

Start crop Point



End crop Point



MSA length = 6356



Start crop Point

End crop Point

0

10

0

1000

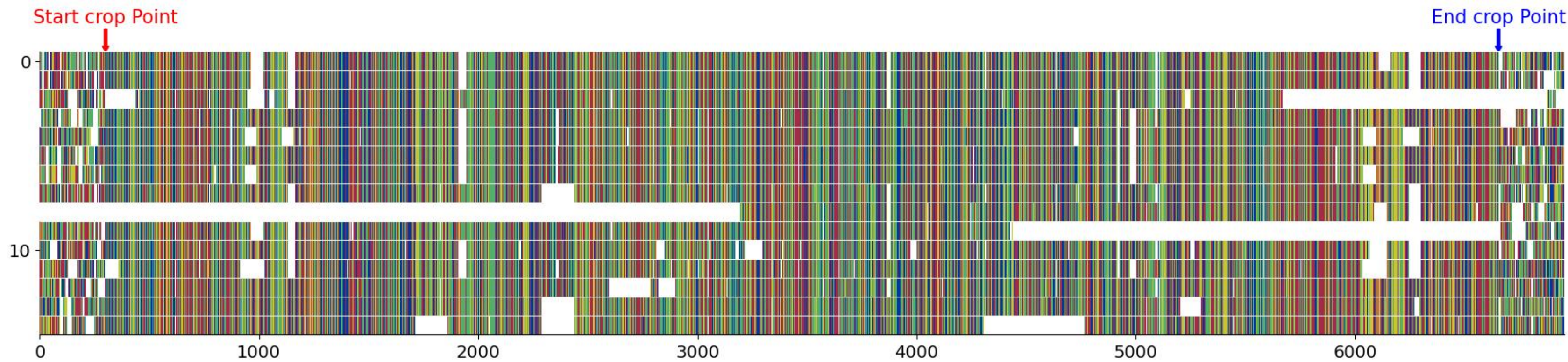
2000

3000

4000

