

Start crop Point



End crop Point



MSA length = 5847



Start crop Point



End crop Point



0

10

0

1000

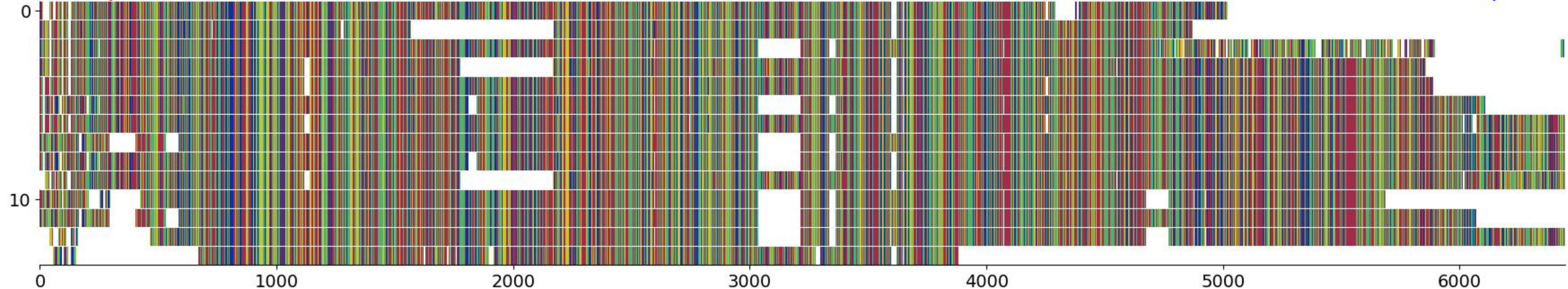
2000

3000

4000

