

Start crop Point

End crop Point

MSA length = 5827



Start crop Point

End crop Point

0

10

0

1000

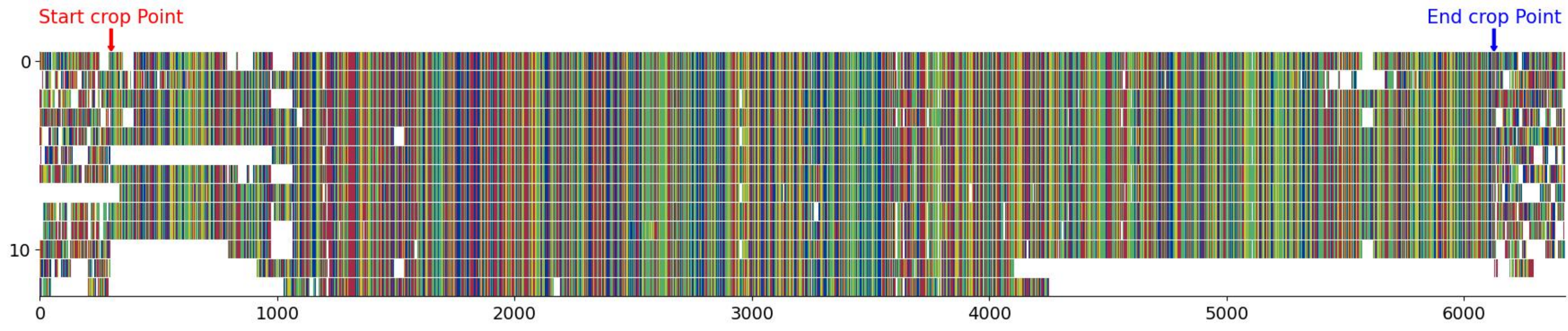
2000

3000

4000

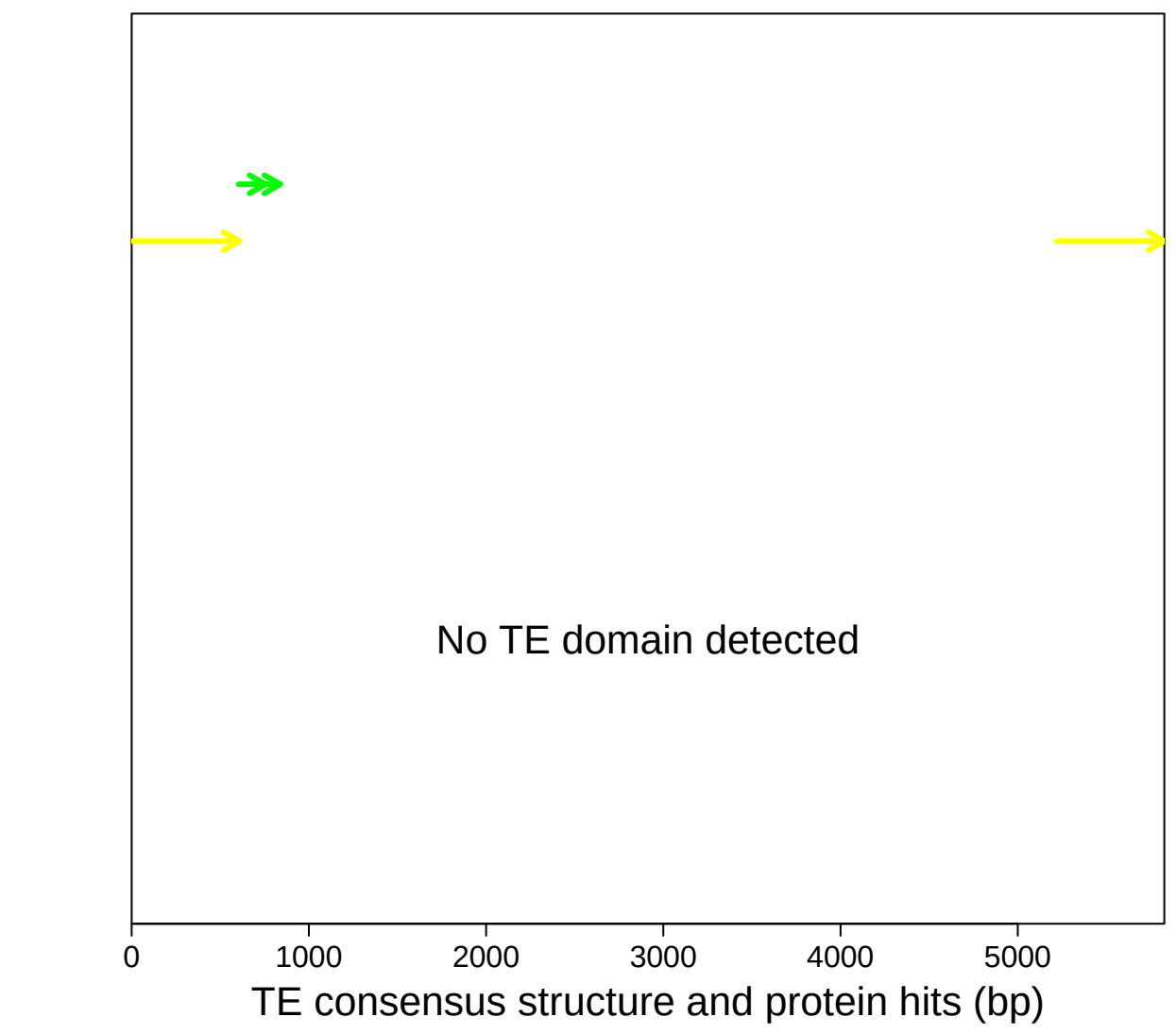