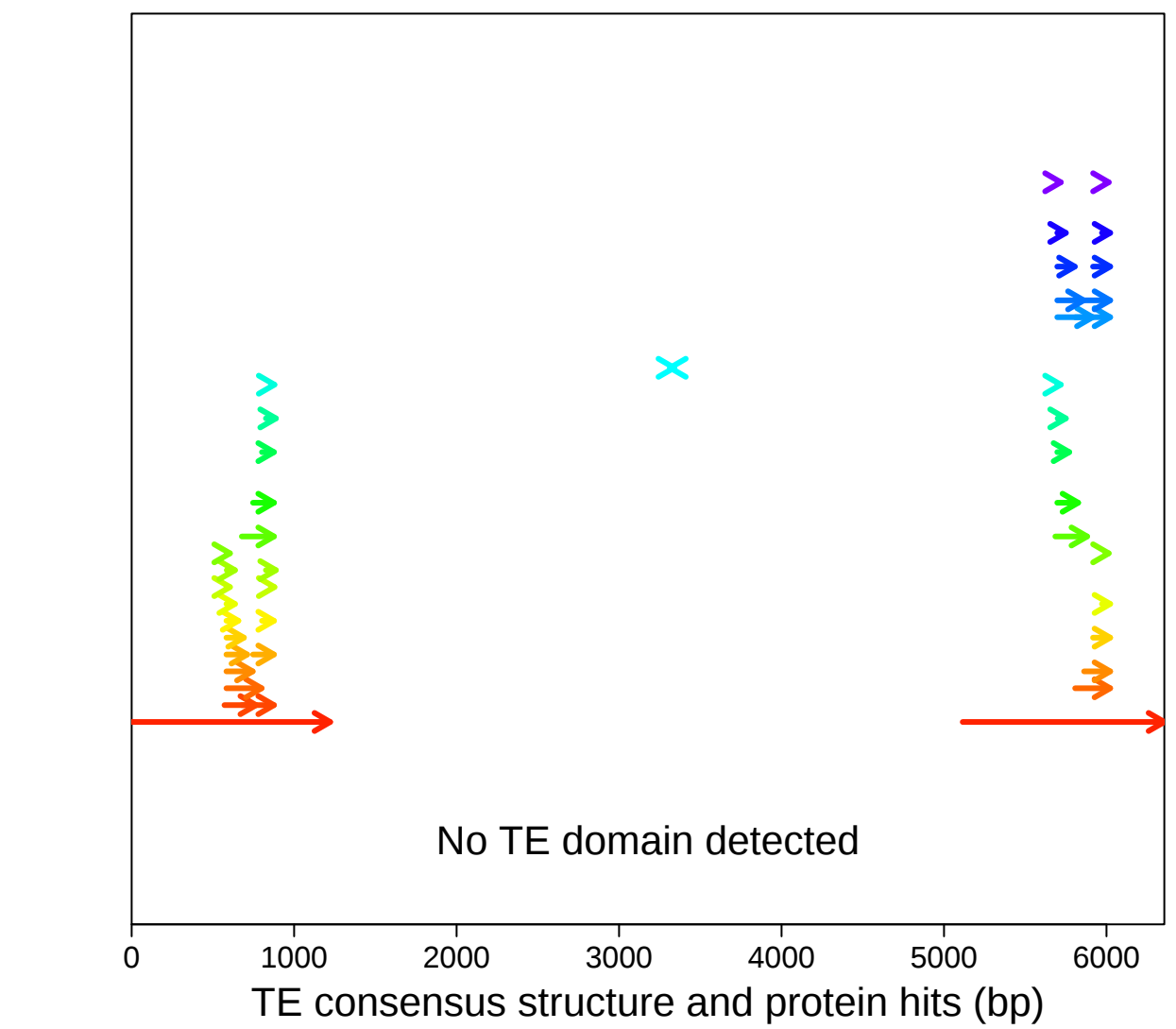
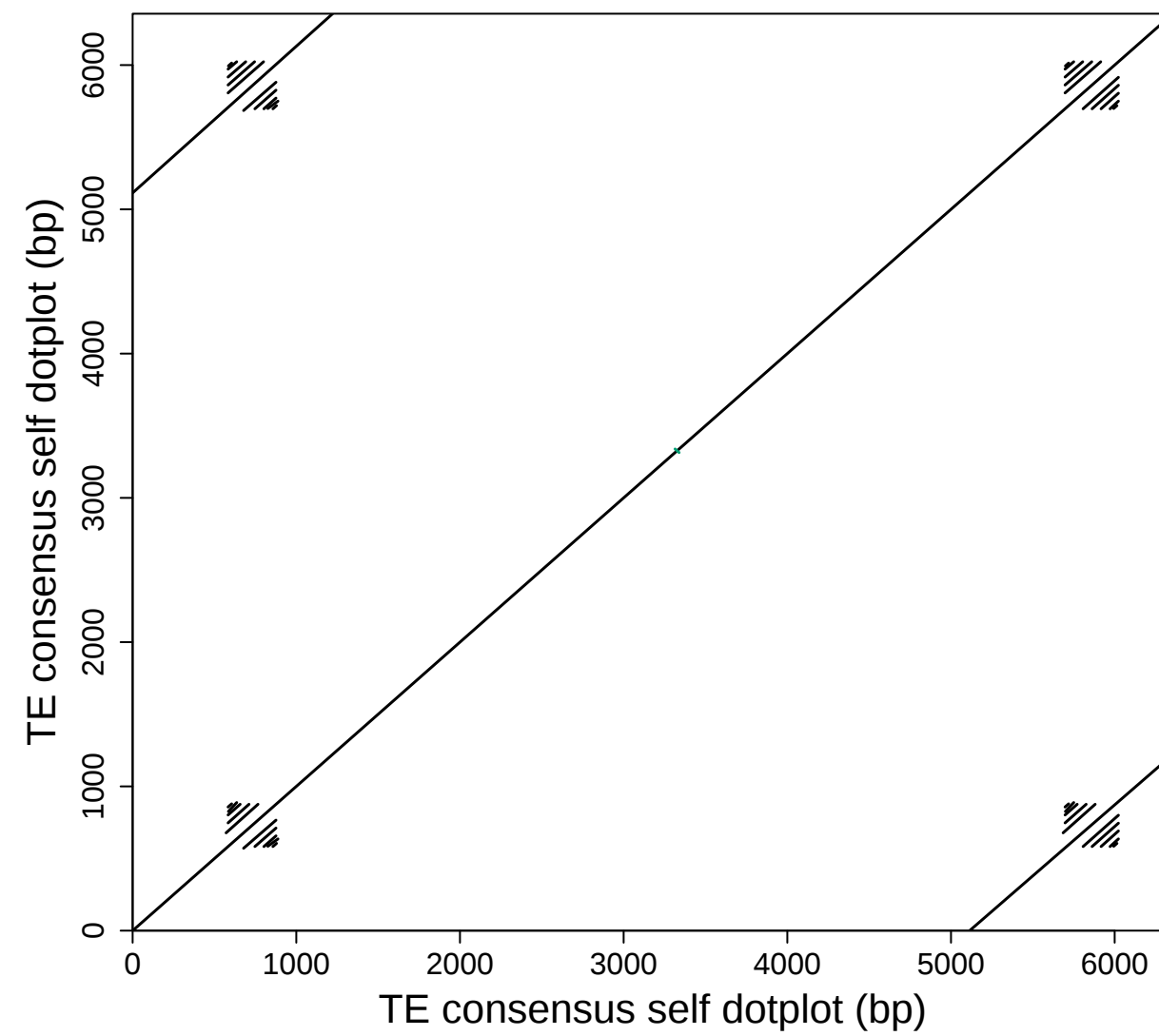
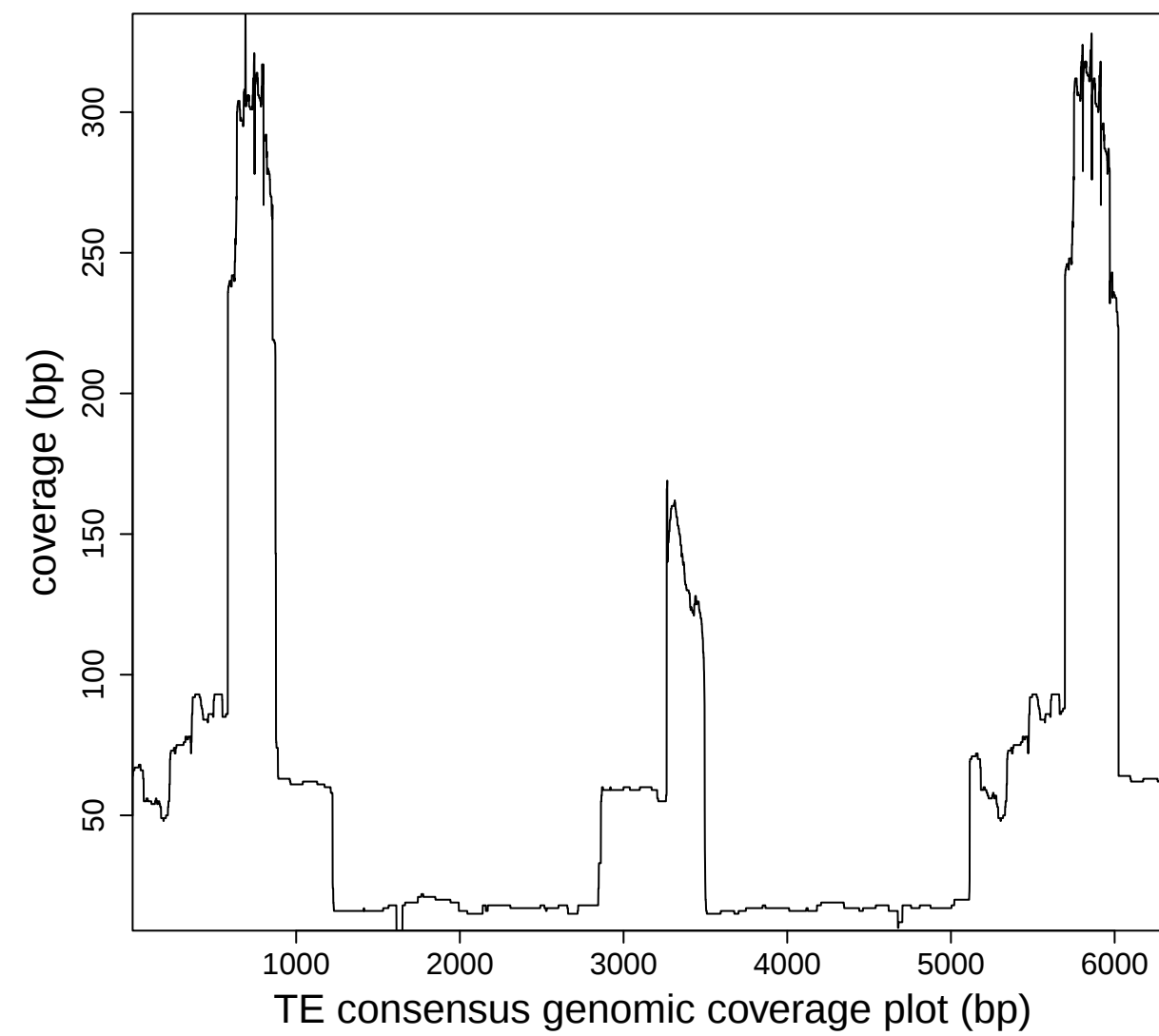
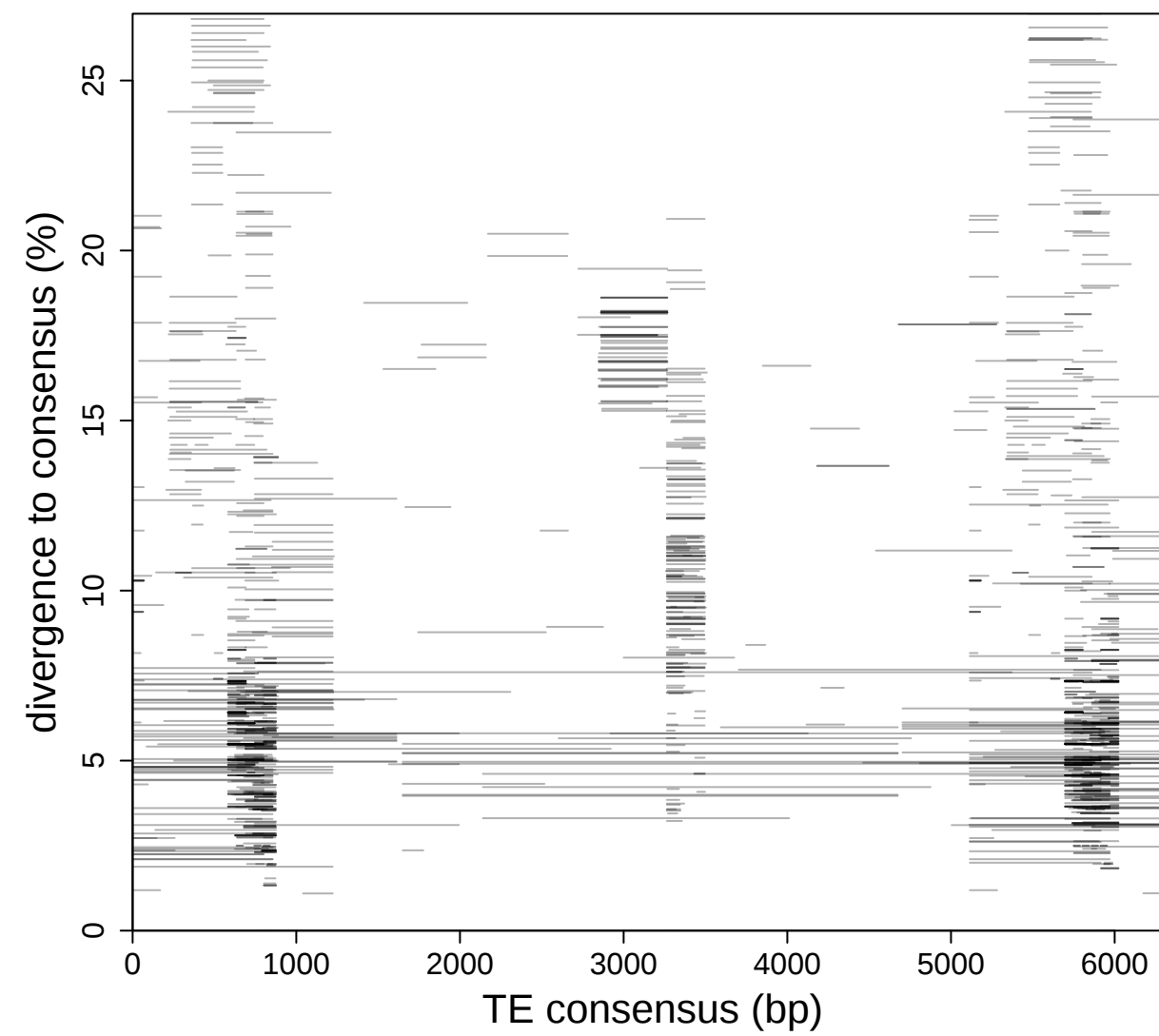
5000