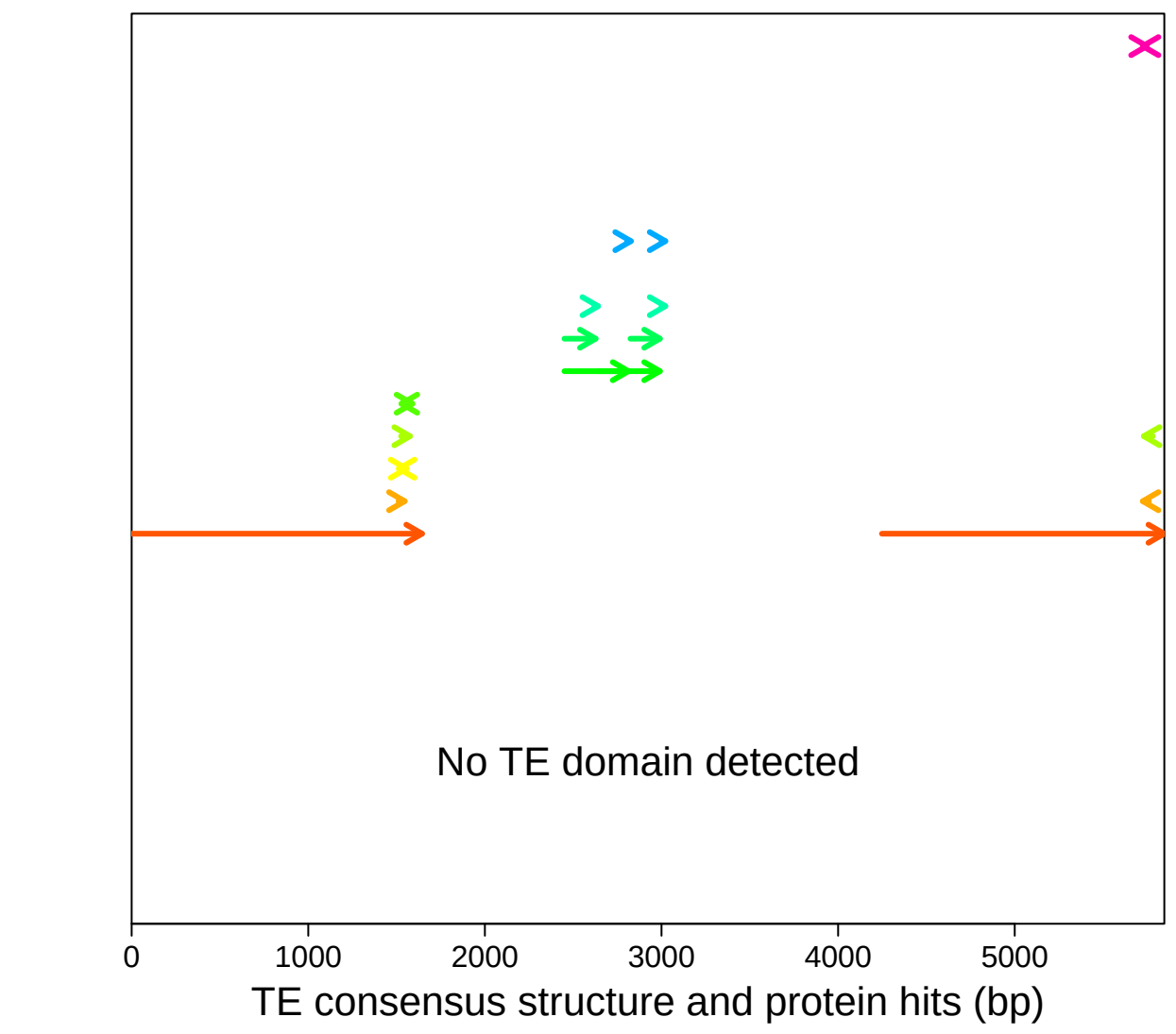
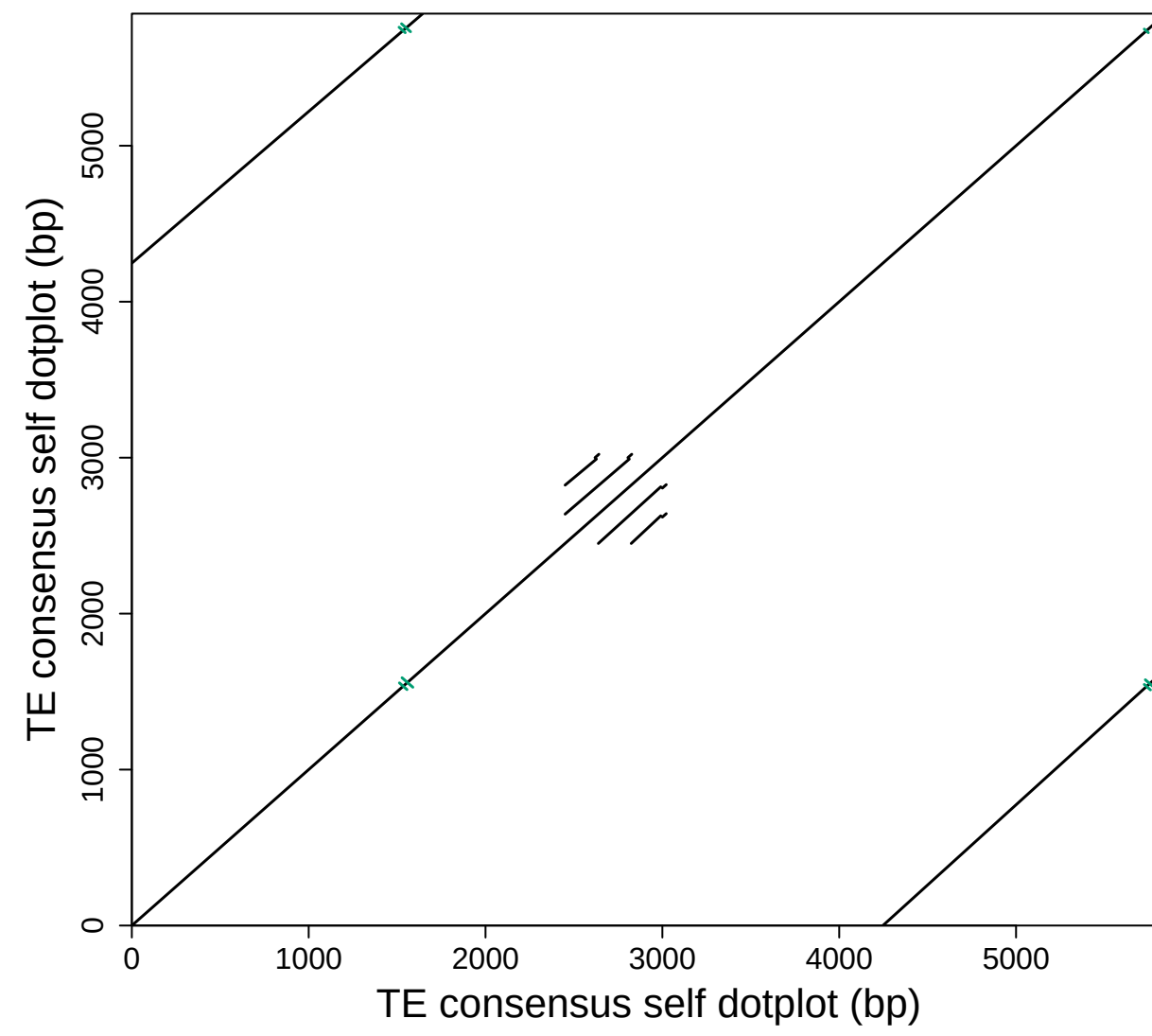