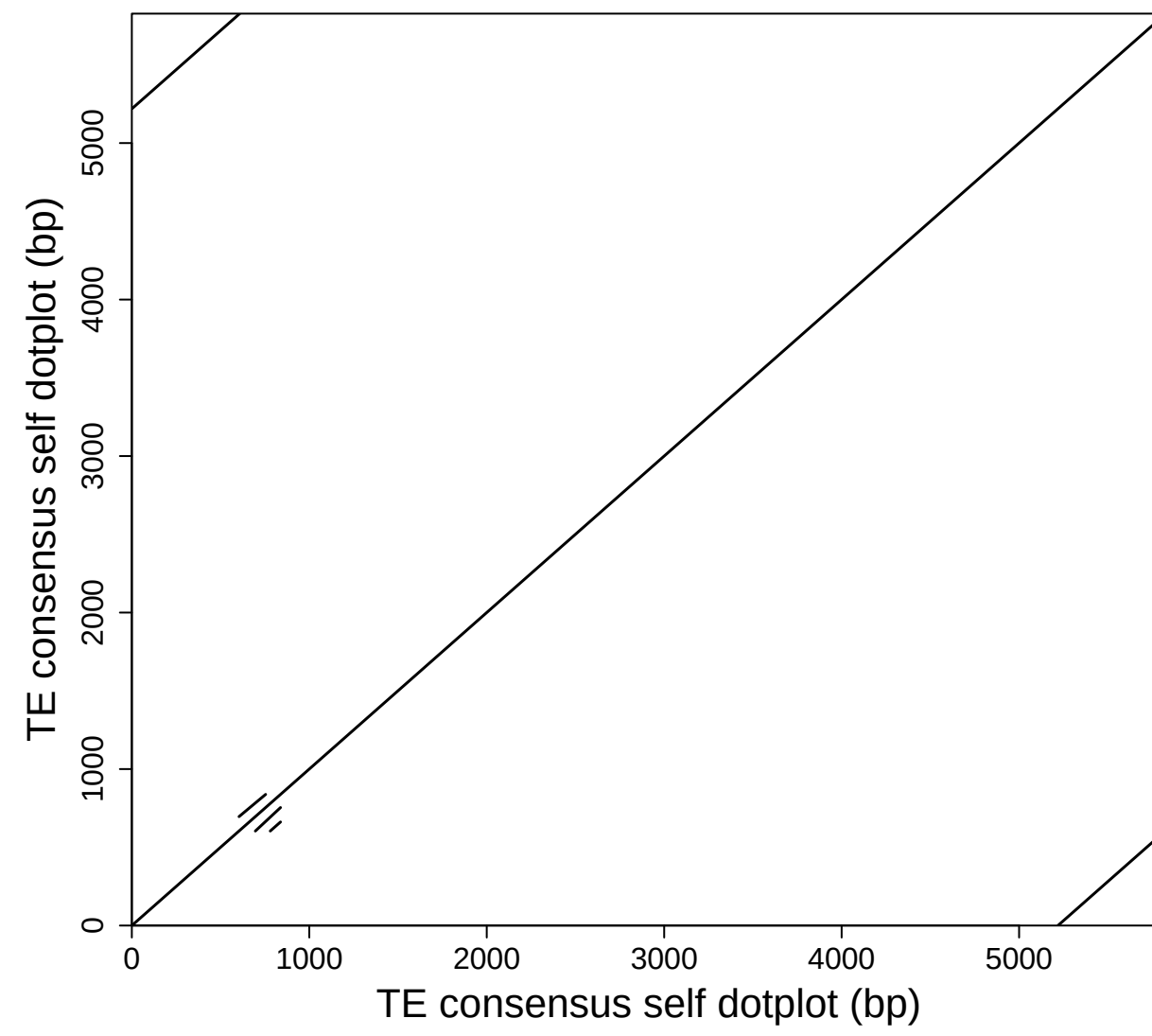
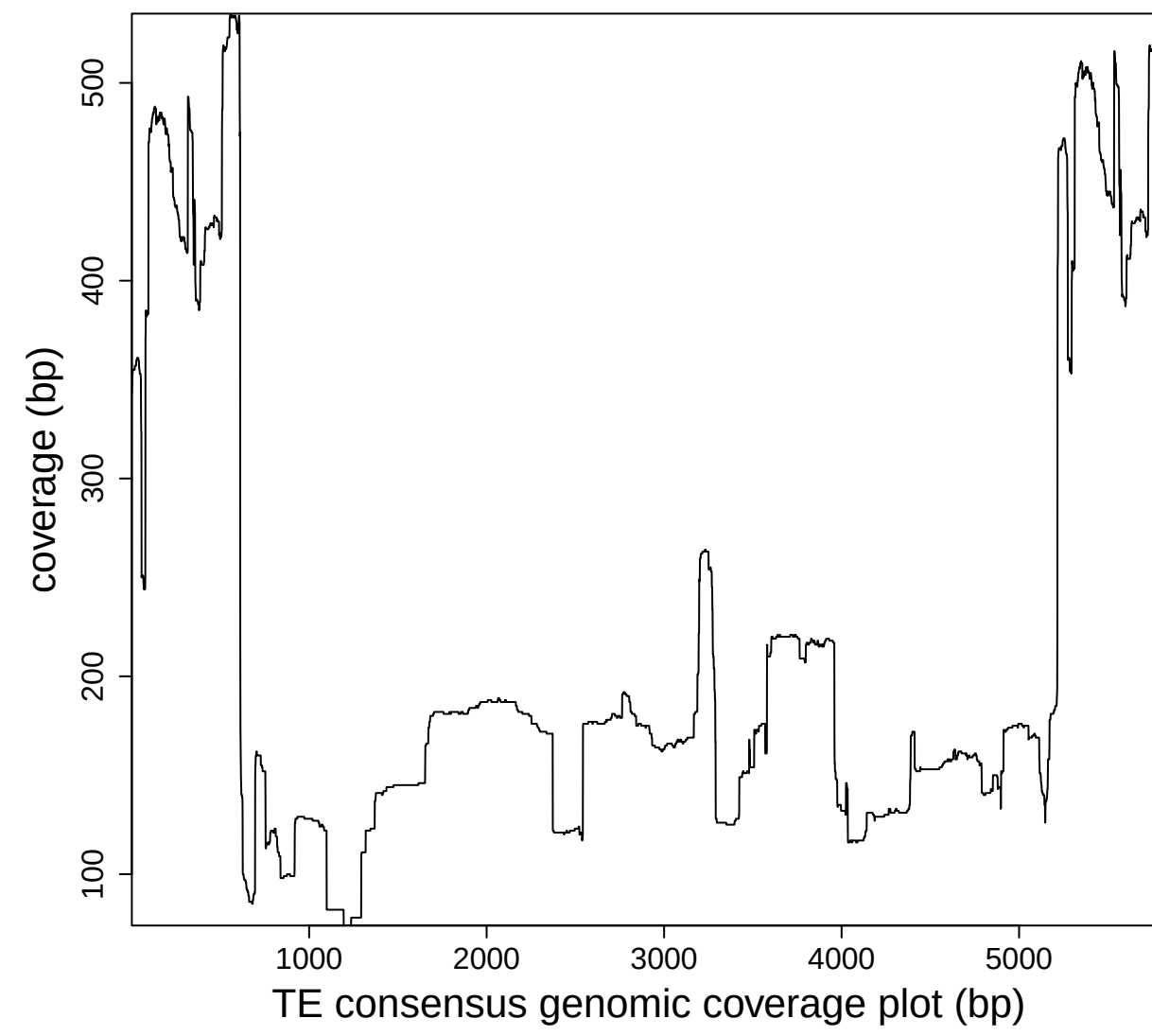
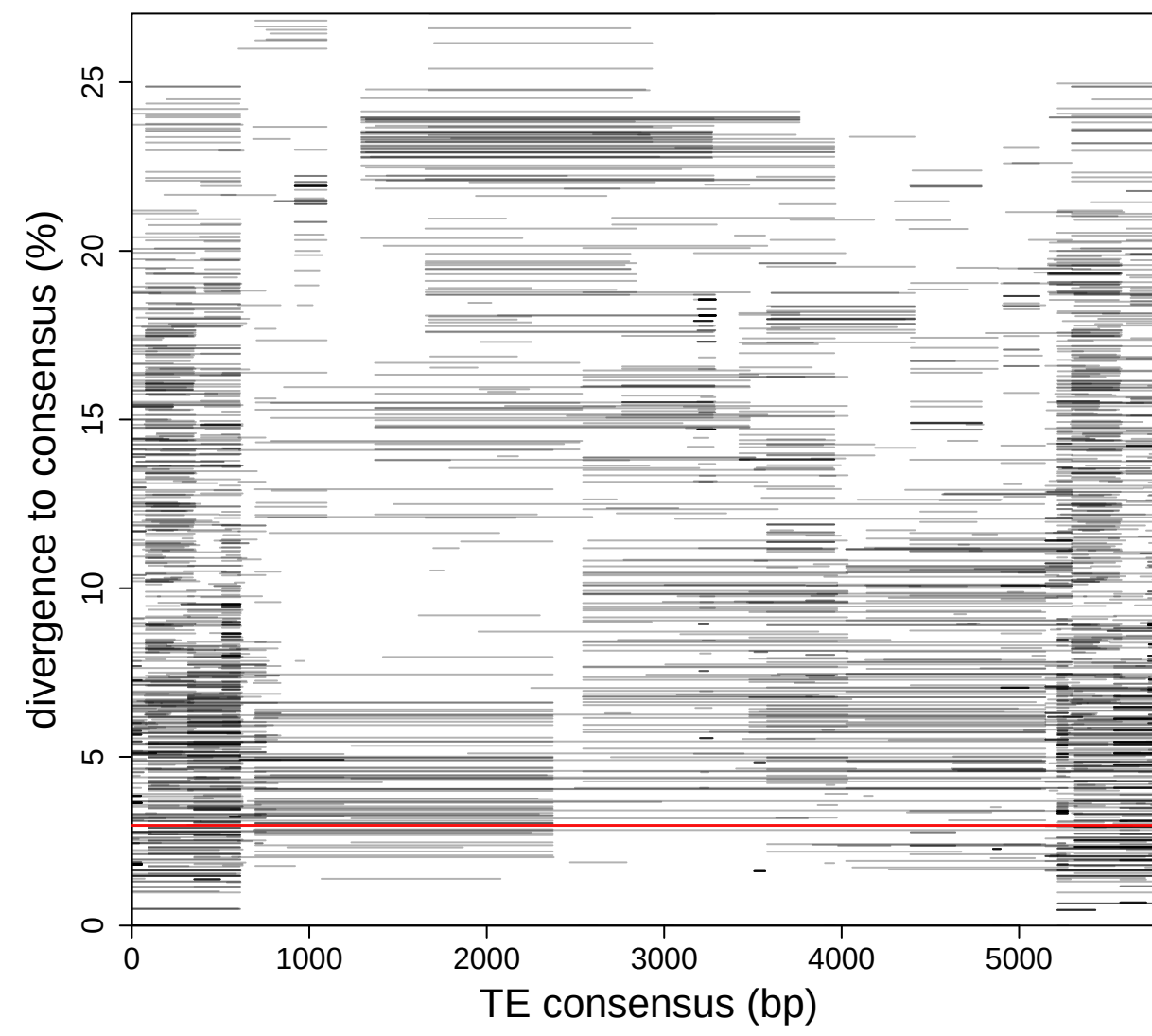
5000

