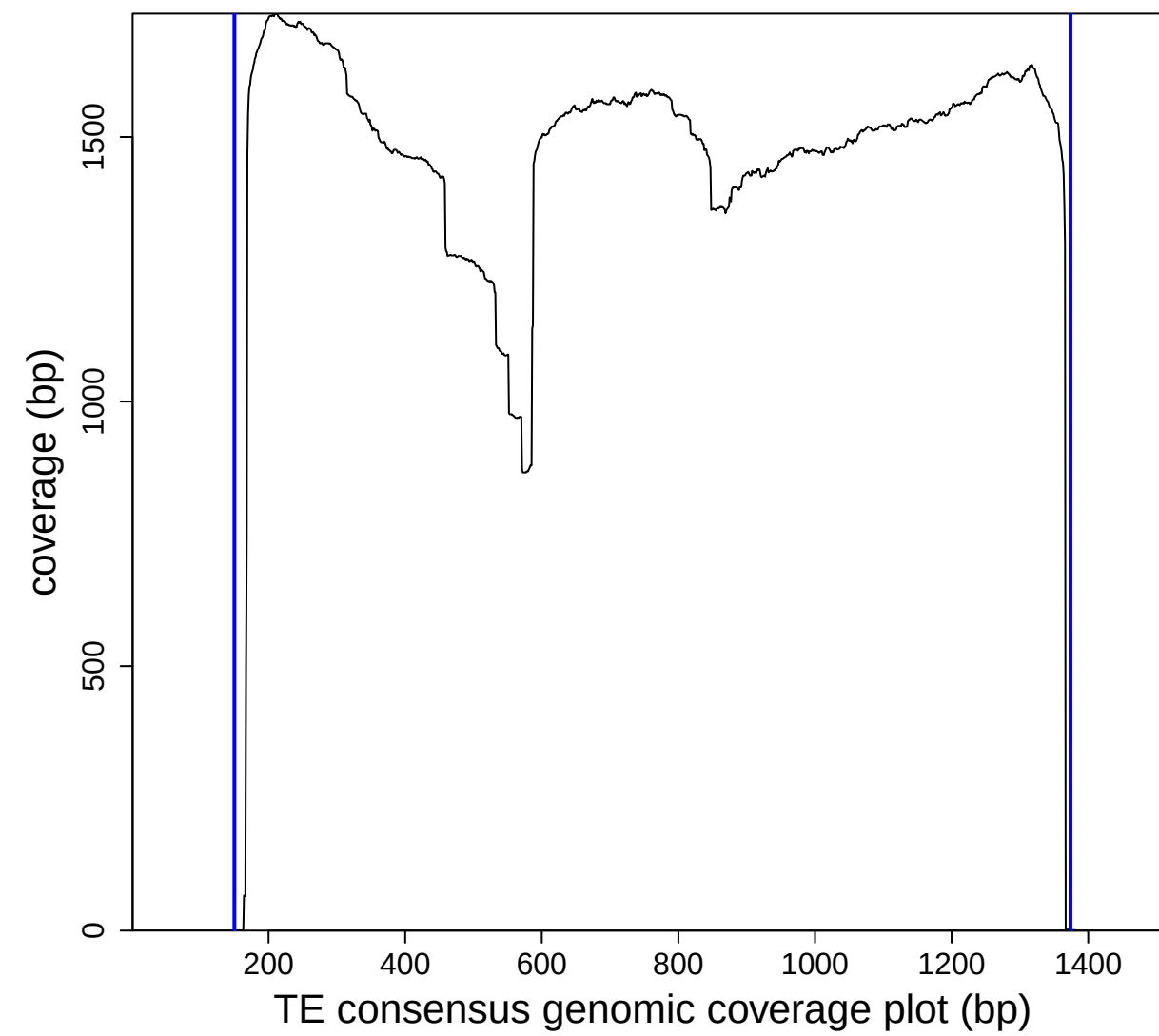
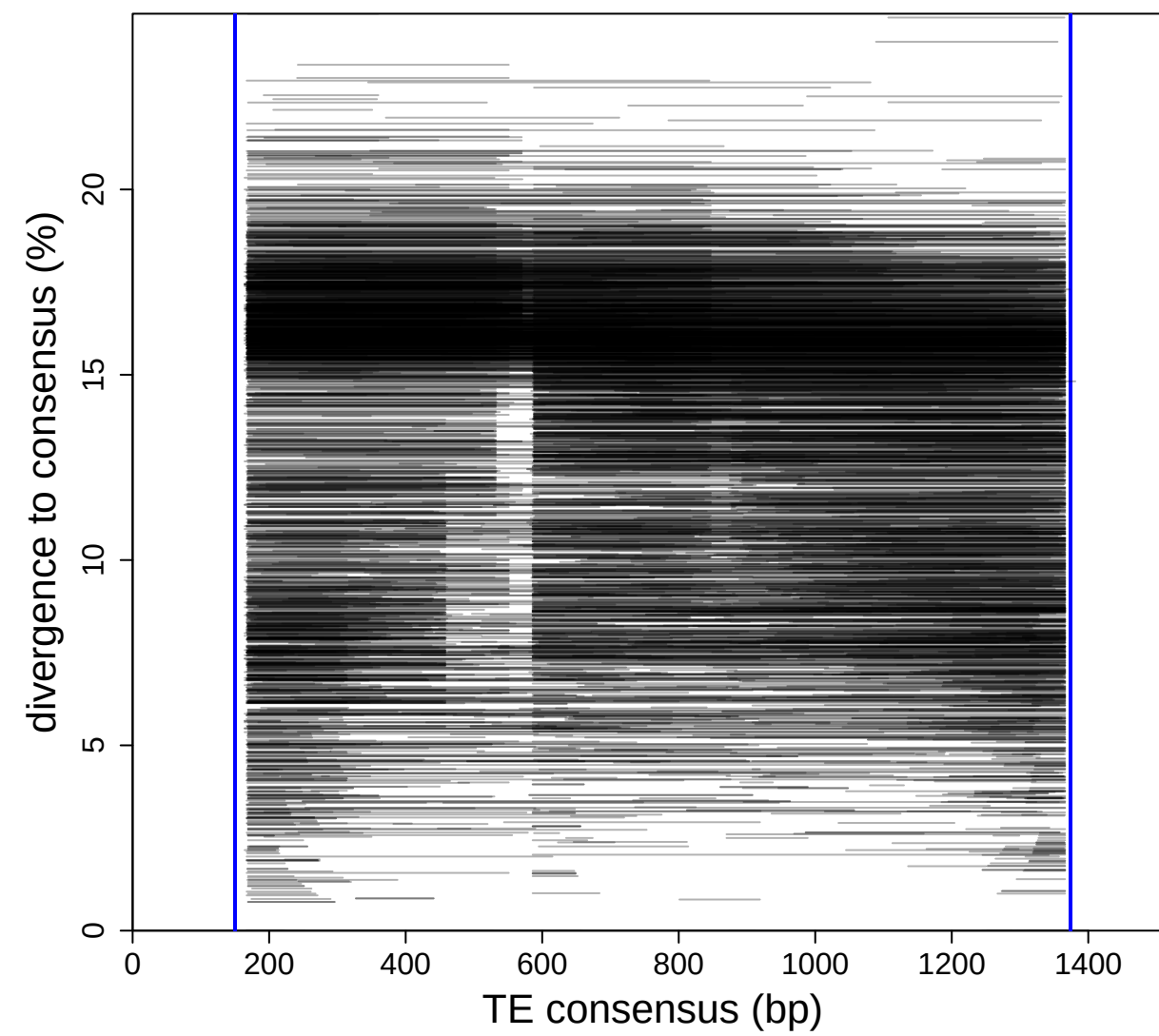
b.bed uf.bed g_3.bed fm_1.bed 0_0 bcln.fa aln.fa cl.fa gs.fa