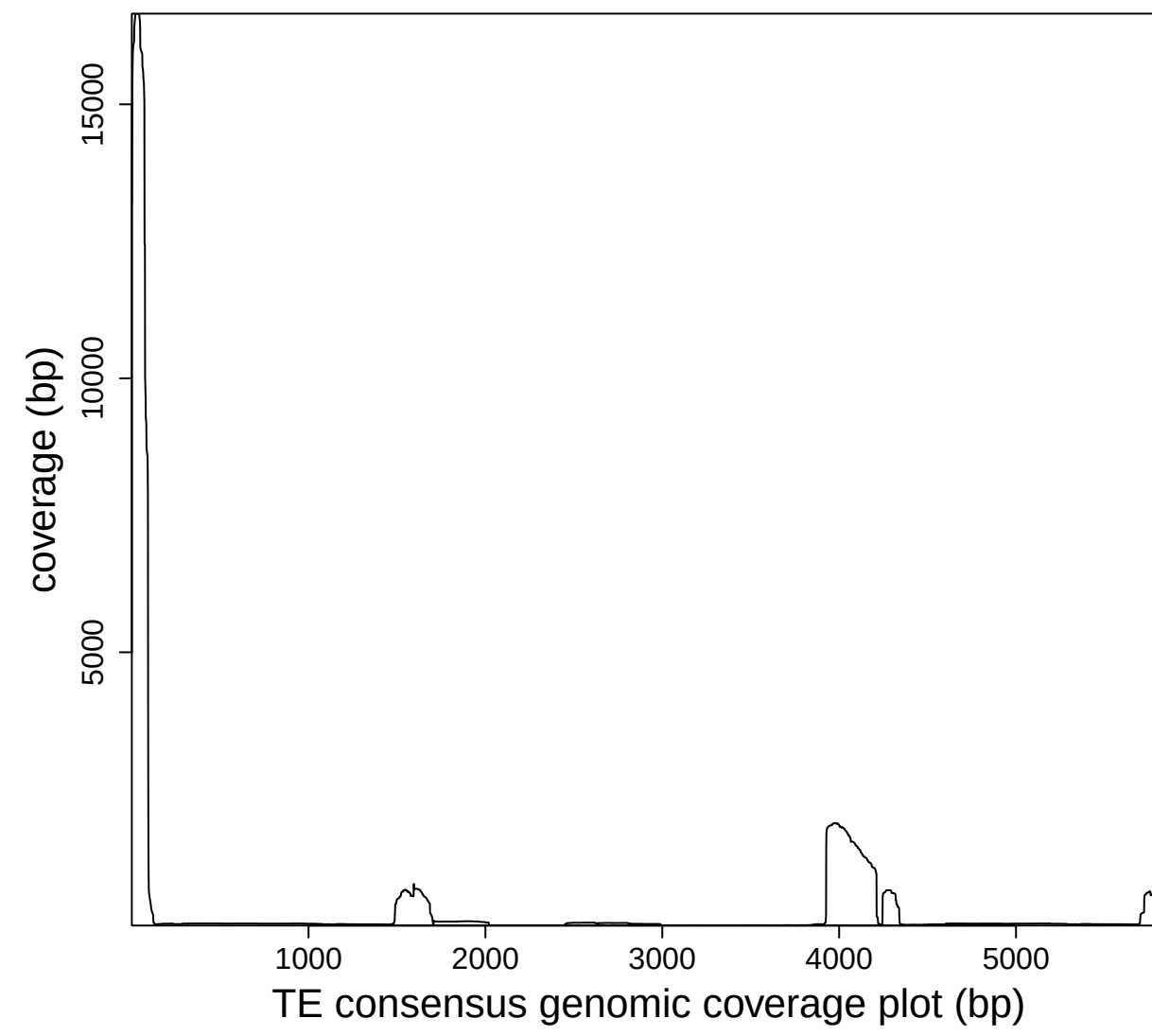
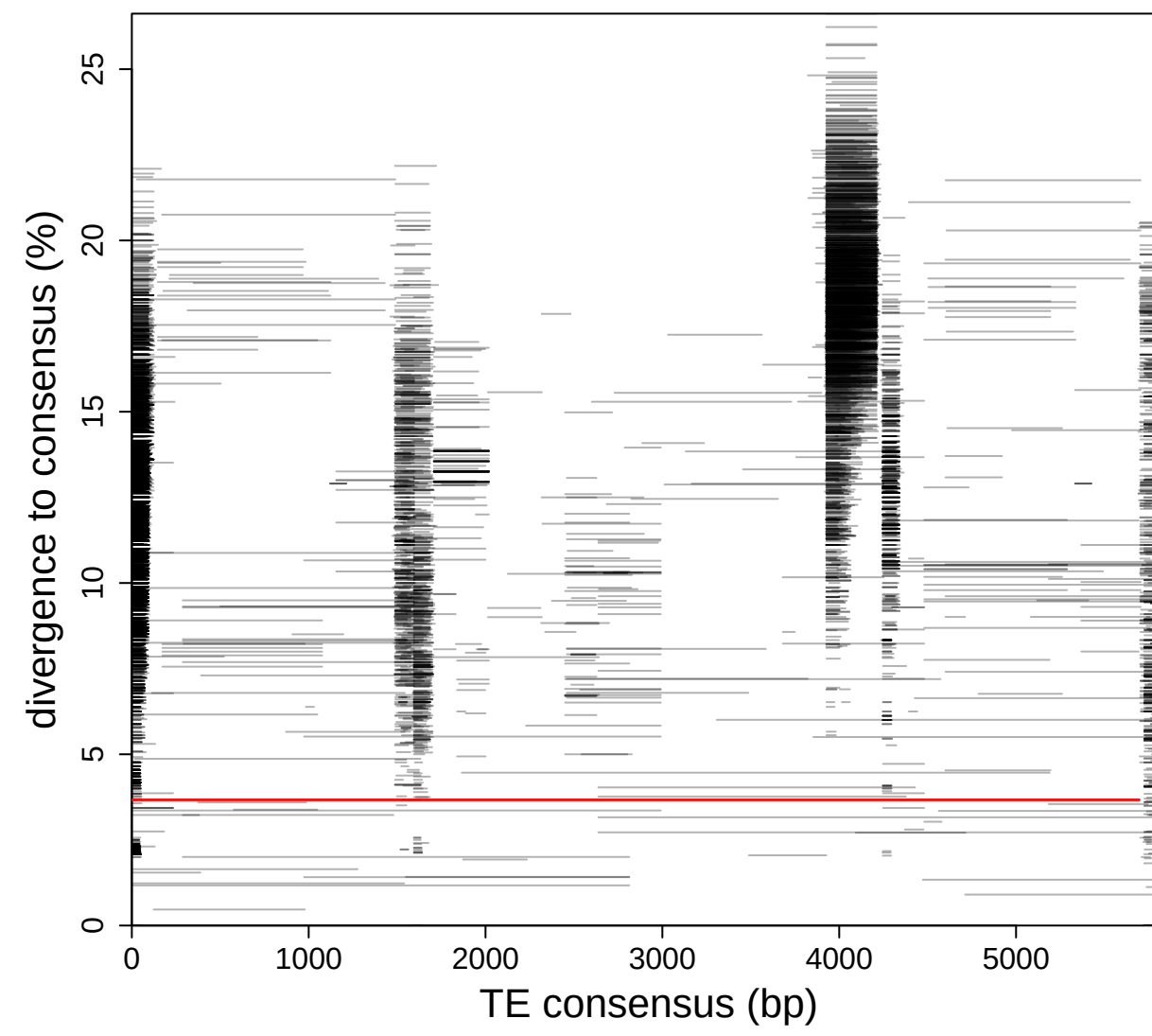
5000

