

Start crop Point



End crop Point



MSA length = 5791



Start crop Point



End crop Point



0

10

0

1000

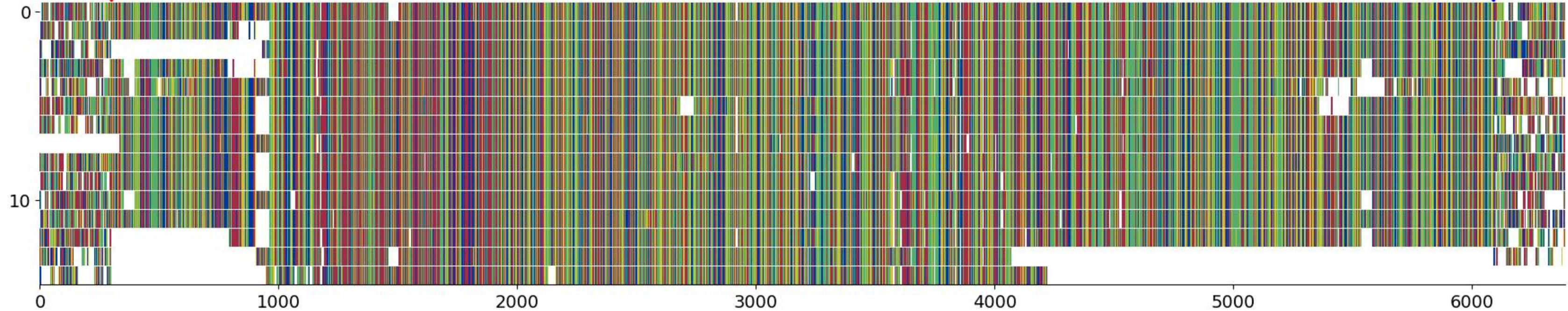
2000

3000

4000

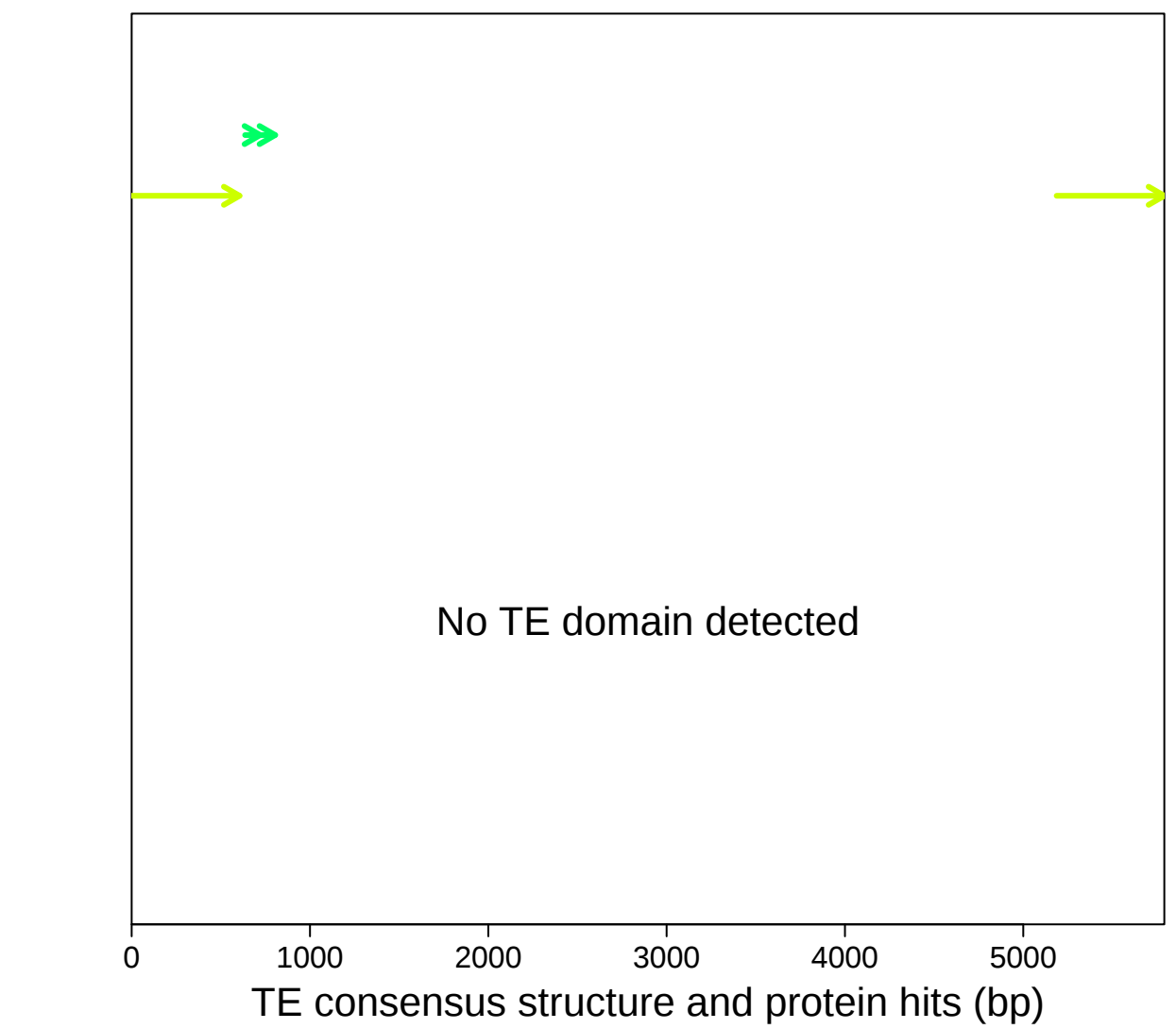