size: 1223bp; fragments: 3761; full length: 435 (≥ 1100.7 bp)

After TEtrimmer 1223 bp



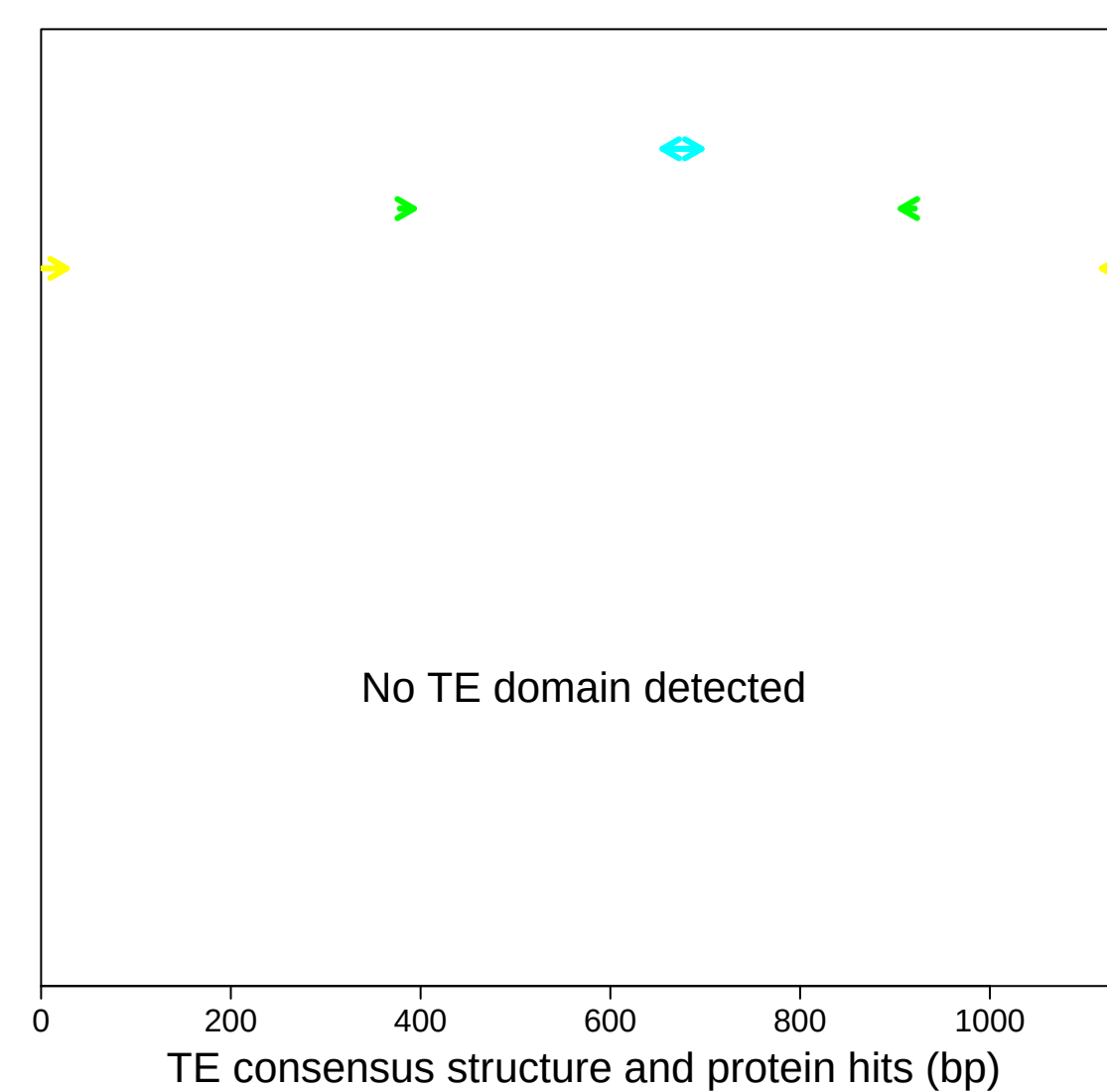