6000



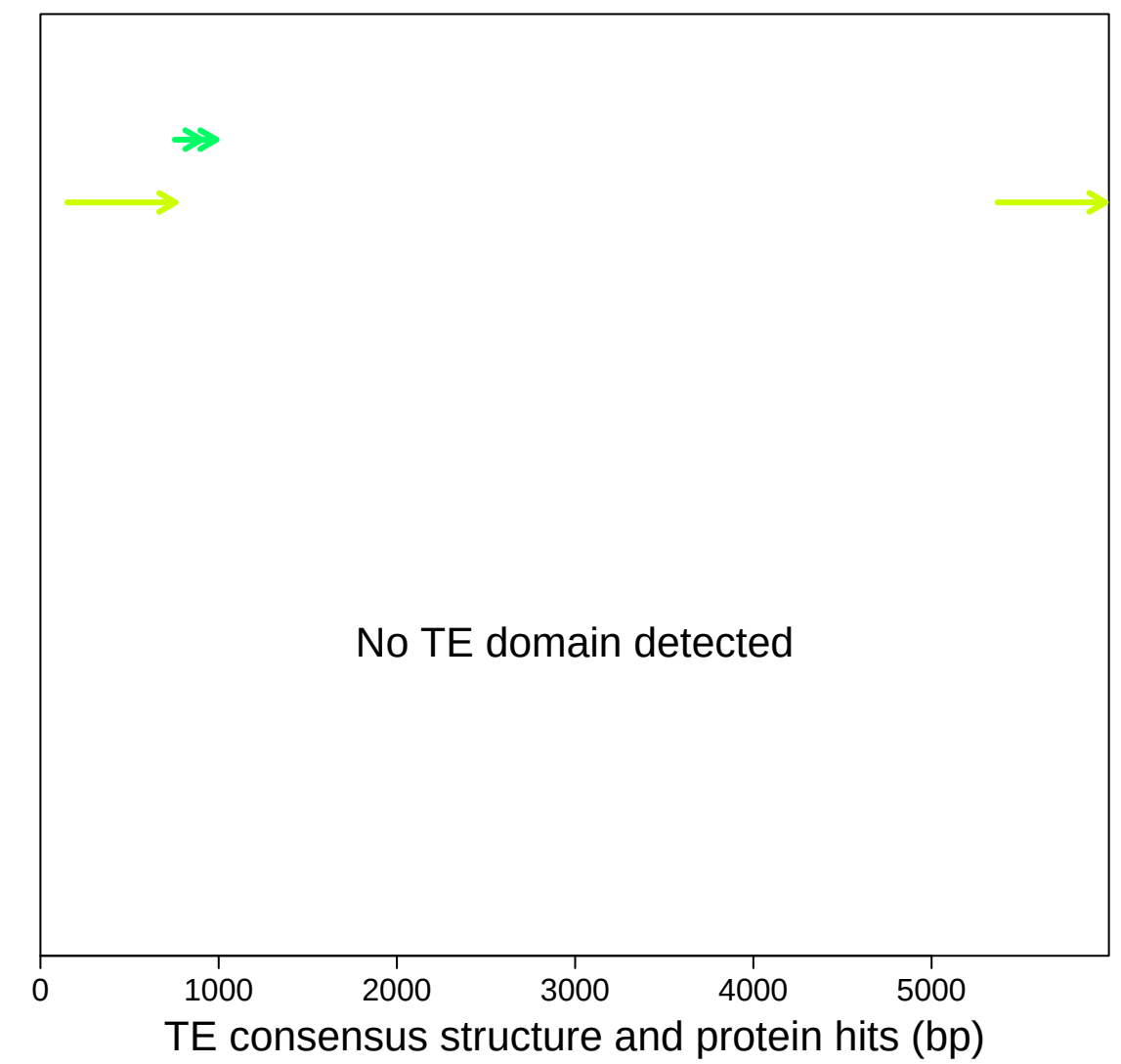
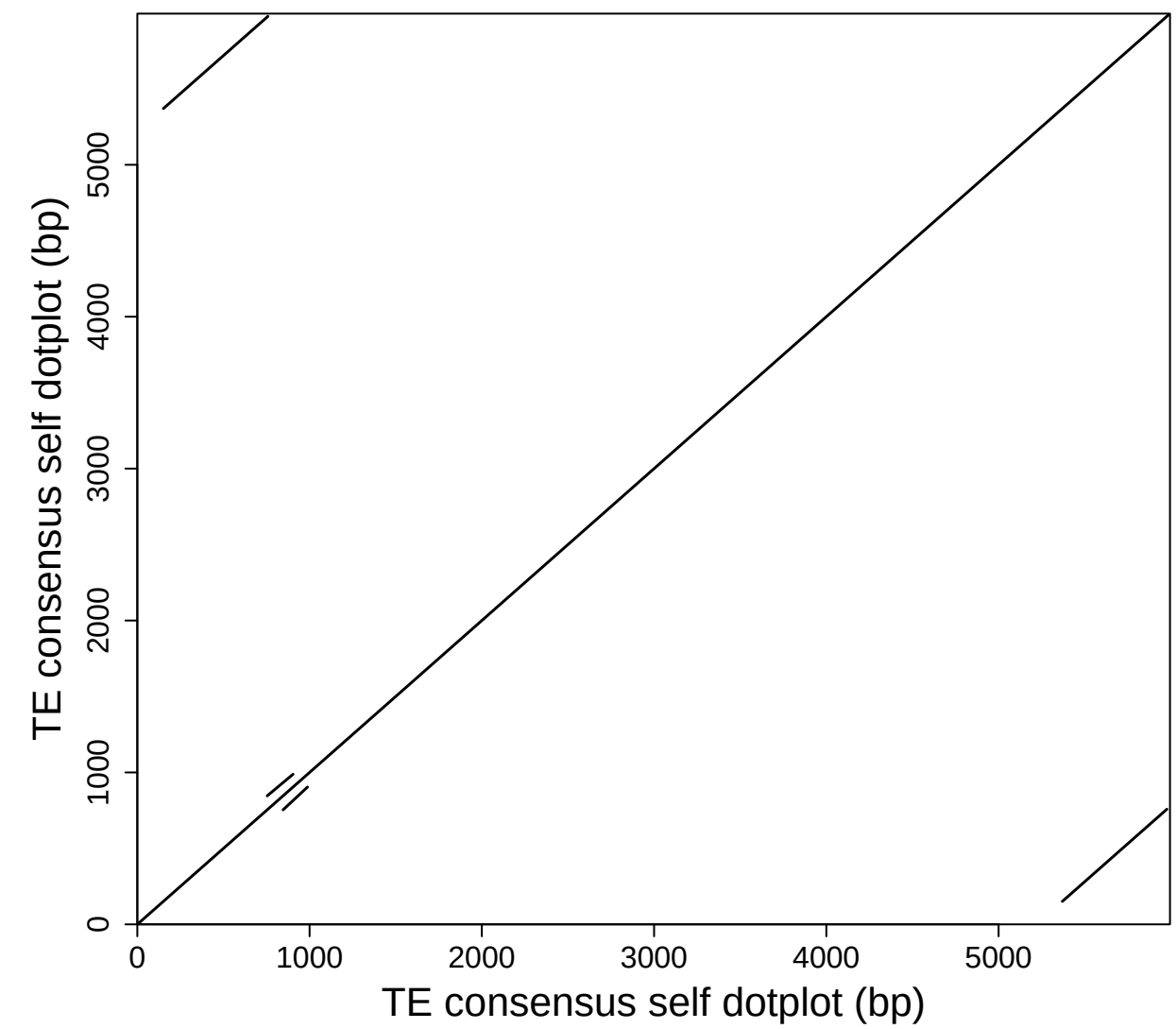
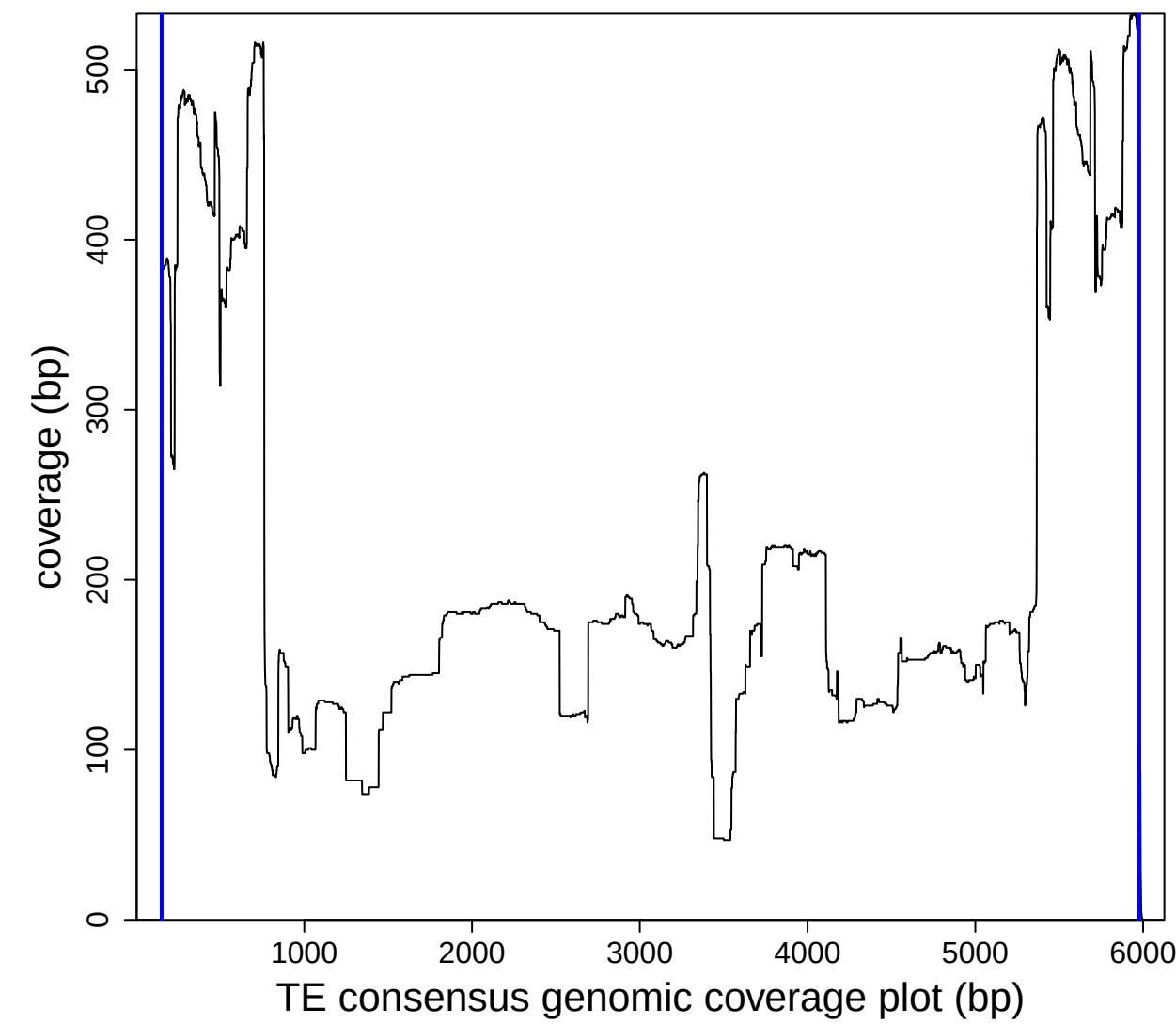