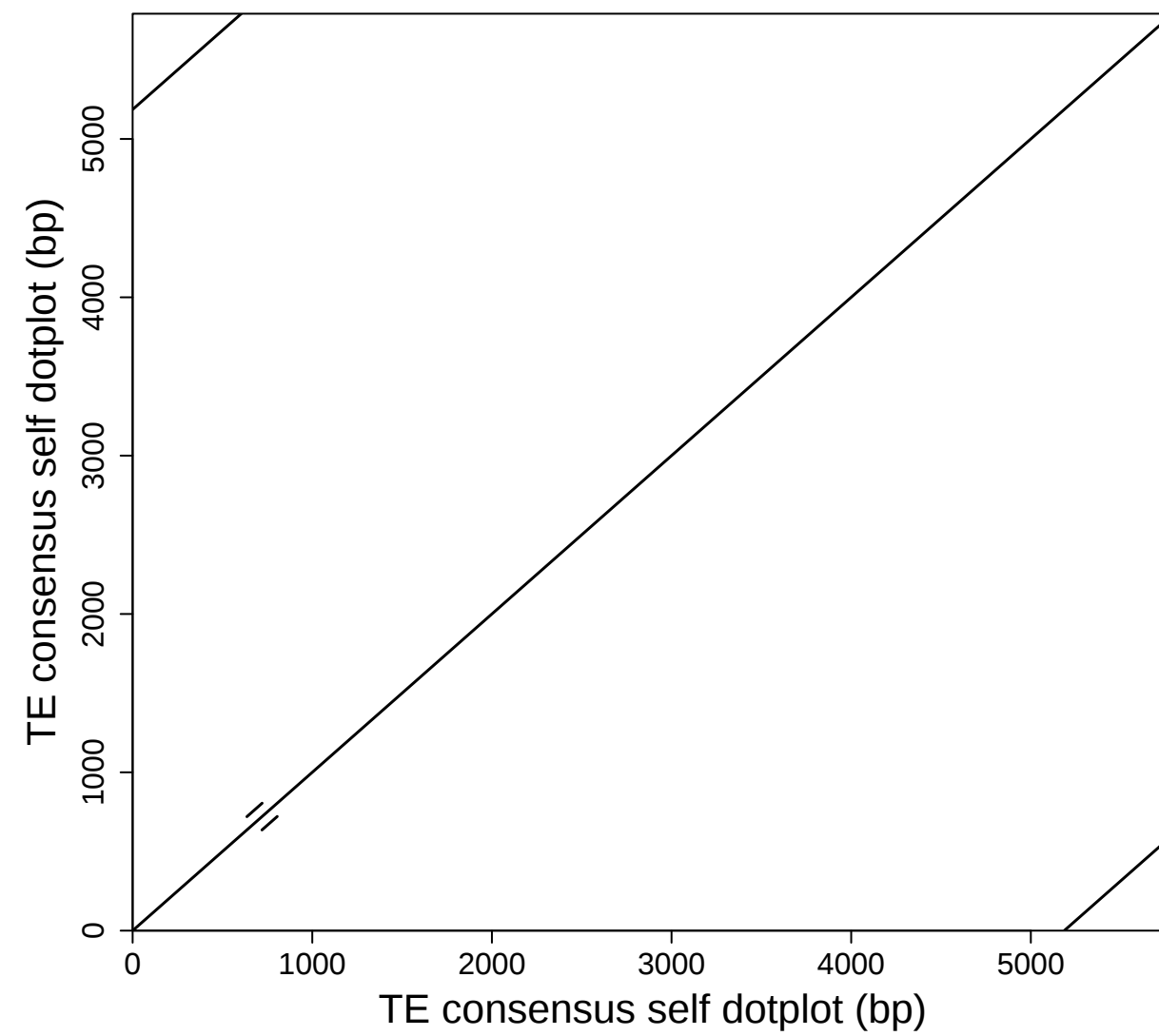
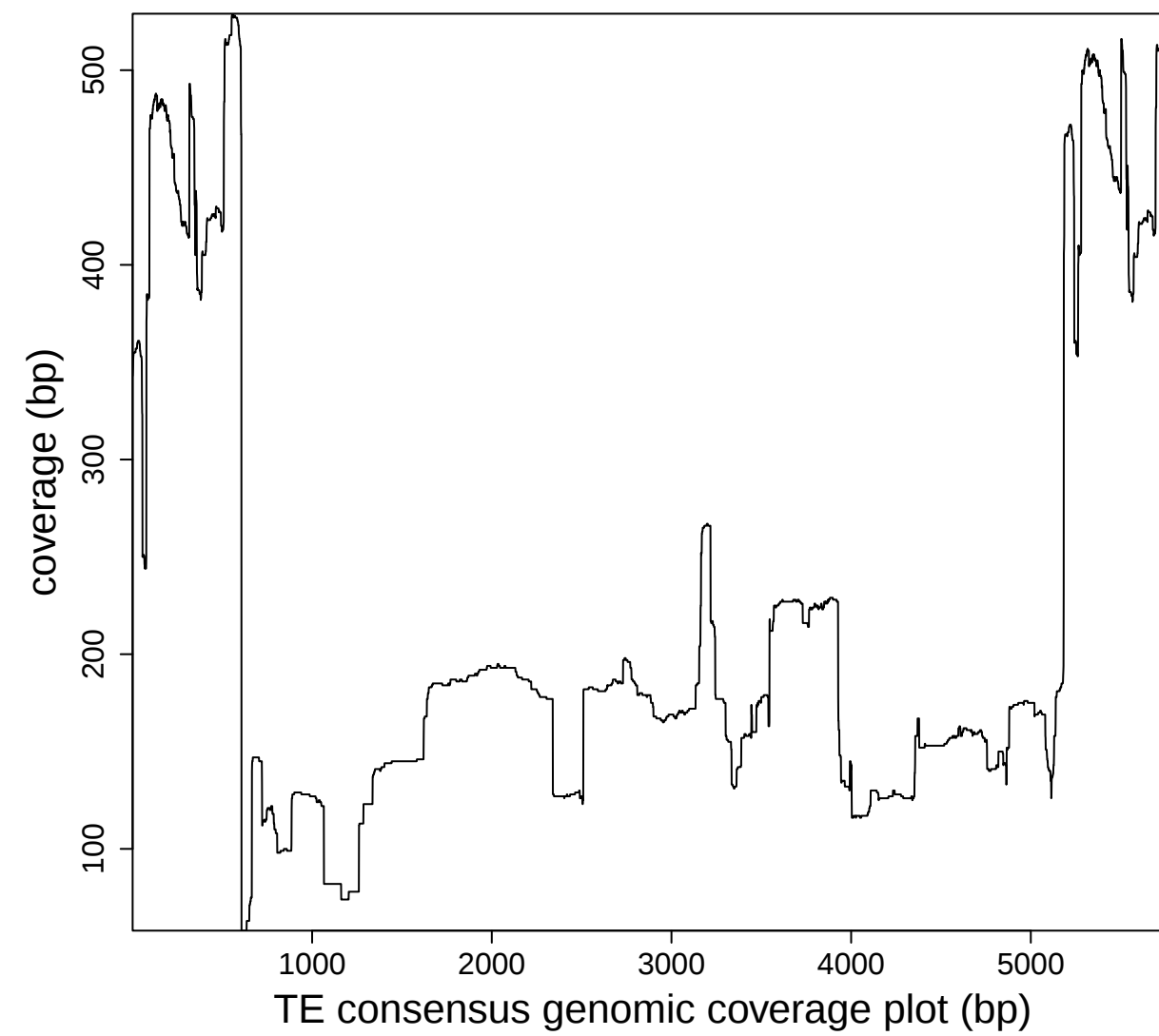
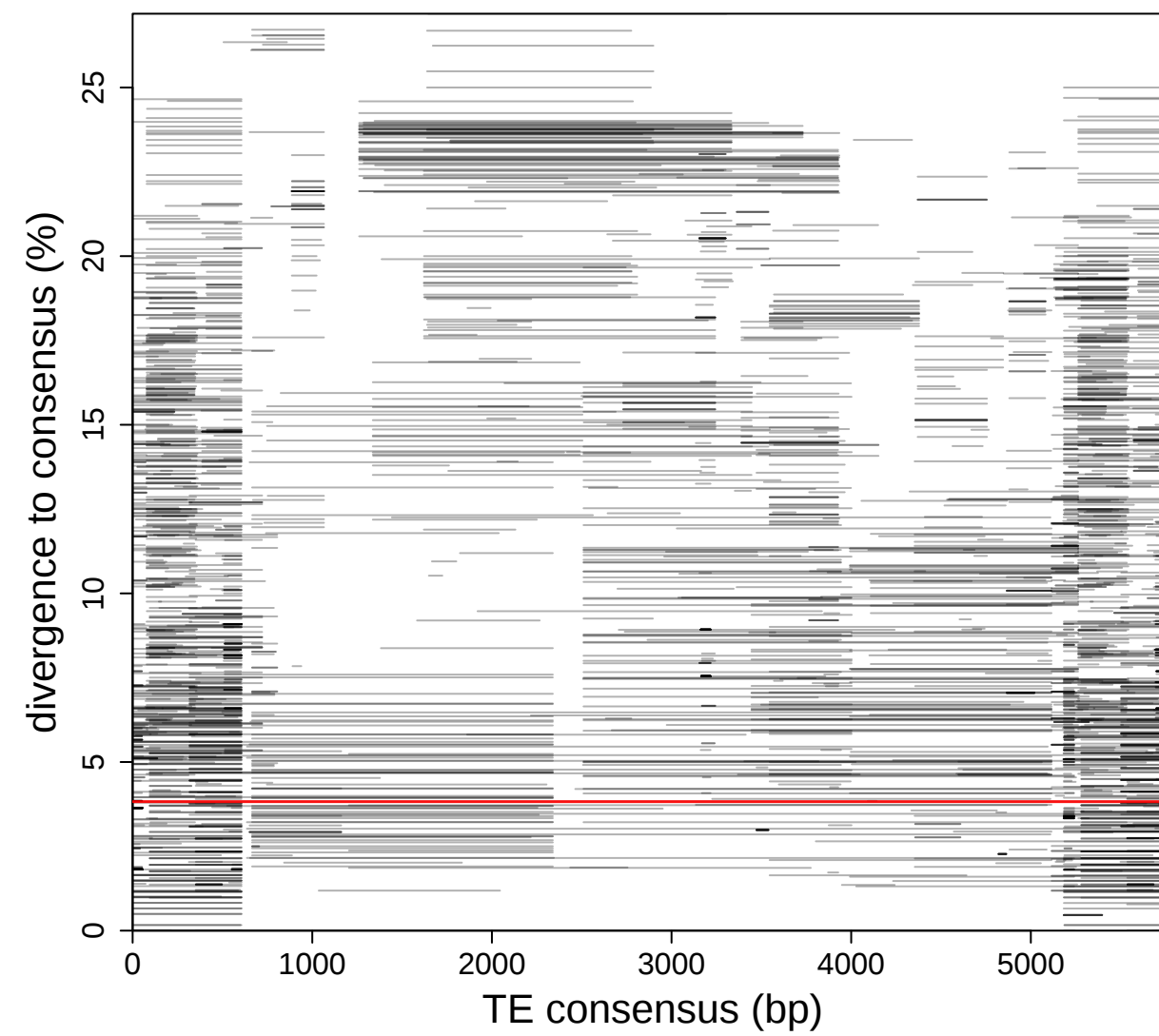
5000