6000



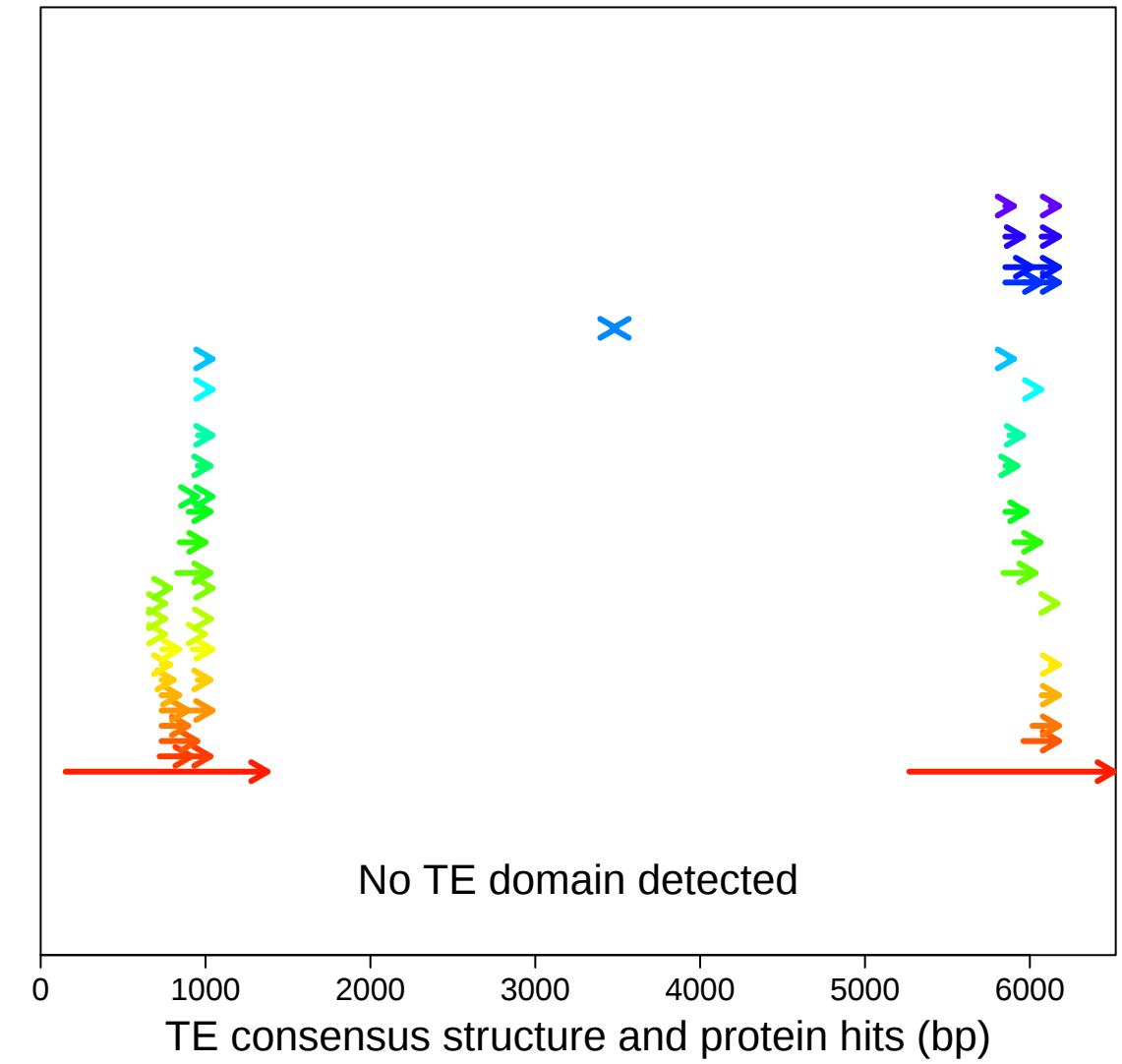
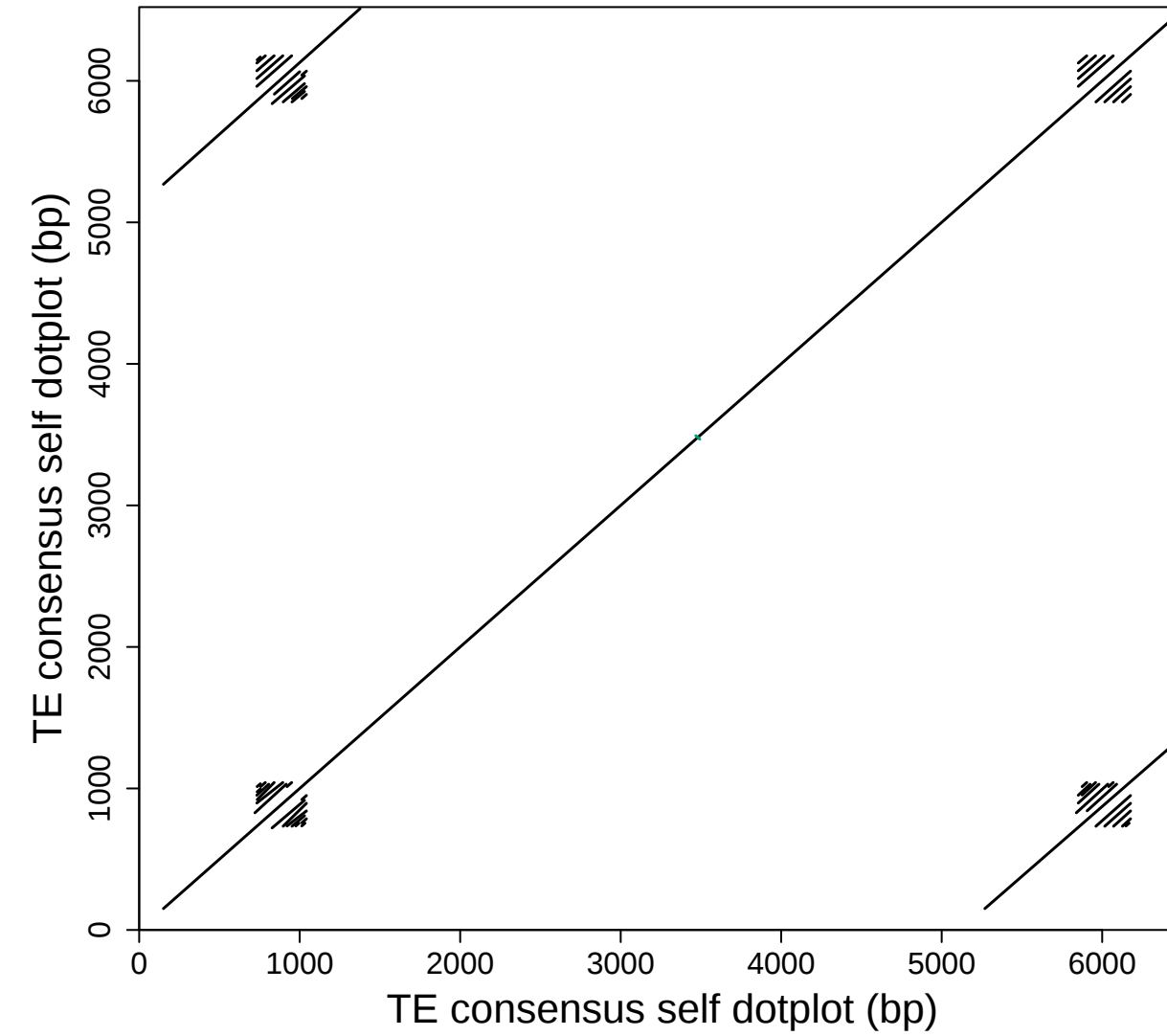
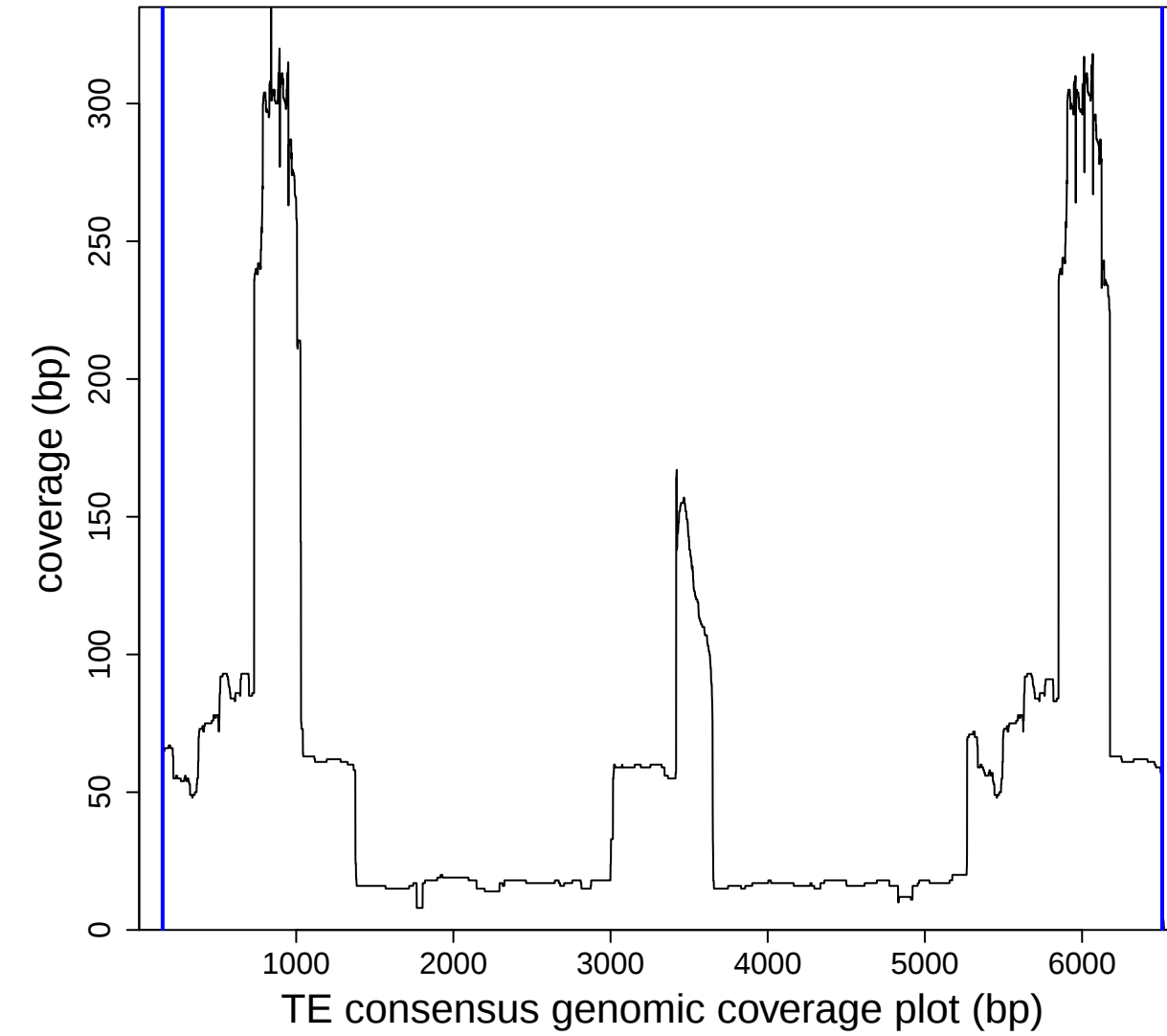
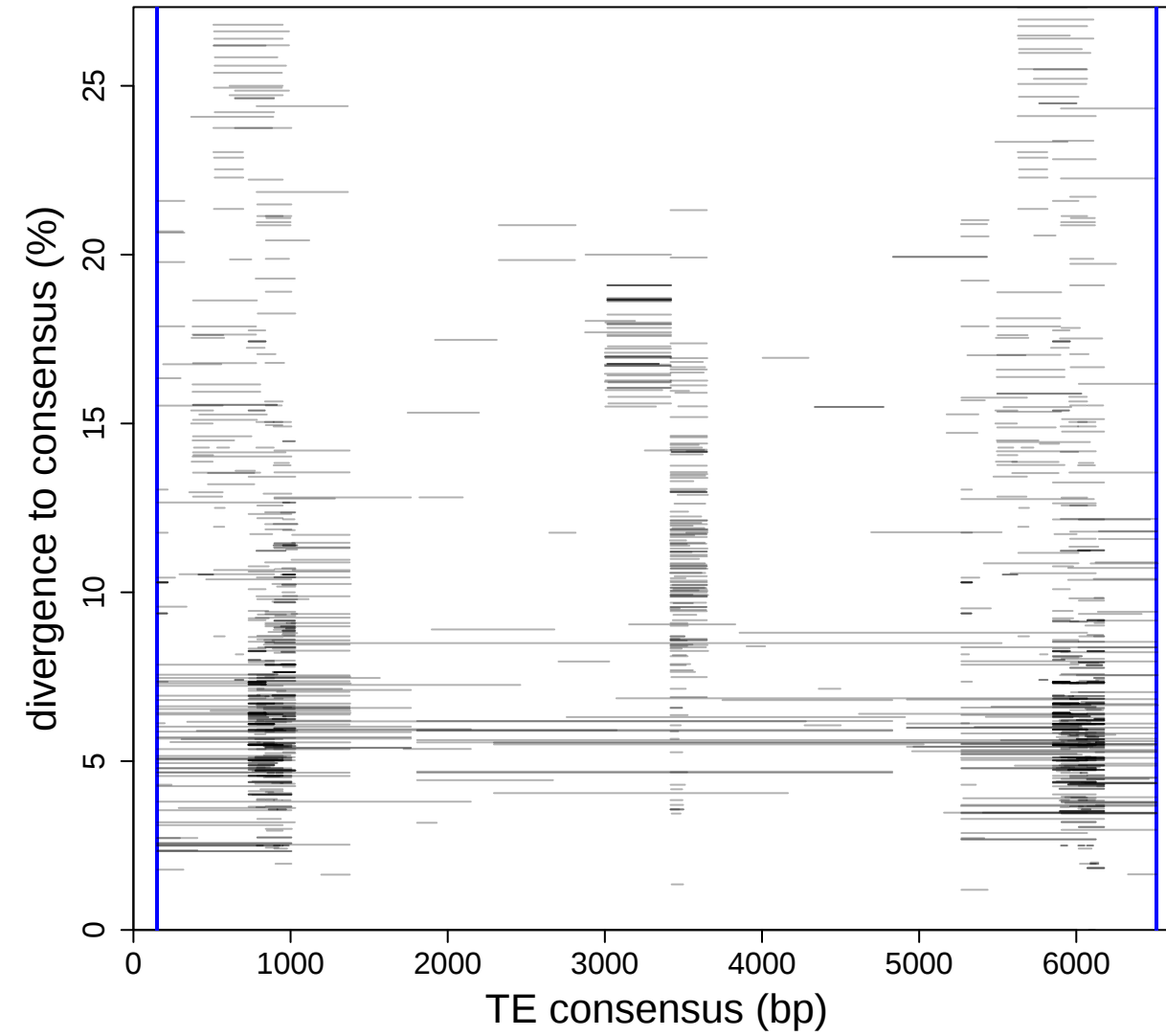
ed_uf.bed_g_3.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.fa_ce.
size: 6356bp; fragments: 1425; full length: 0 (>=5720.4bp)