6000



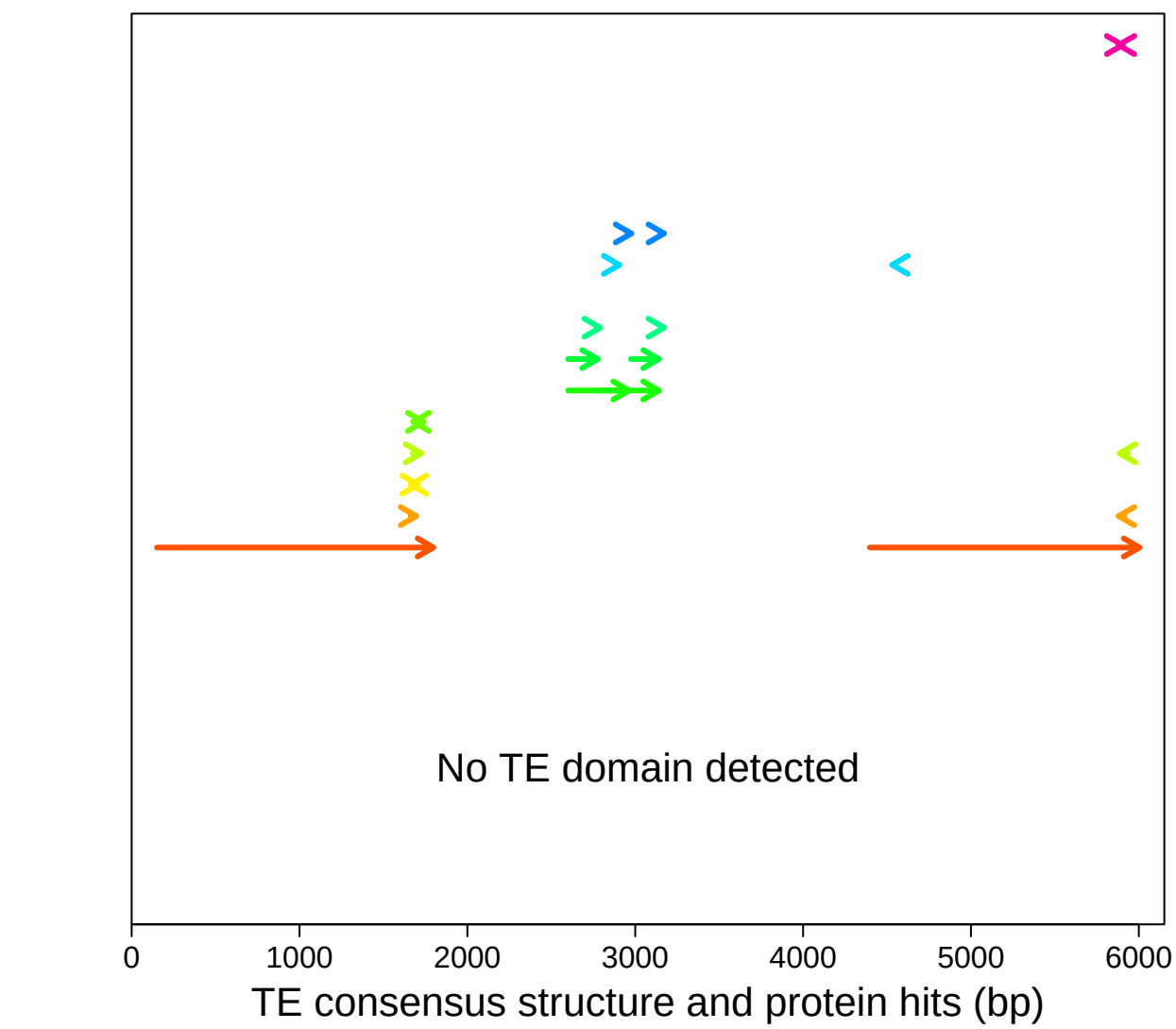