6000



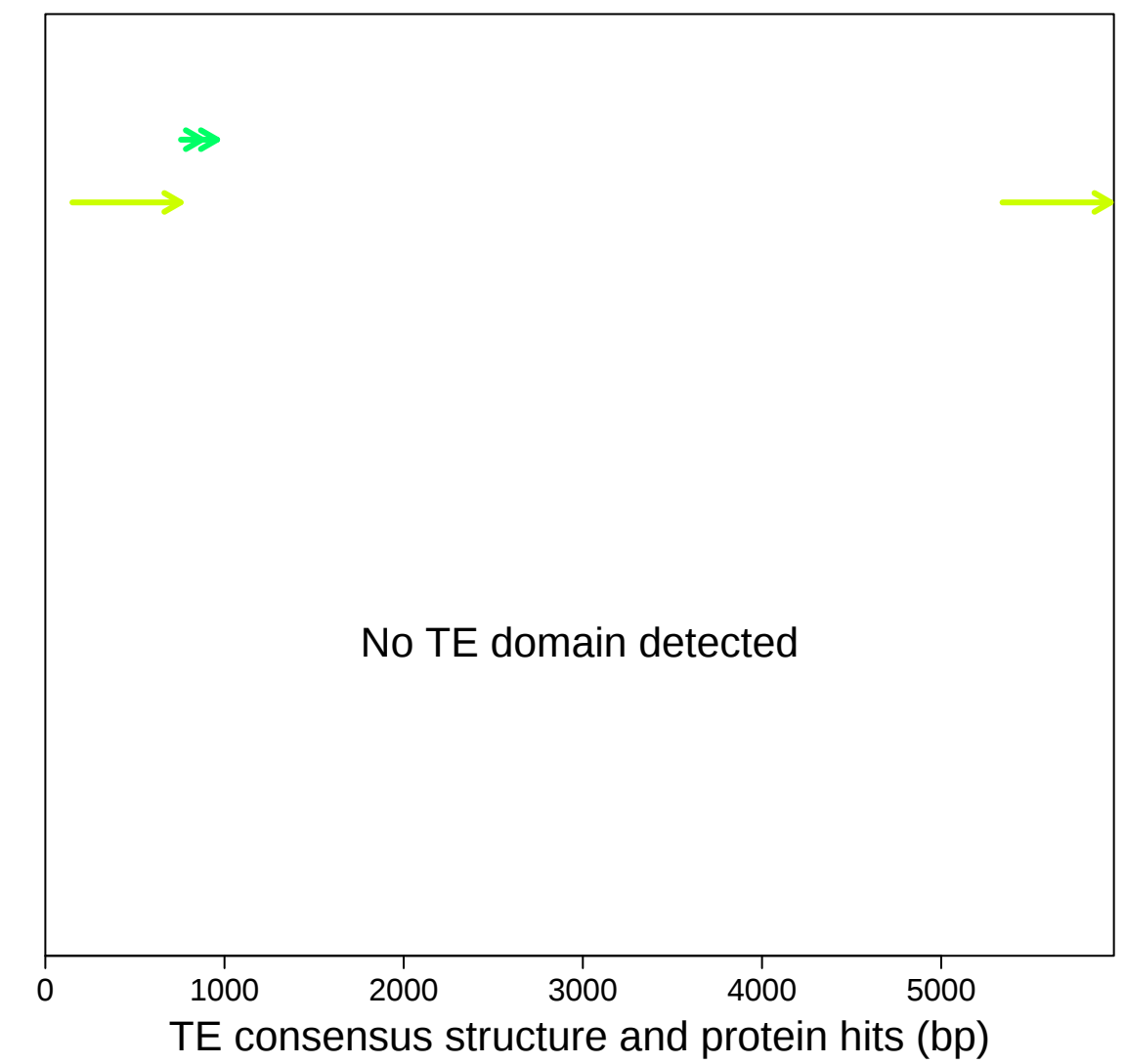
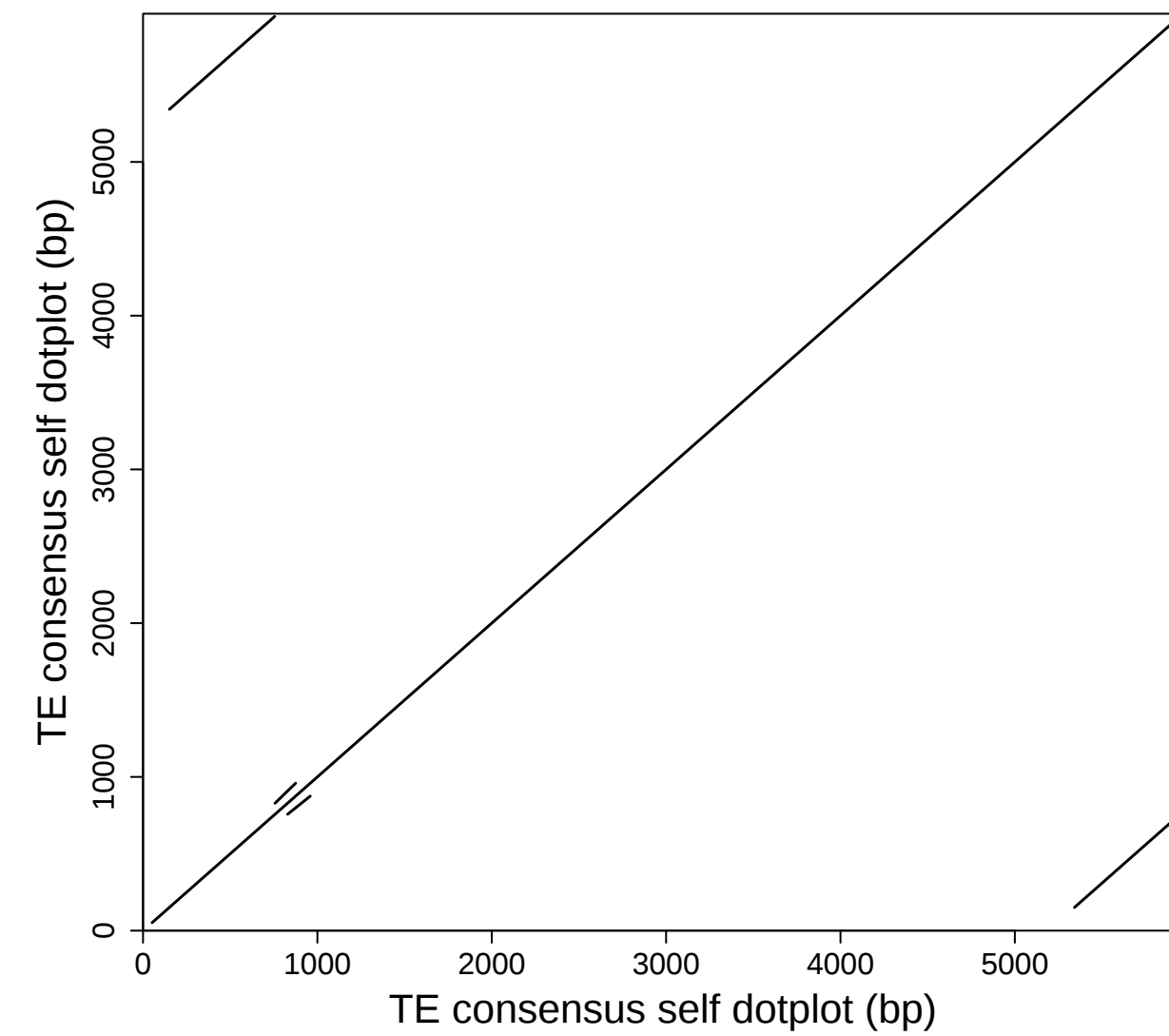
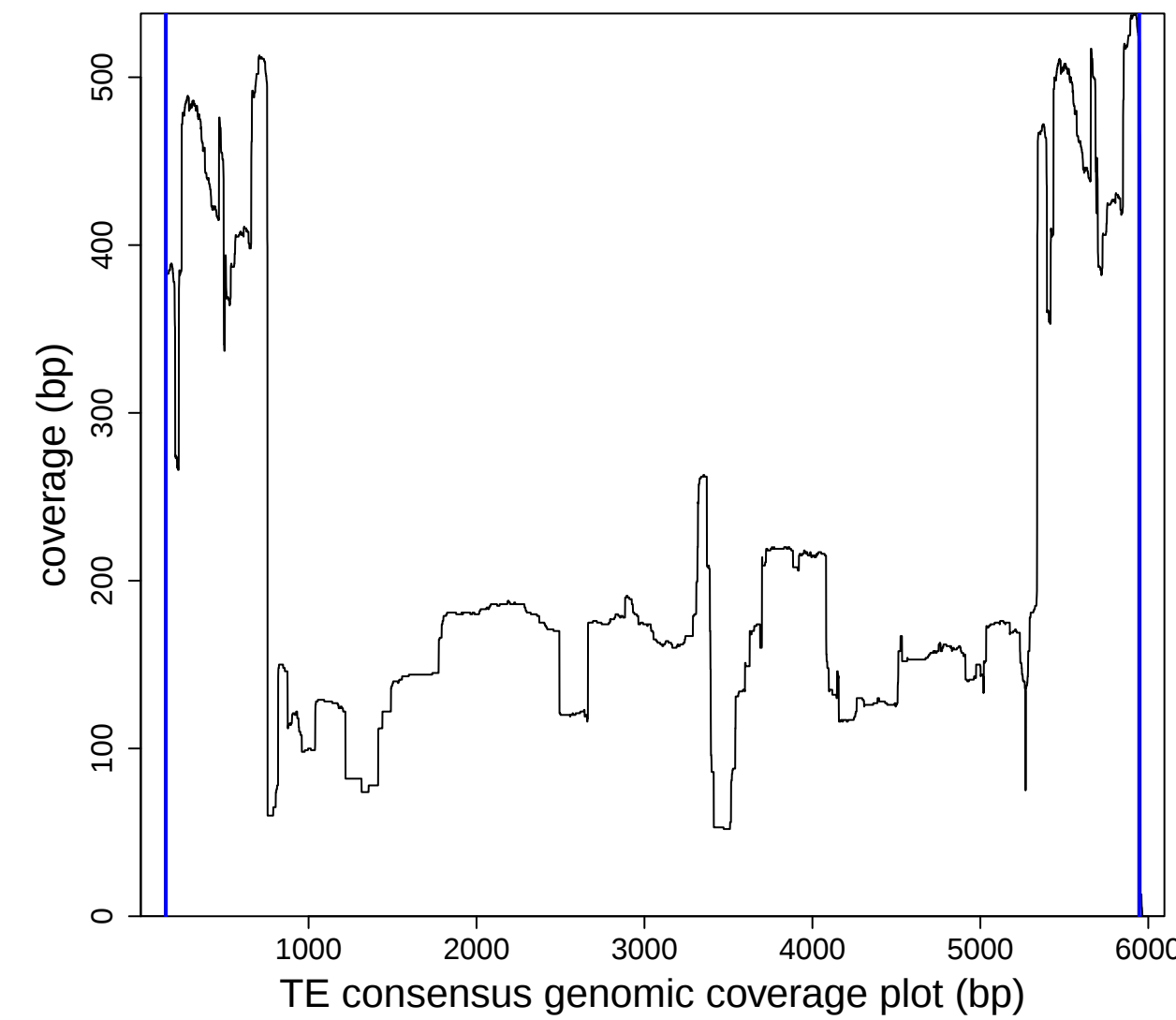
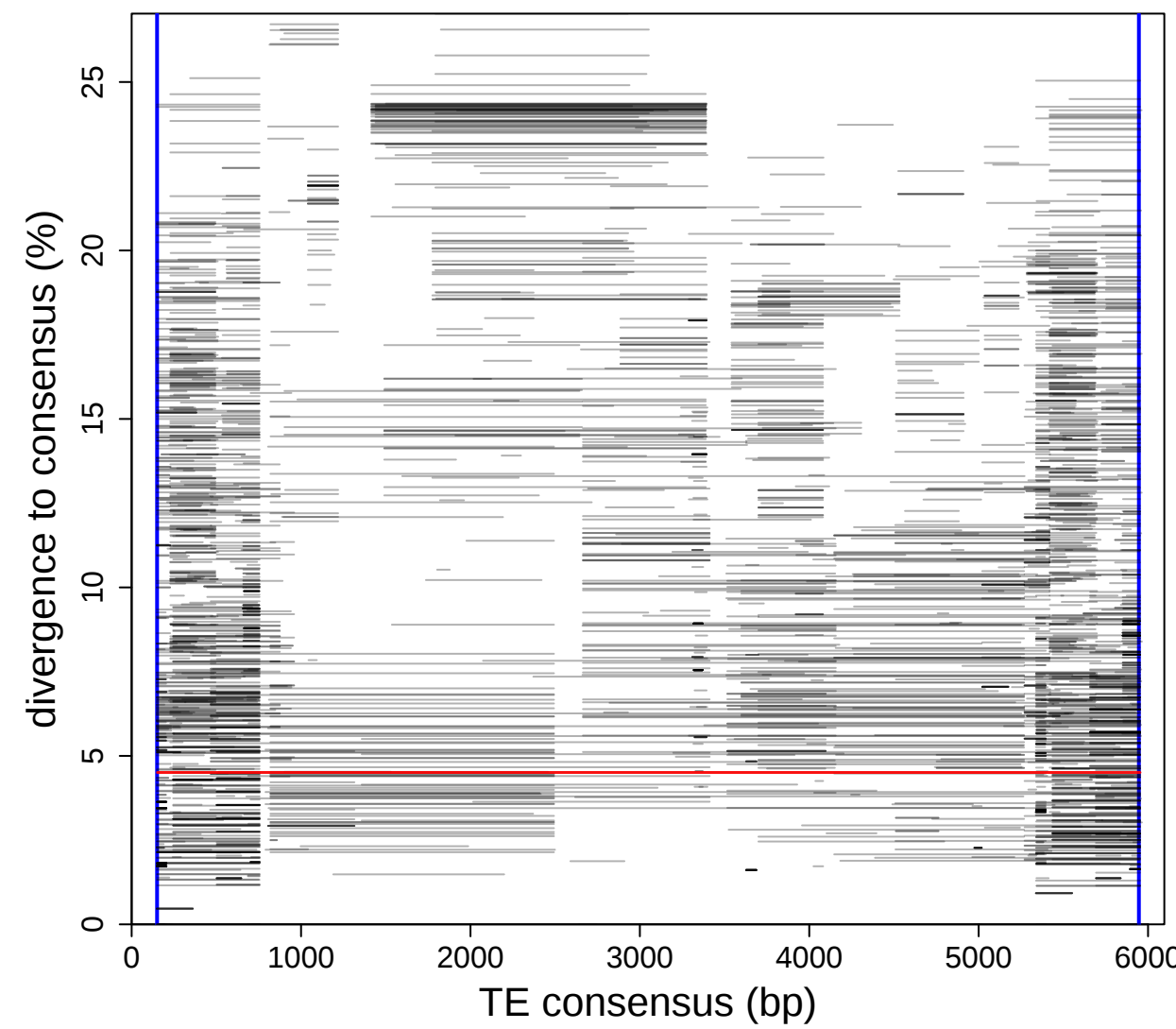
ed_uf.bed_g_2.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.fa_ce.
size: 5791bp; fragments: 2792; full length: 1 (>=5211.9bp)