After TEtrimmer 6356 bp



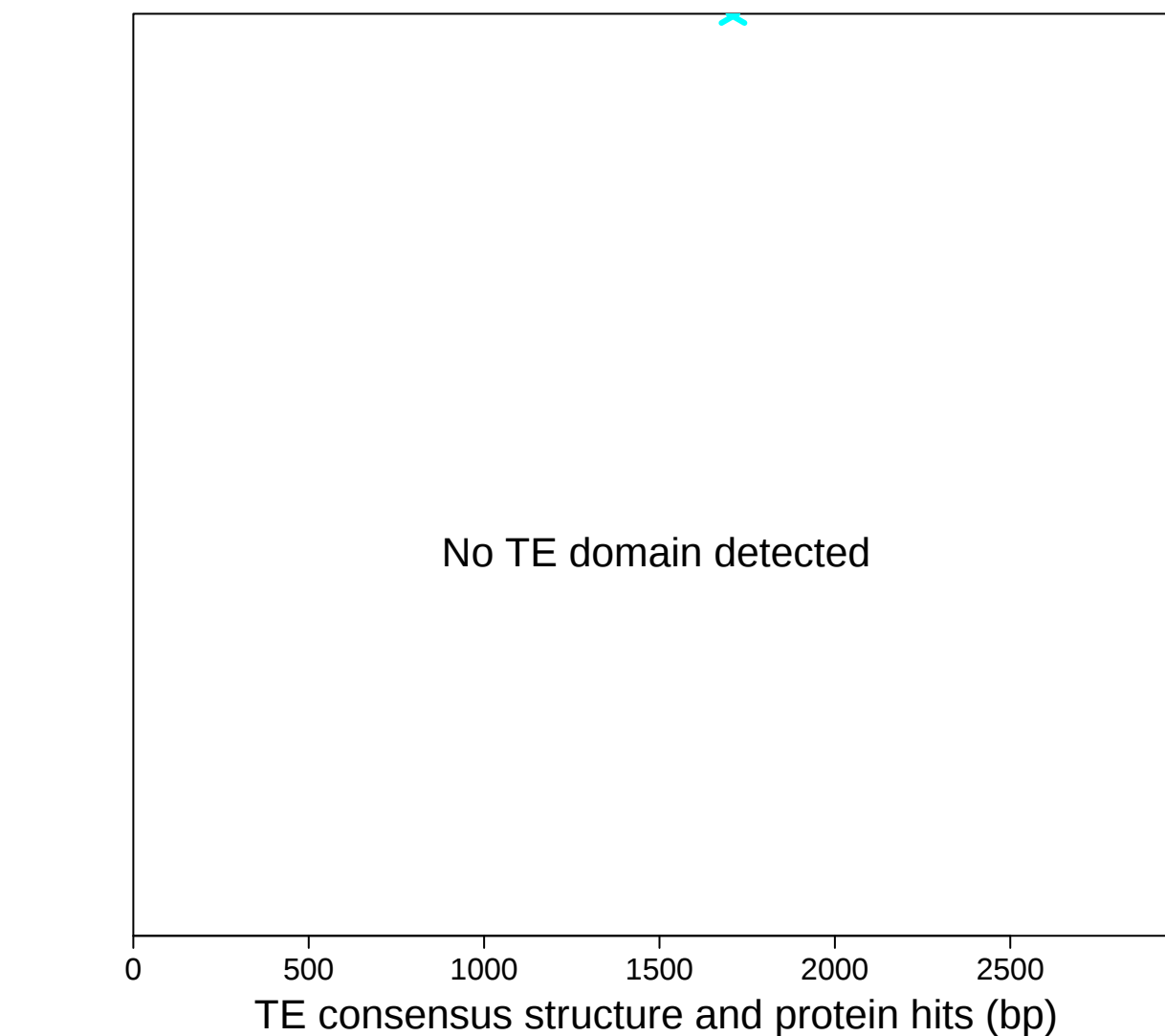
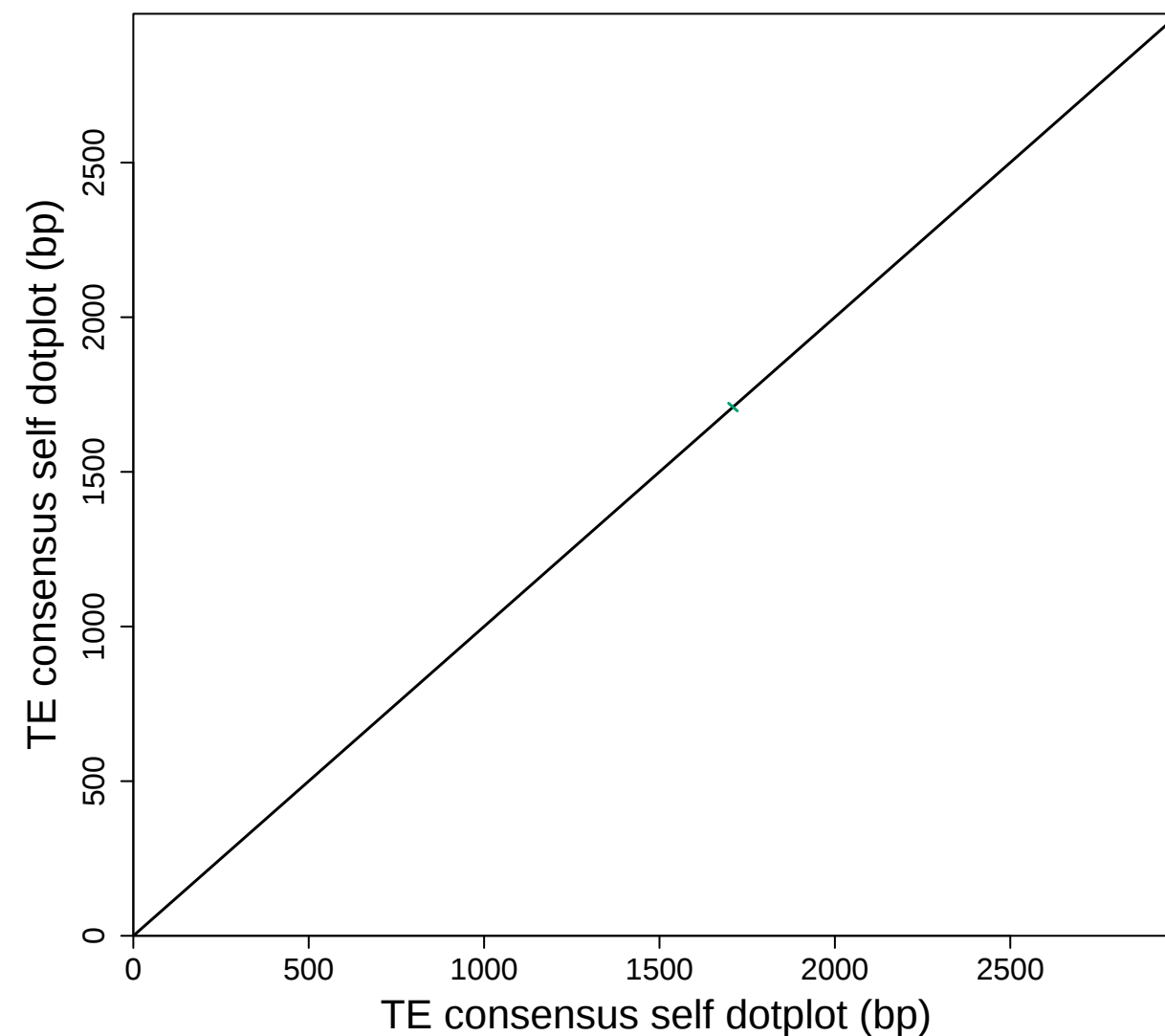
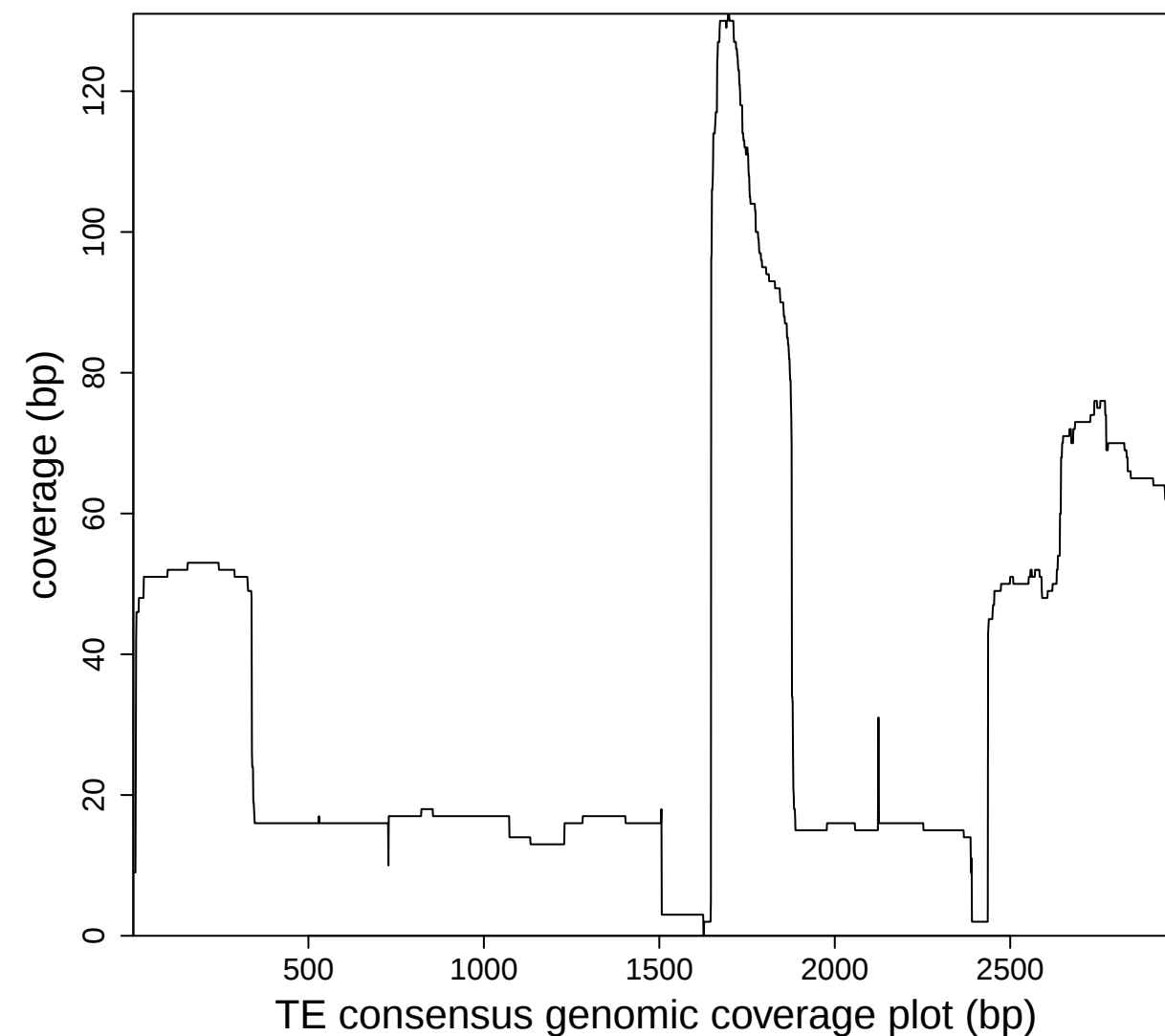