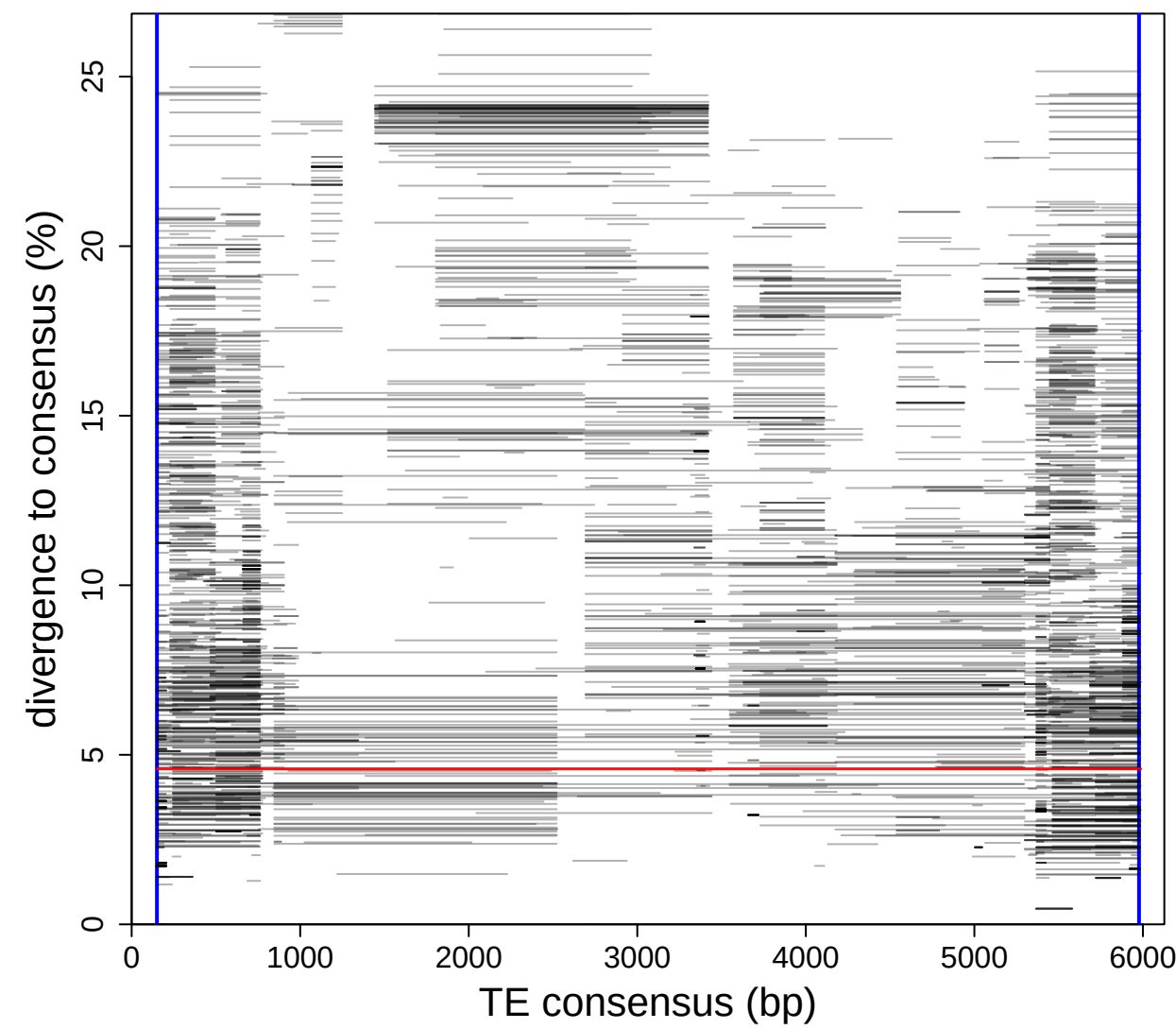
ed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.fa_ce.
size: 5827bp; fragments: 2799; full length: 1 (>=5244.3bp)