After TEtrimmer 5791 bp



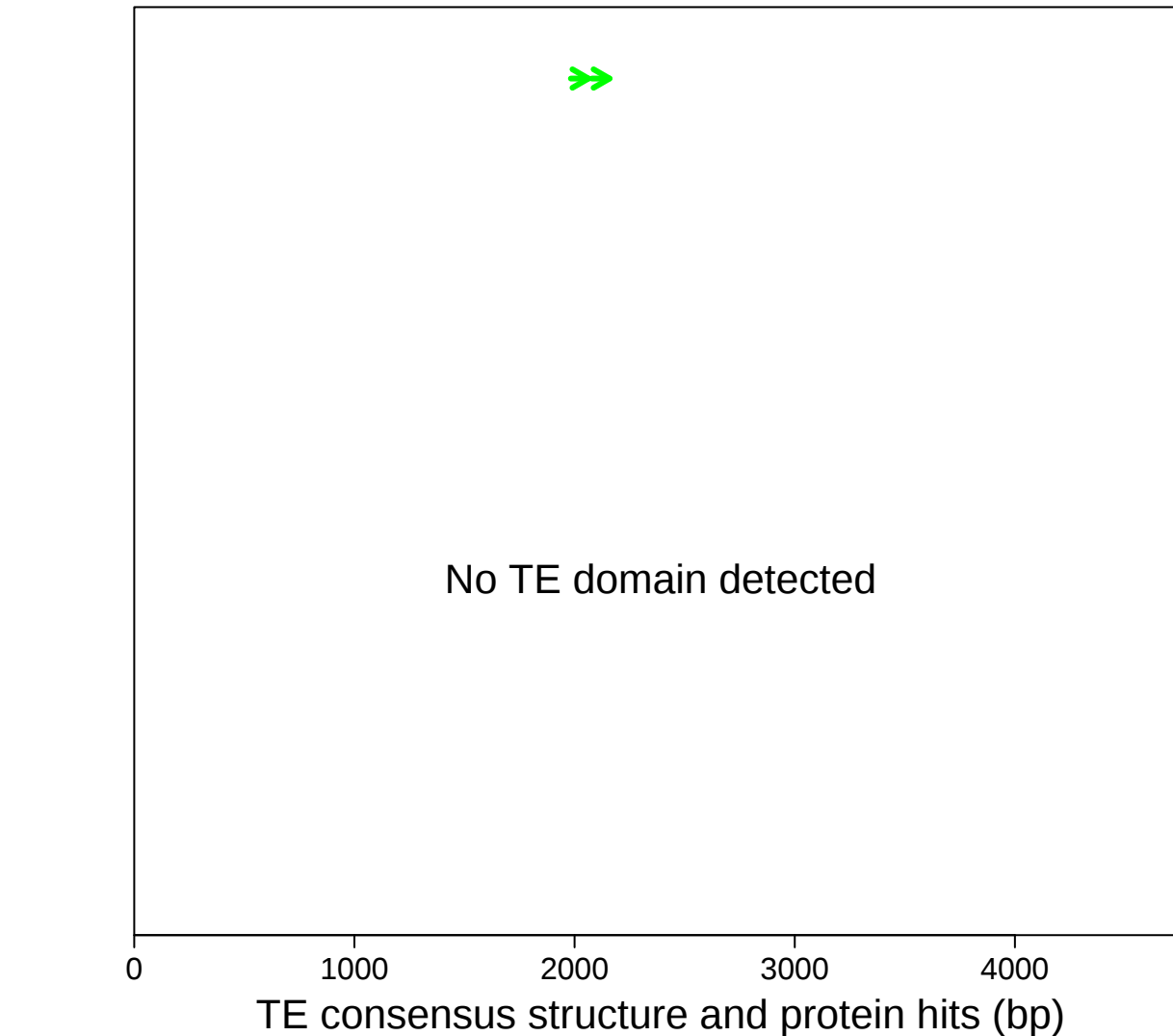
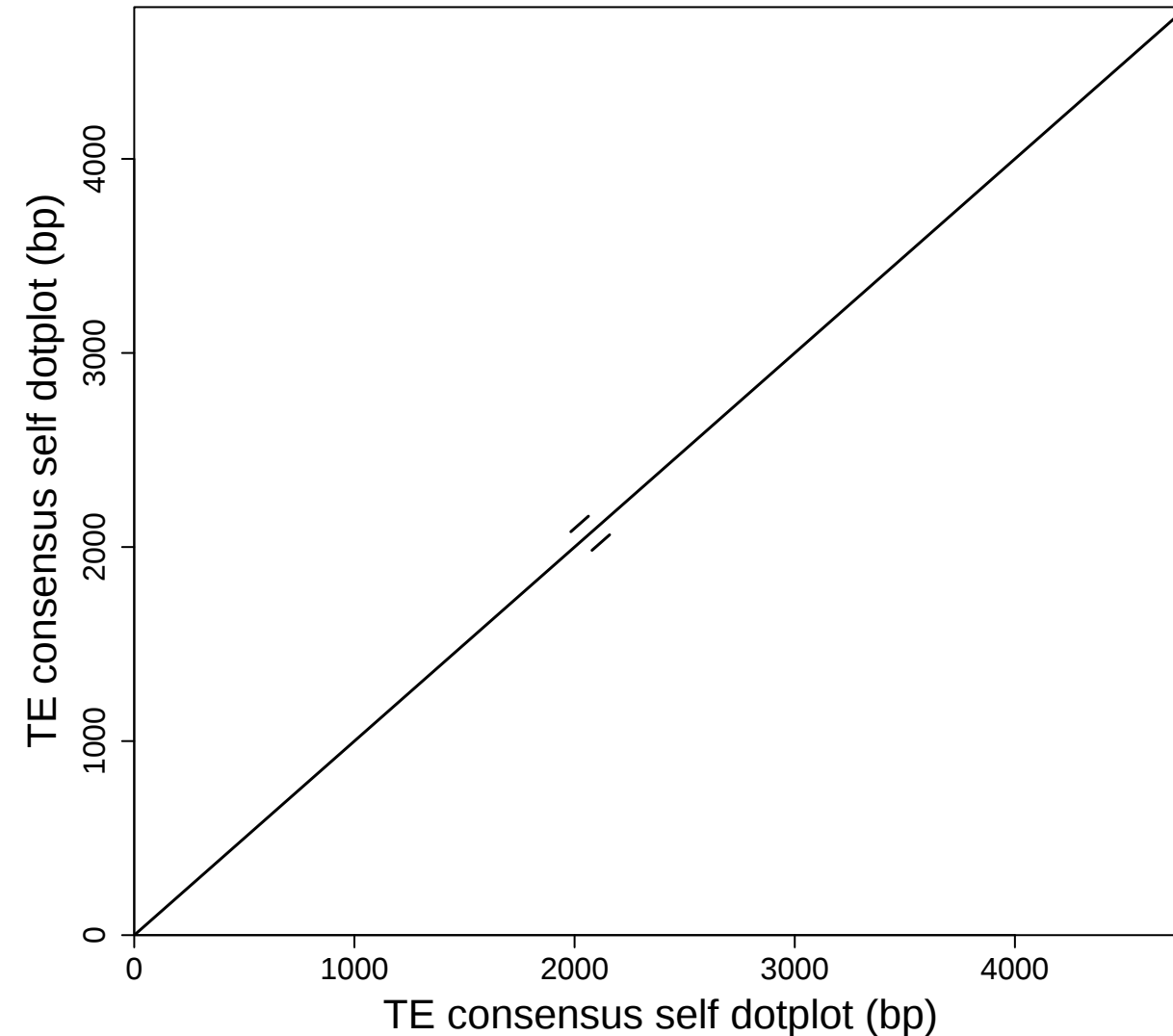
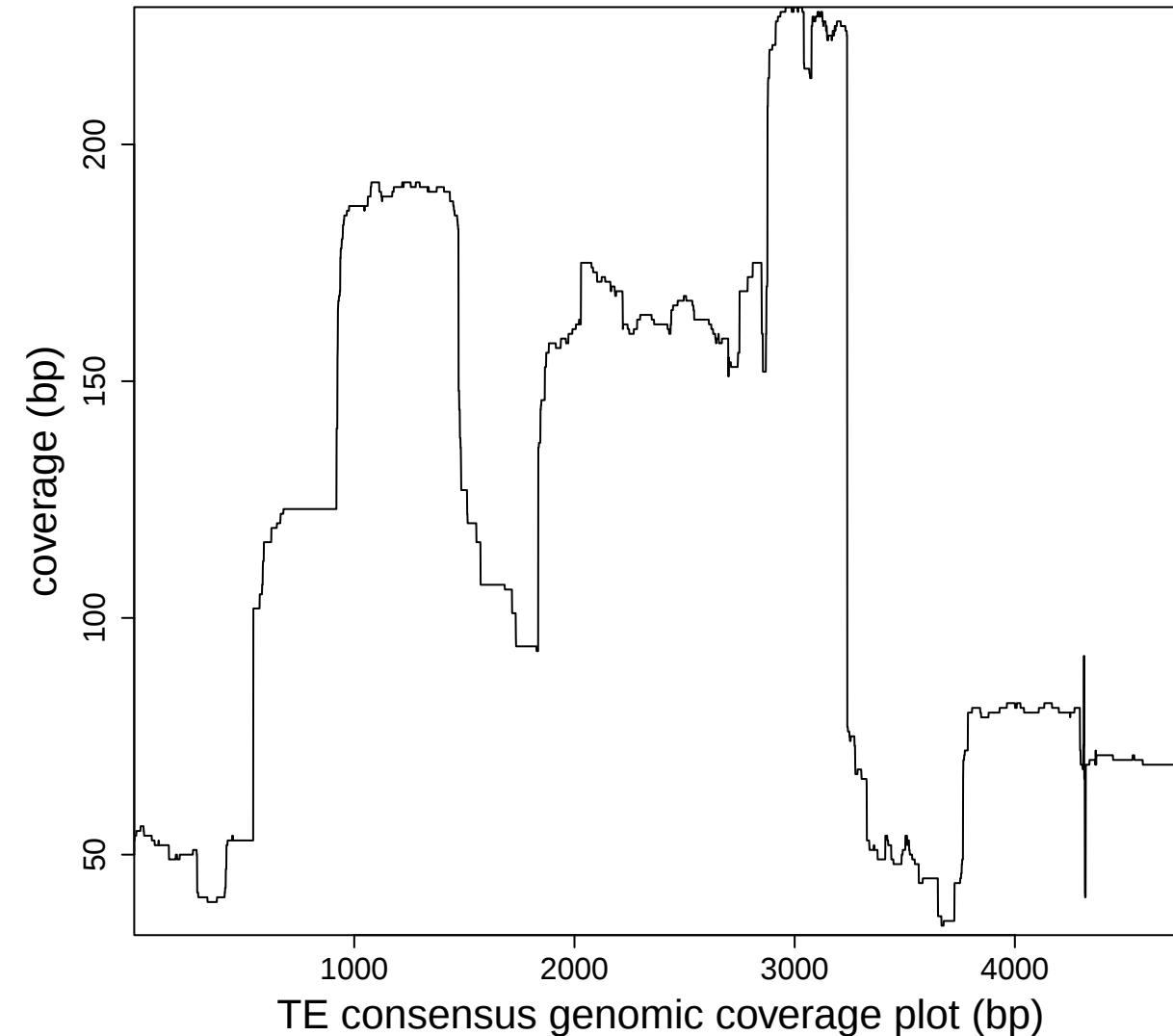