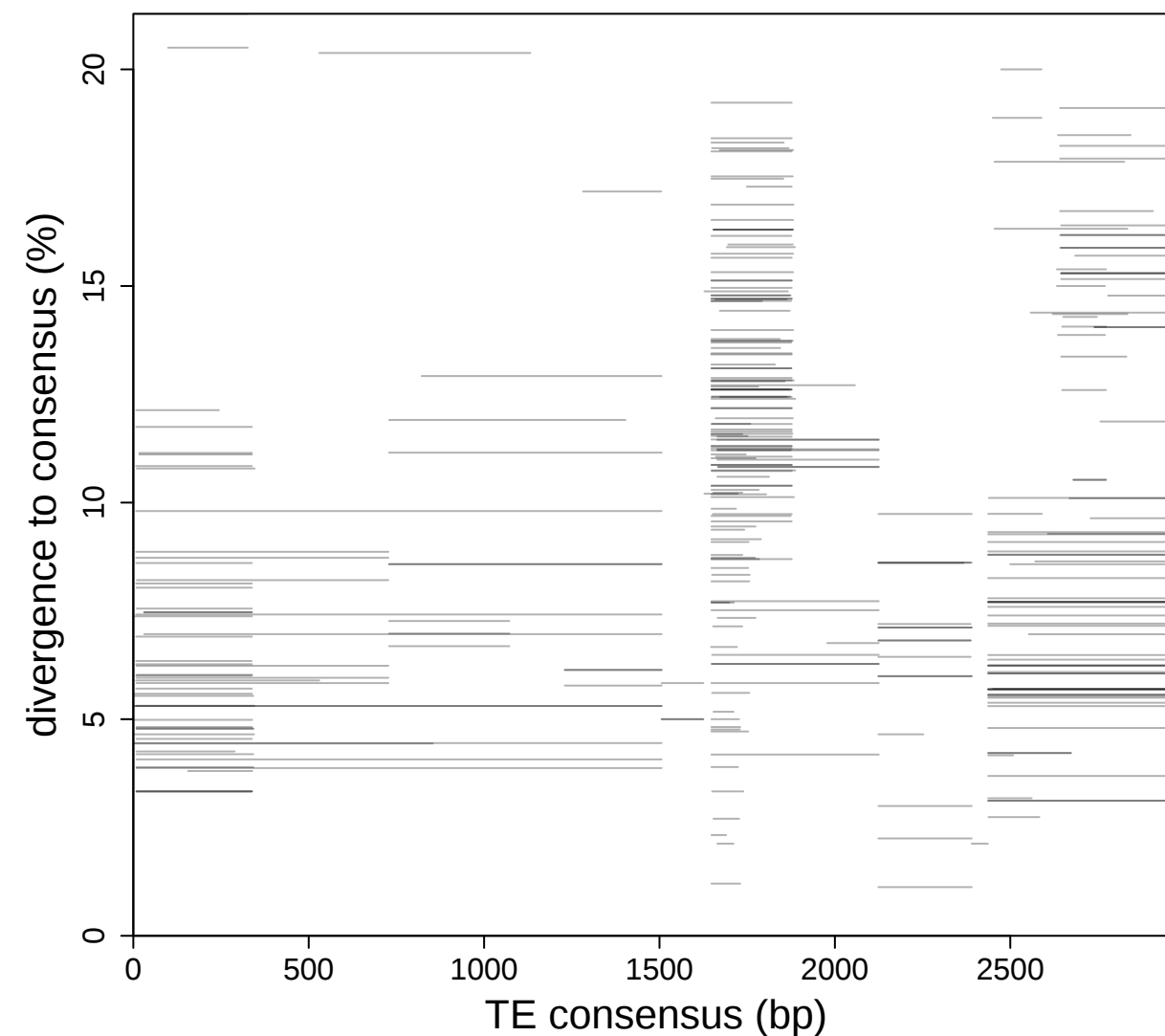
ly_653.fasta.b.bed_uf.bed_g_3.bed_fm_1.bed_0_0_bcln.fa_aln.fa
size: 6660bp; fragments: 1381; full length: 0 (>=5994bp)