After TETrimmer 5827 bp



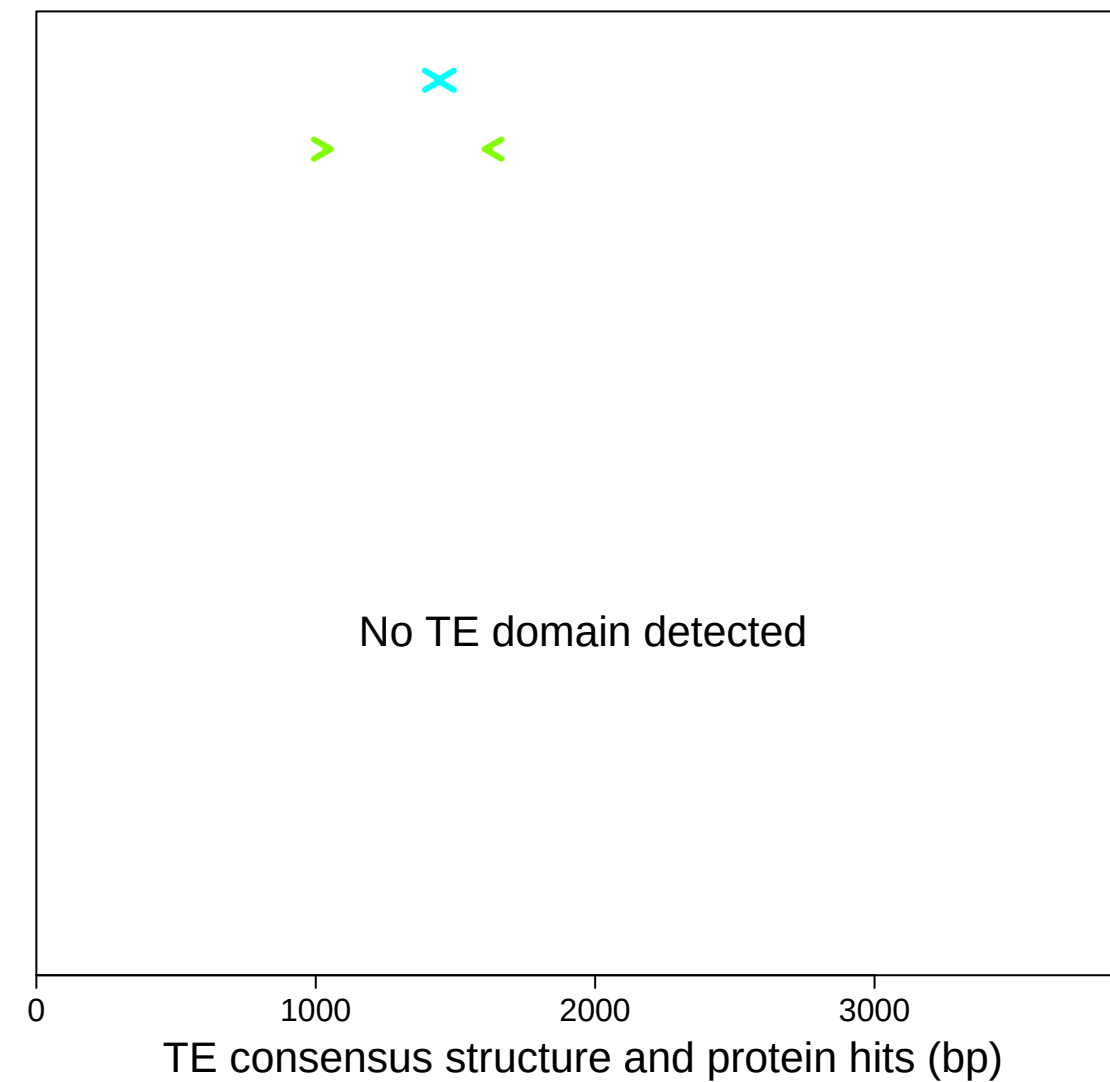
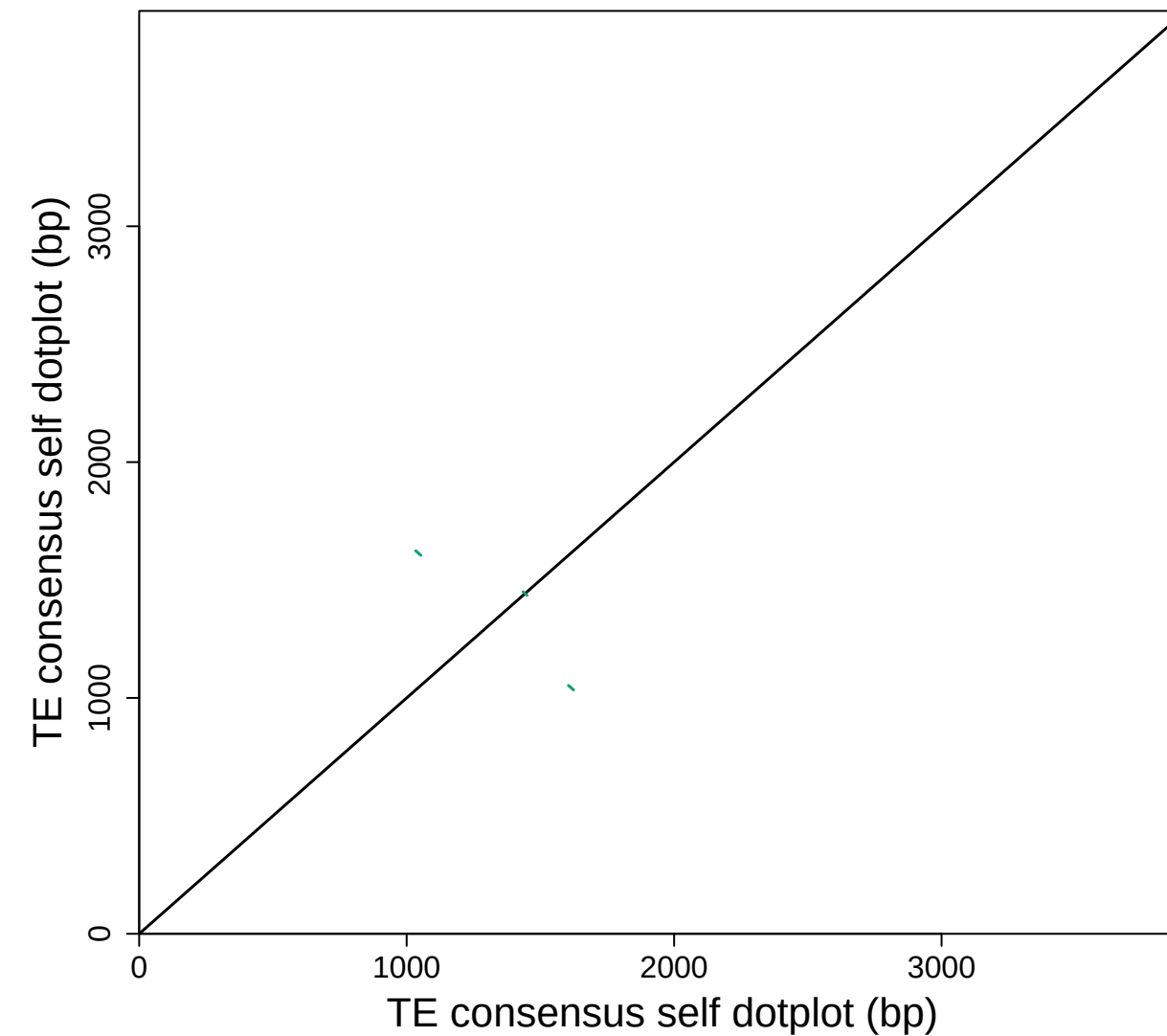
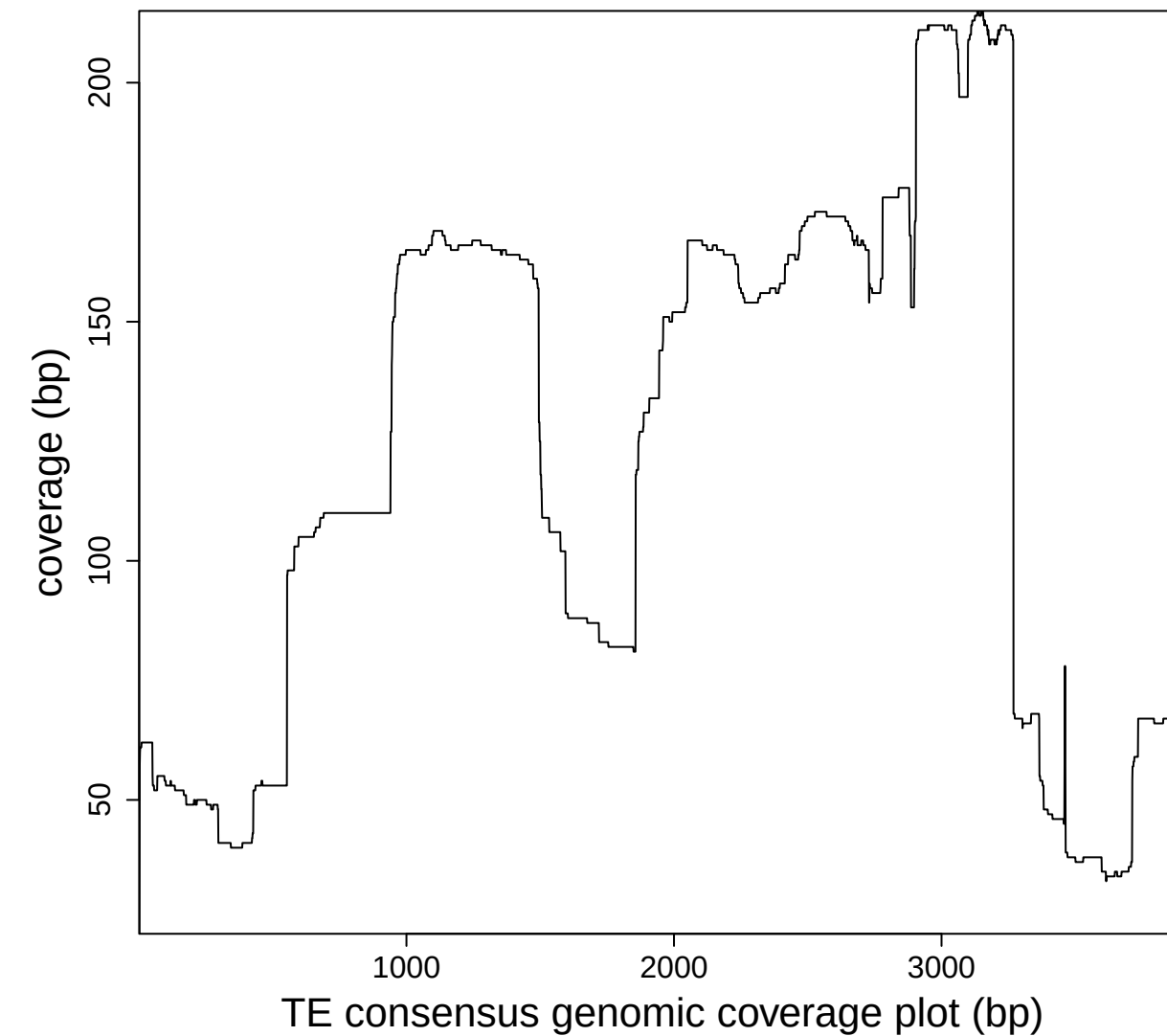