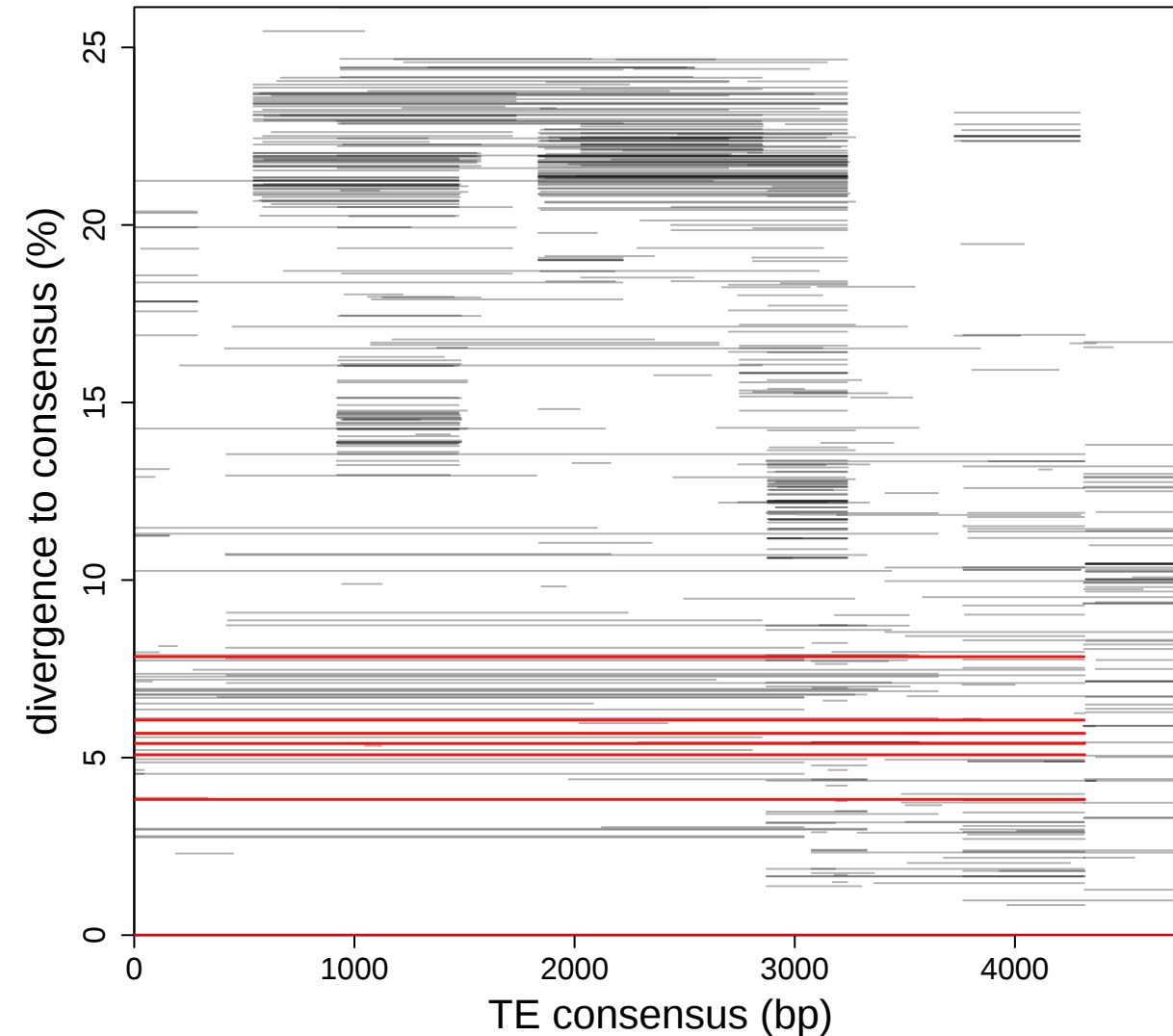
ly_852.fasta.b.bed_uf.bed_g_2.bed_fm_1.bed_0_0_bcln.fa_aln.fa
size: 6096bp; fragments: 2859; full length: 1 (>=5486.4bp)