After TEtrimmer Extended plot Blue lines are boundaries

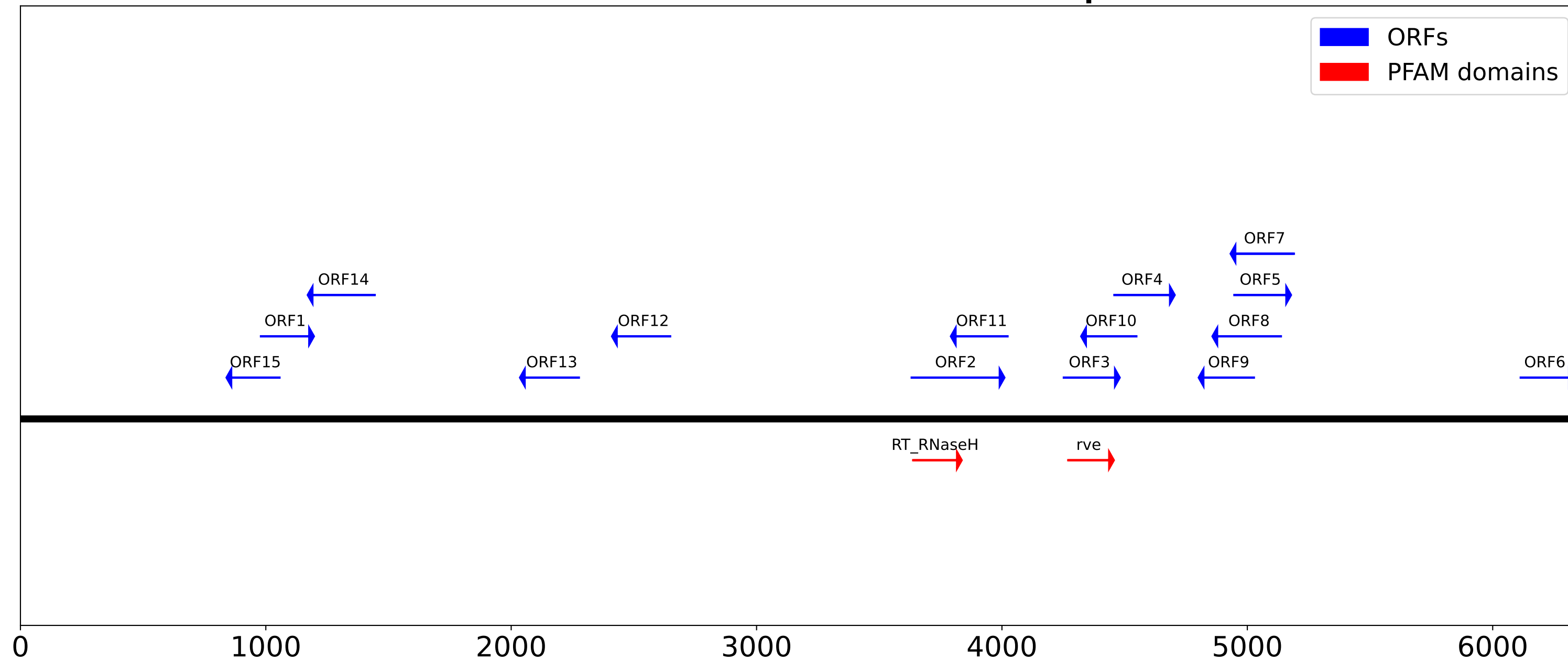


TE: ltr_1_family_653
size: 2981bp; fragments: 307; full length: 0 (≥ 2682.9 bp)