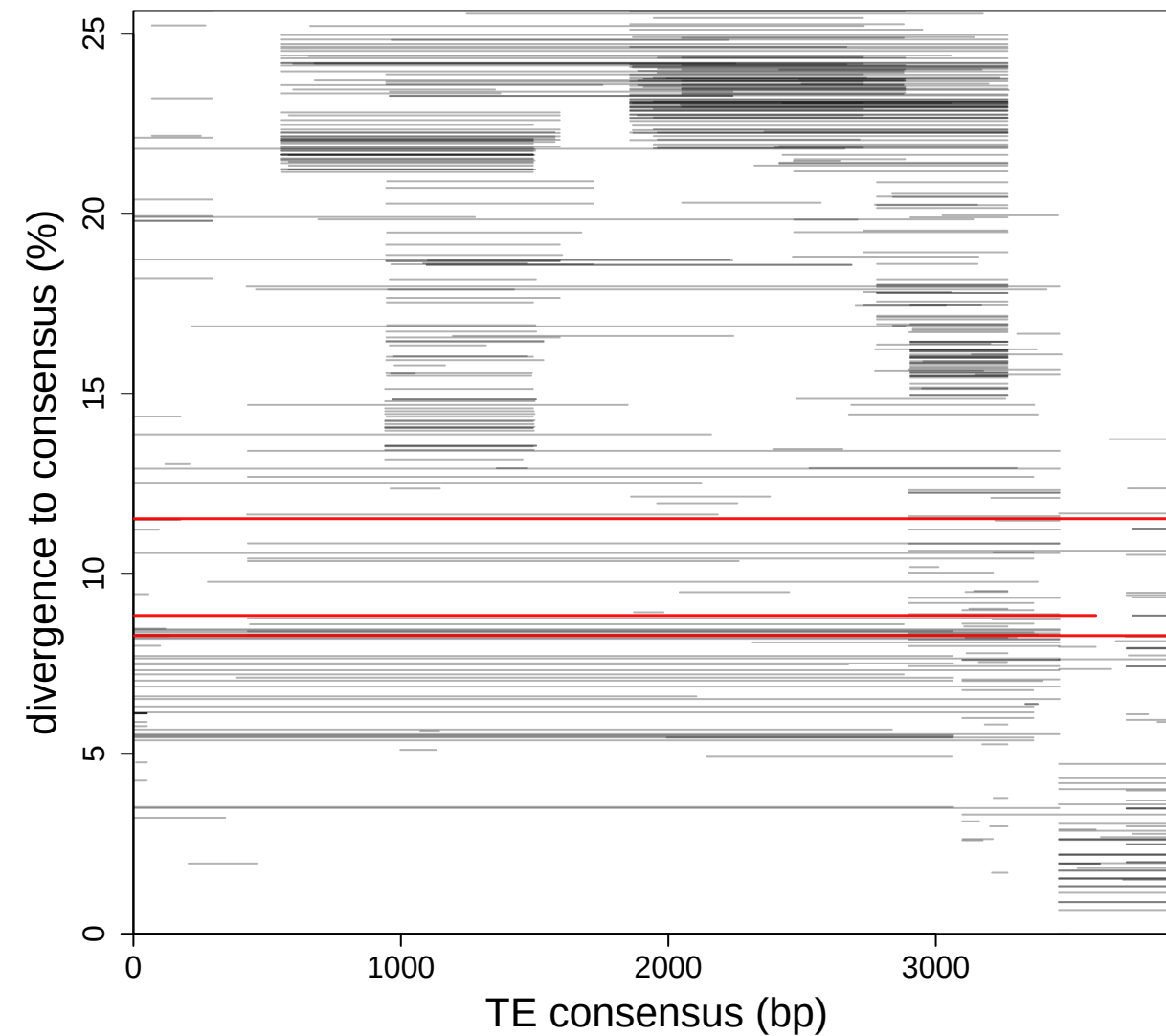
ly_431.fasta.b.bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa
size: 6127bp; fragments: 2878; full length: 1 (>=5514.3bp)