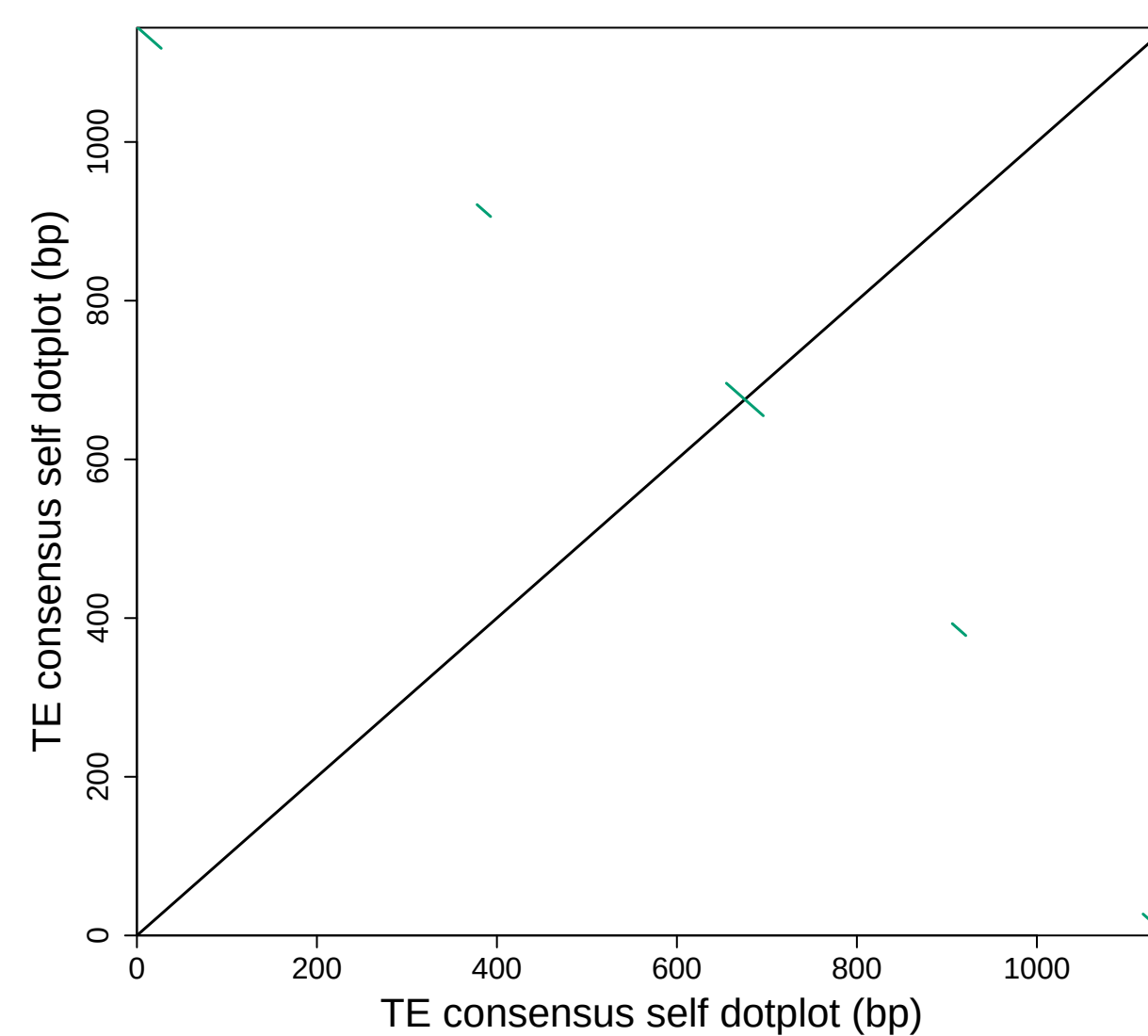
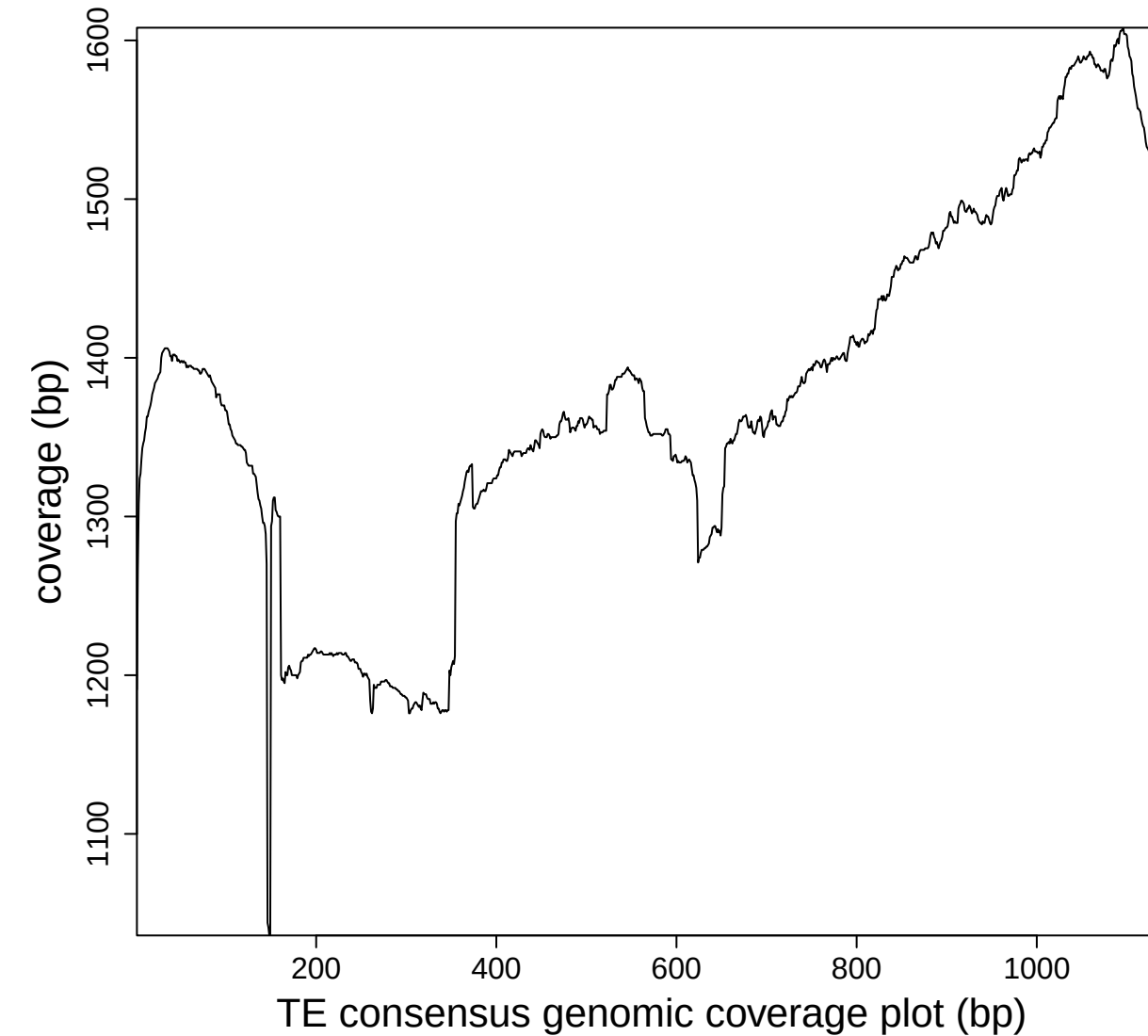
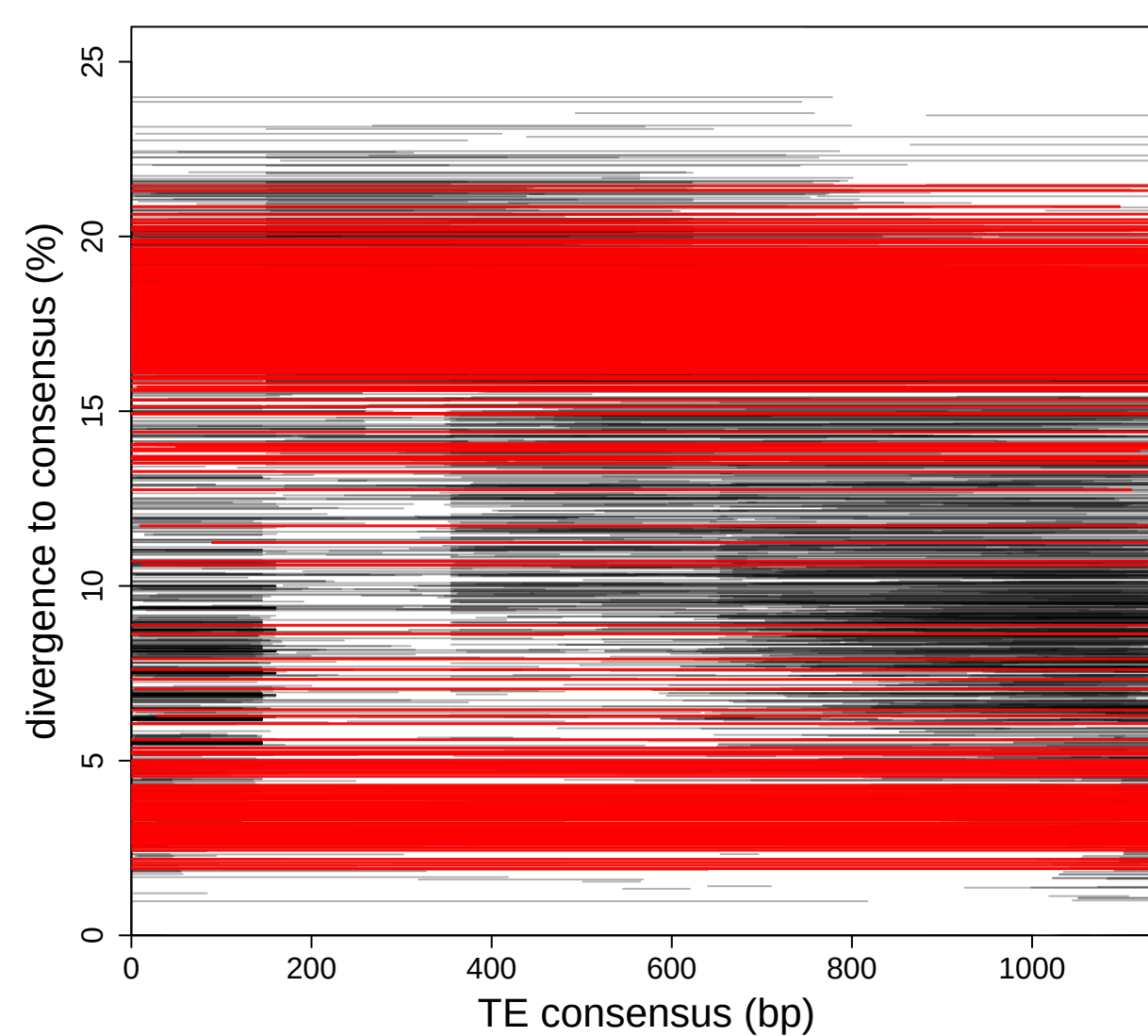