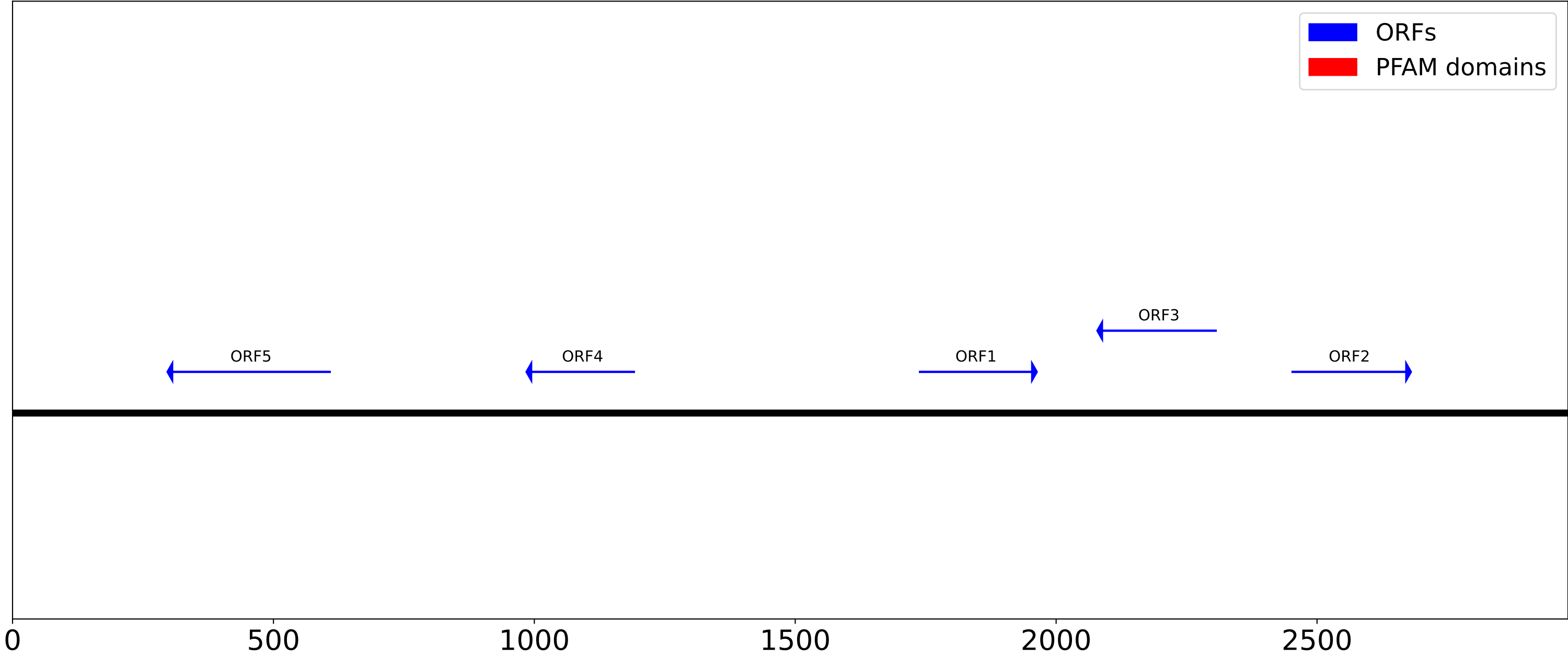
Before TEtrimmer 2981 bp



After TEtrimmer ORF and PFAM domain plot



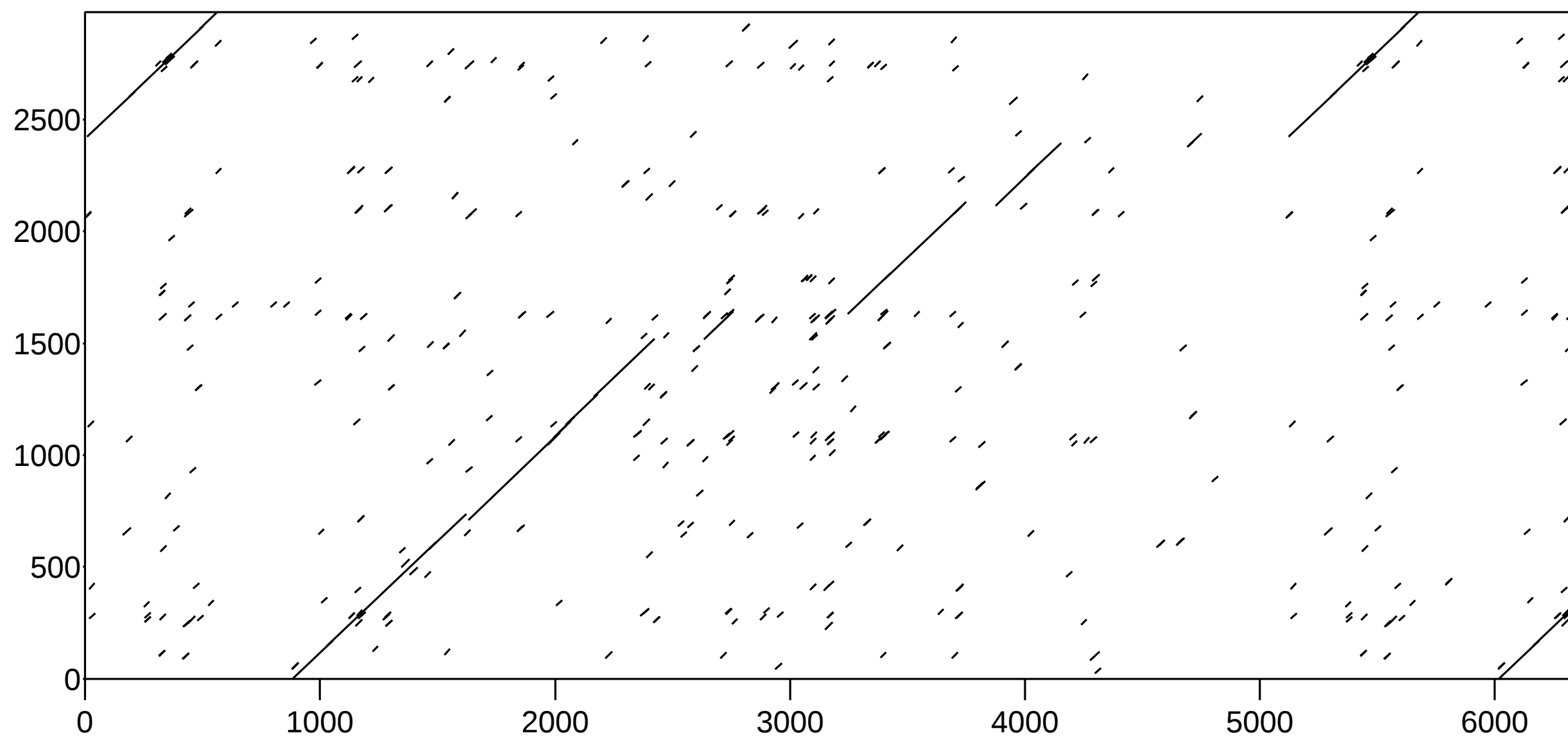
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)