After TEtrimmer Extended plot Blue lines are boundaries

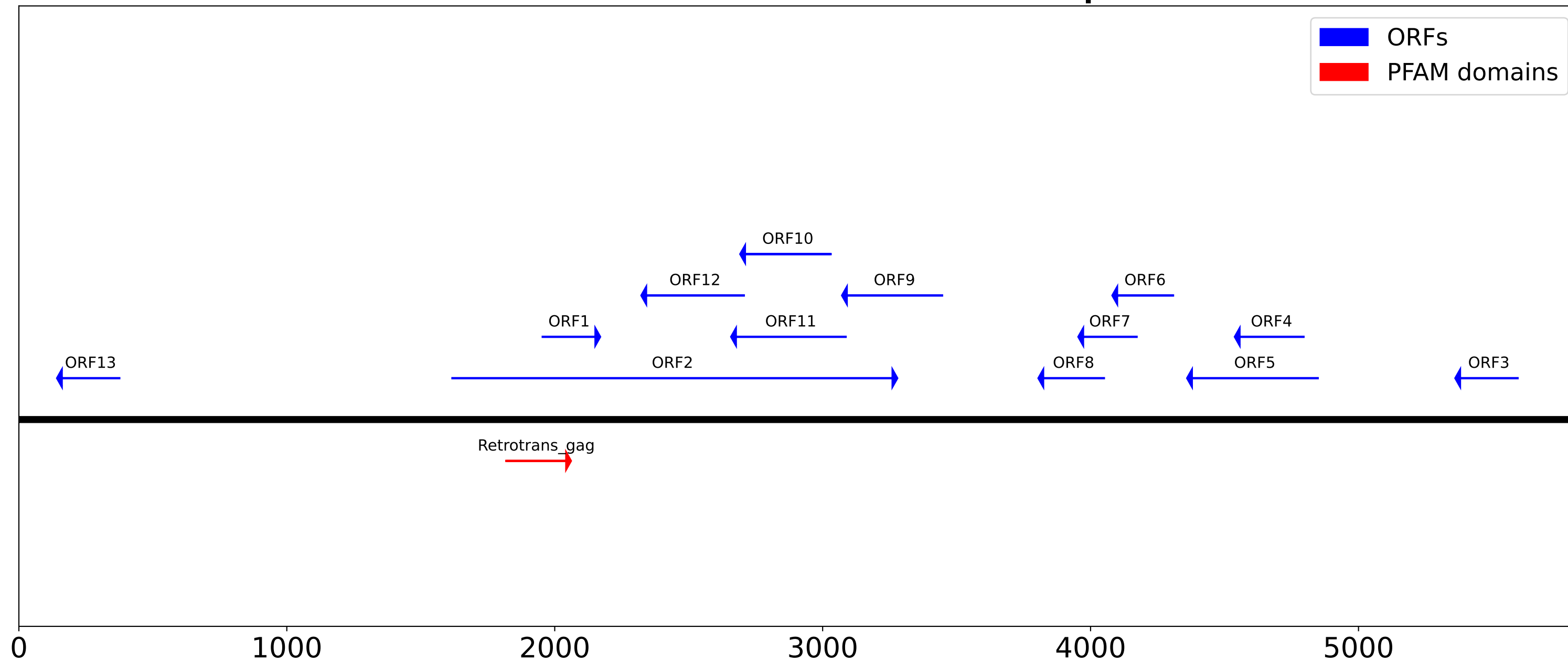


TE: ltr_1_family_431
size: 3913bp; fragments: 506; full length: 3 (>=3521.7bp)

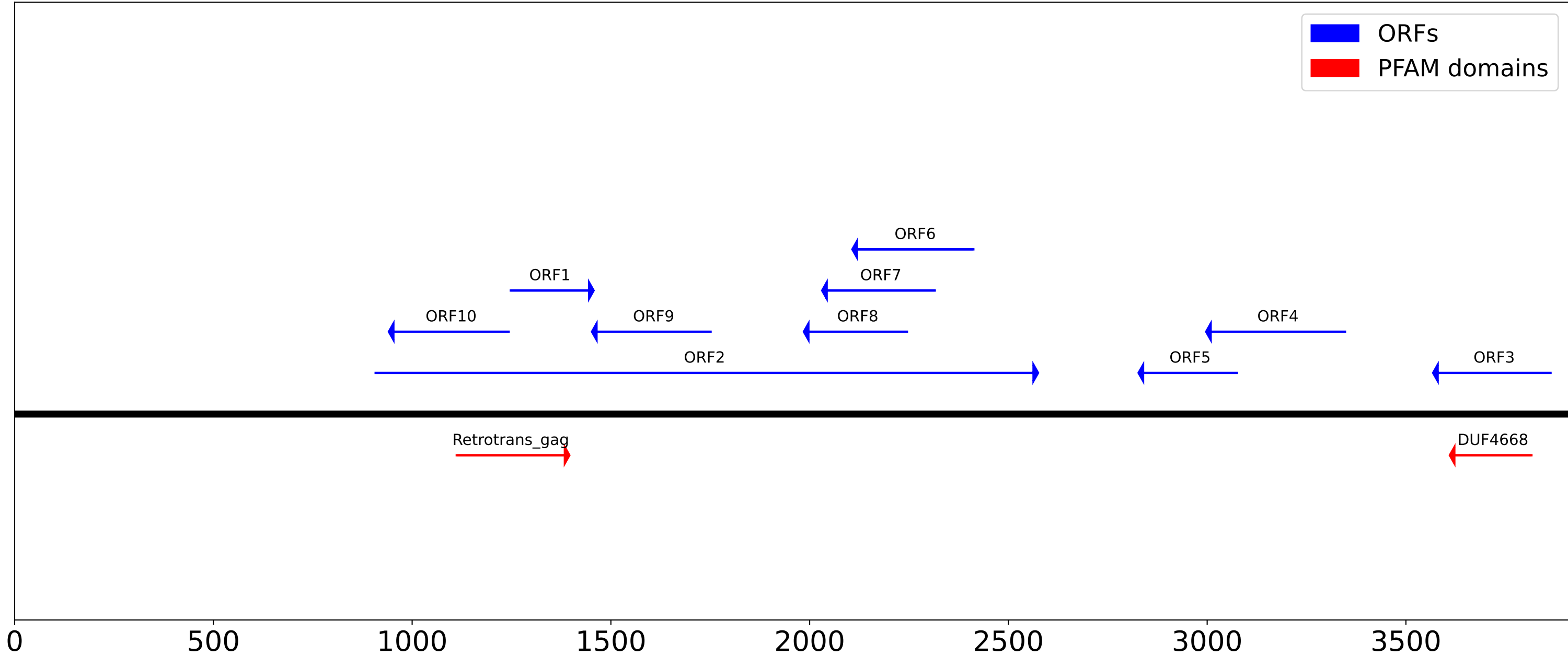
Before TEtrimmer 3913 bp



After TEtrimmer ORF and PFAM domain plot



Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