l.fasta.b.bed_uf.bed_g_3.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa
size: 1524bp; fragments: 3759; full length: 0 (>=1371.6bp)

After TEtrimmer Extended plot Blue lines are boundaries

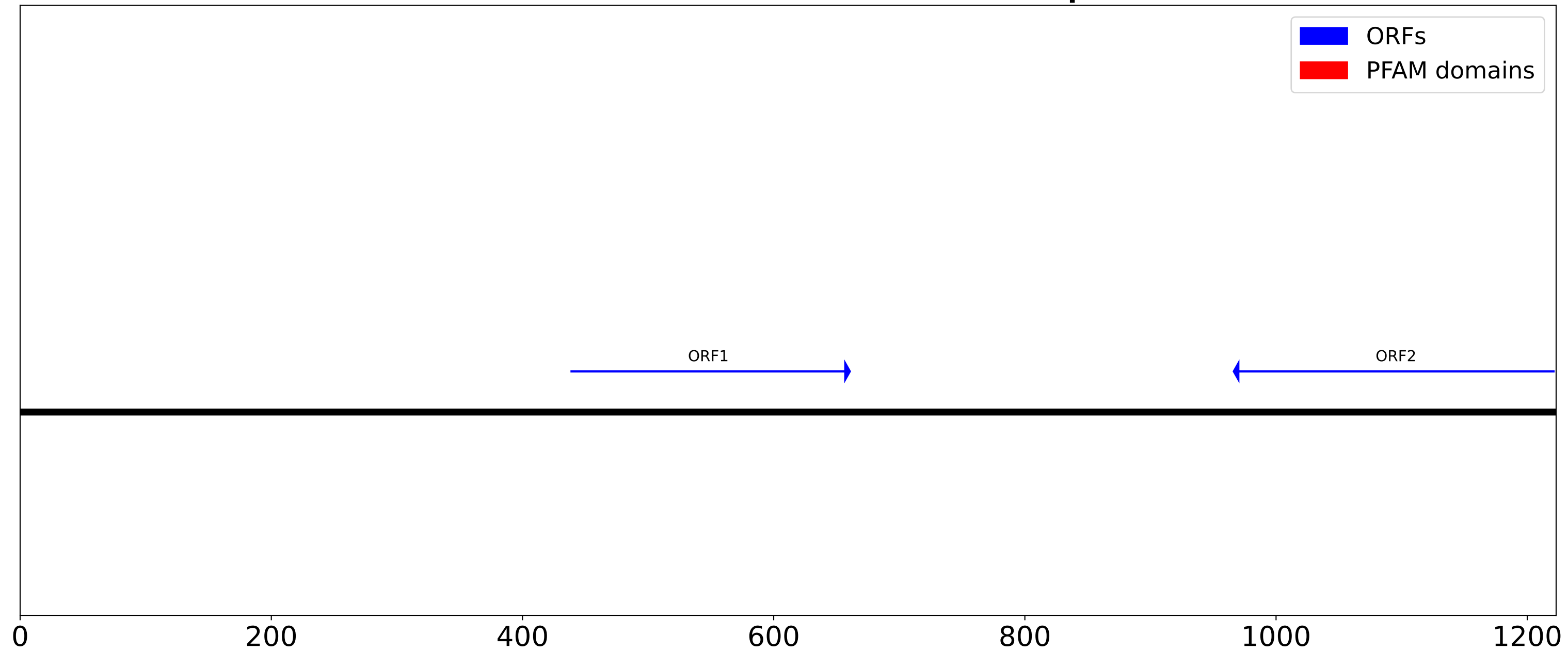