After TEtrimmer Extended plot Blue lines are boundaries

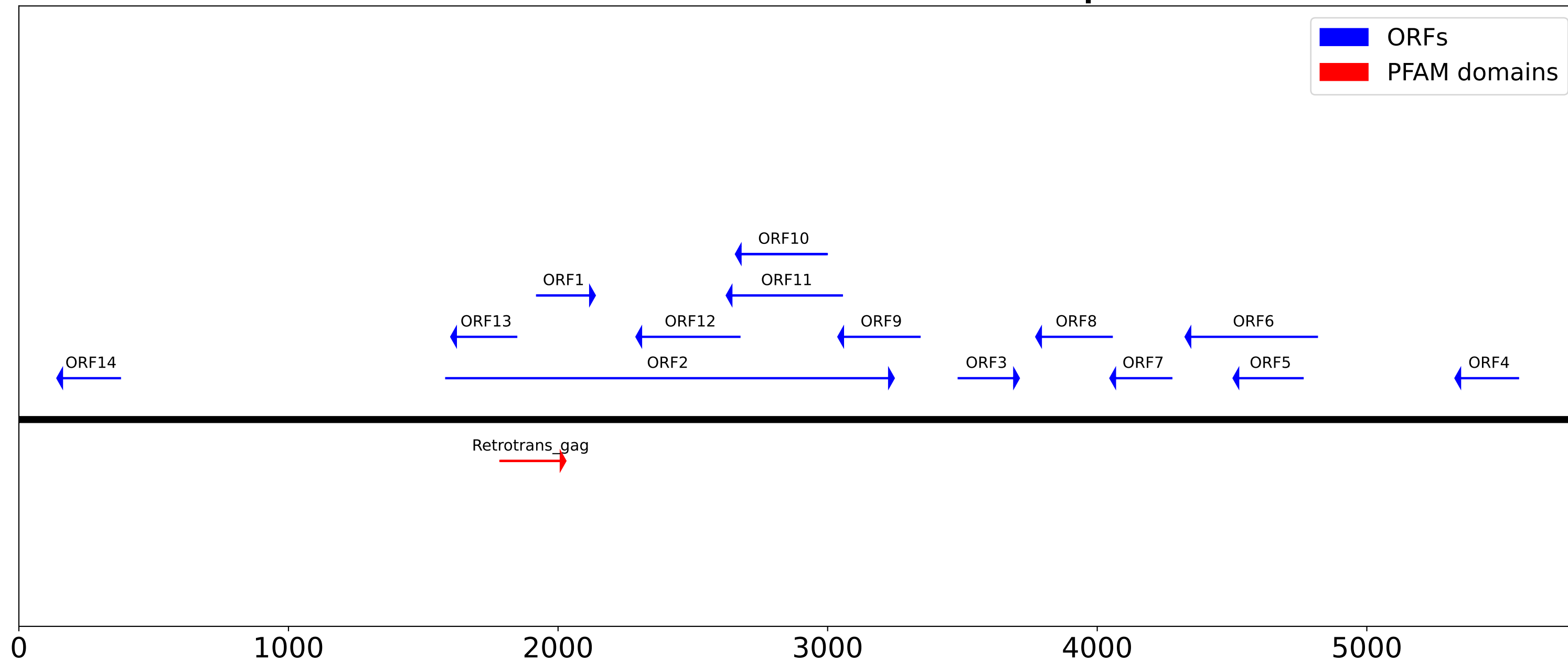


TE: ltr_1_family_852
size: 4782bp; fragments: 592; full length: 7 (≥ 4303.8 bp)

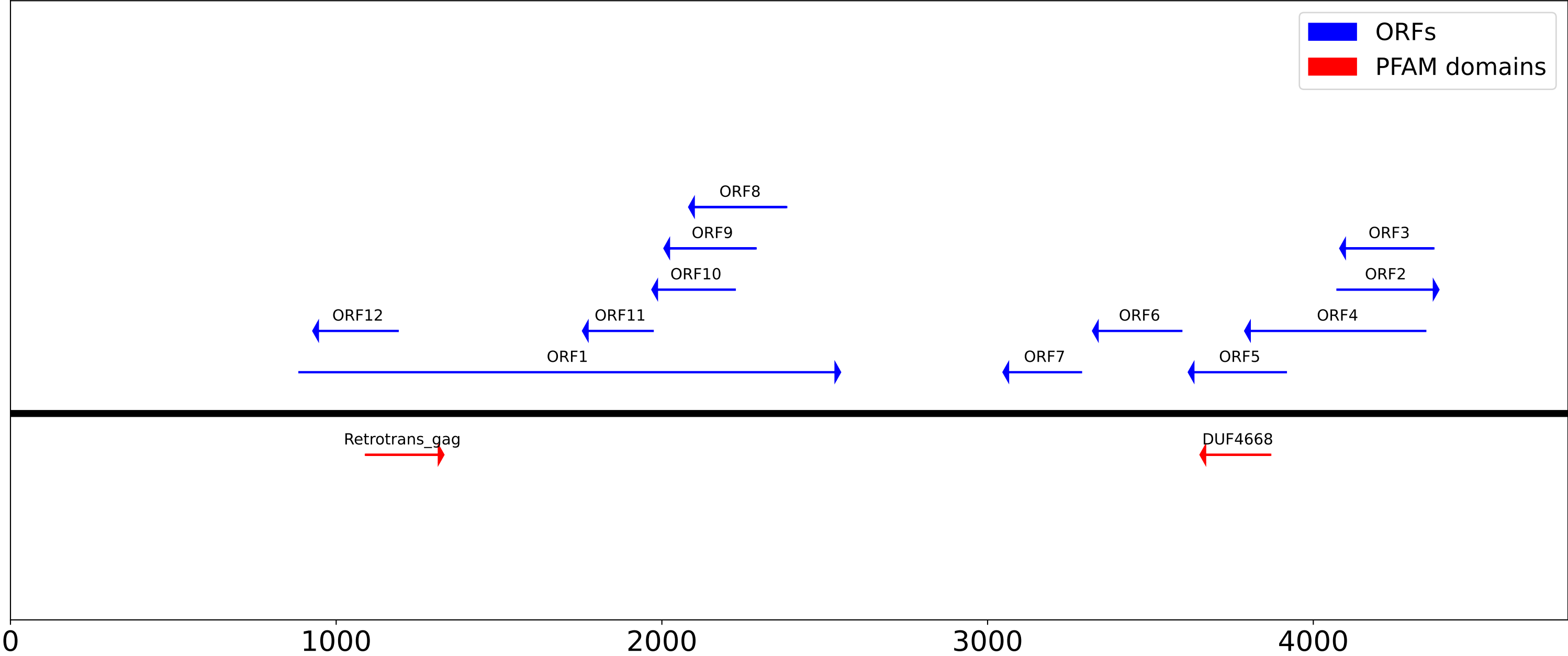
Before TEtrimmer 4782 bp



After TEtrimmer ORF and PFAM domain plot



Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