TE: rnd_1_family_92
size: 1144bp; fragments: 3085; full length: 480 (≥ 1029.6 bp)

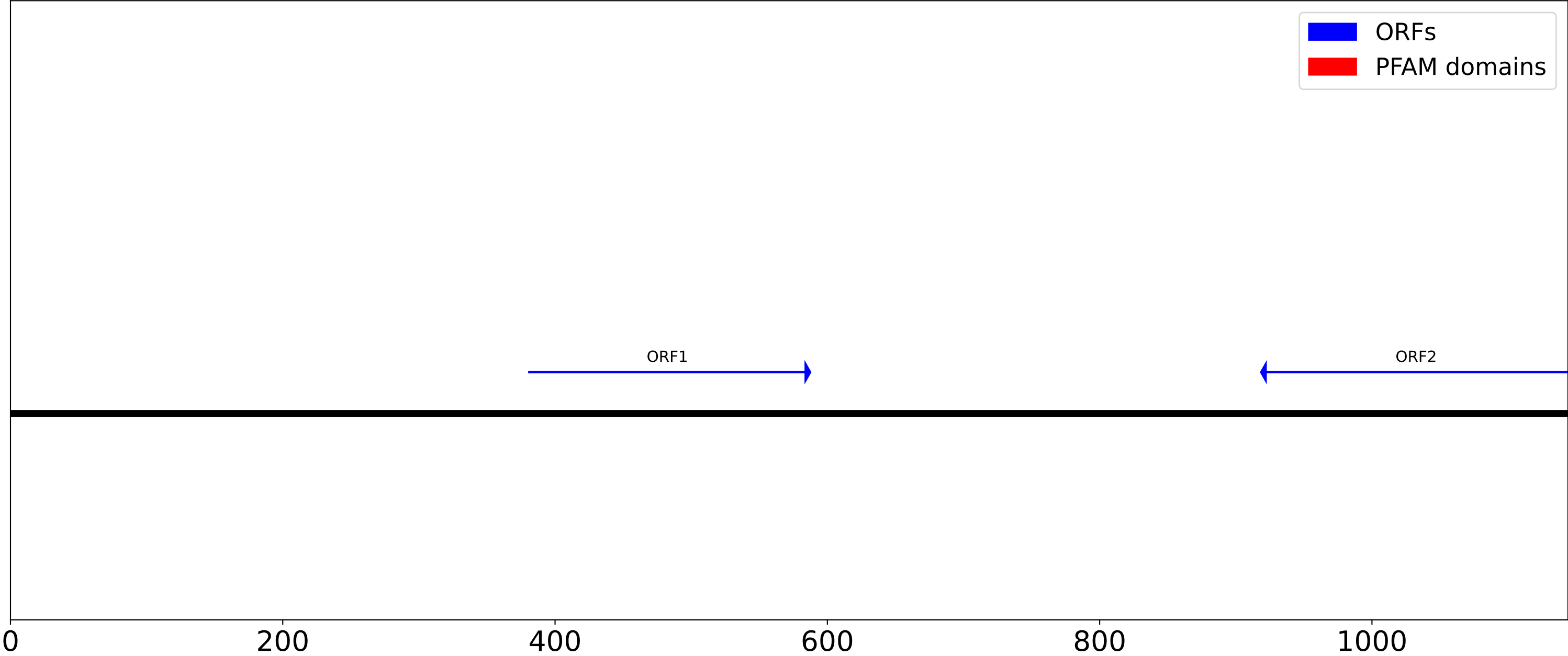
Before TEtrimmer 1144 bp



After TEtrimmer ORF and PFAM domain plot



Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