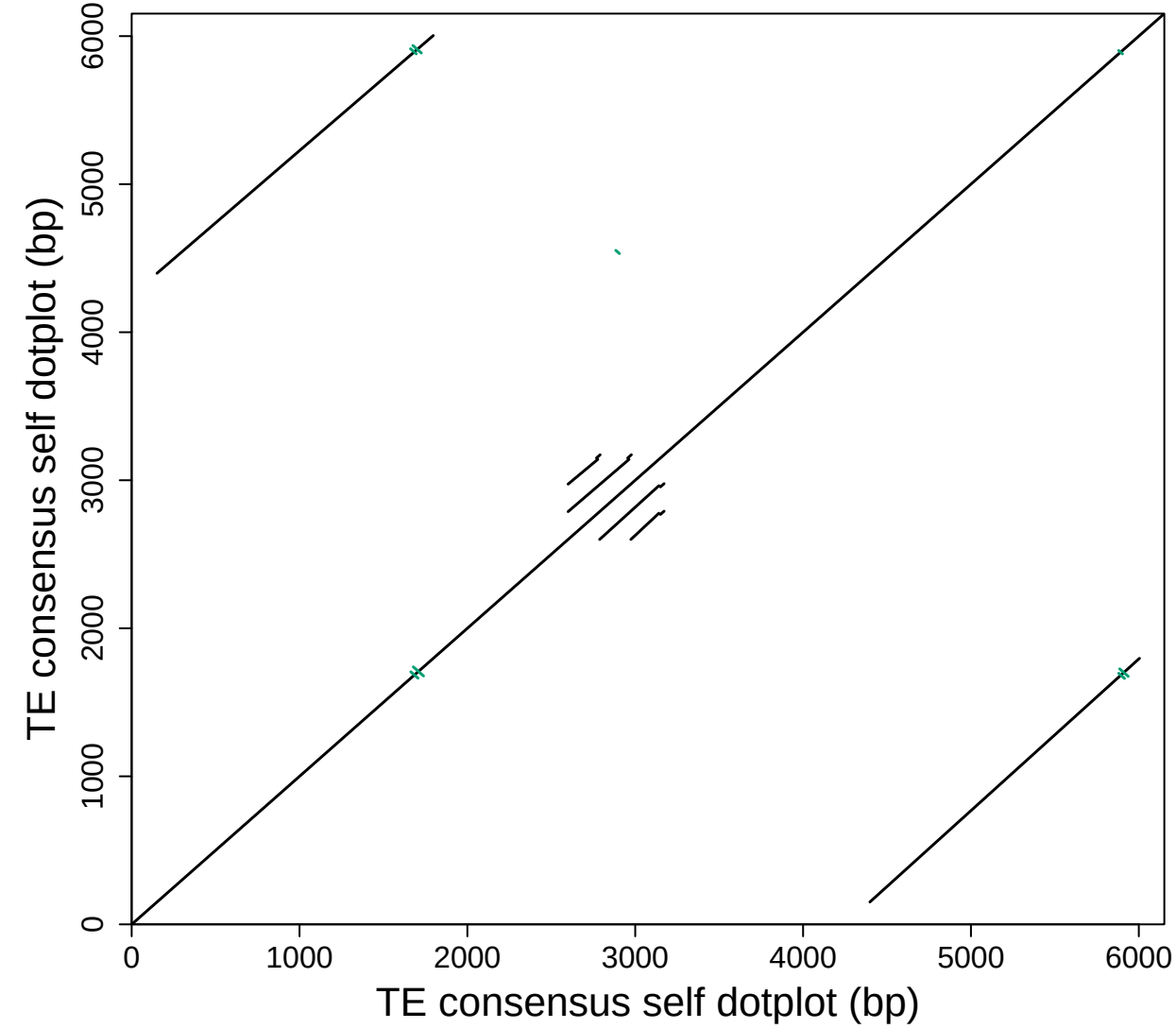
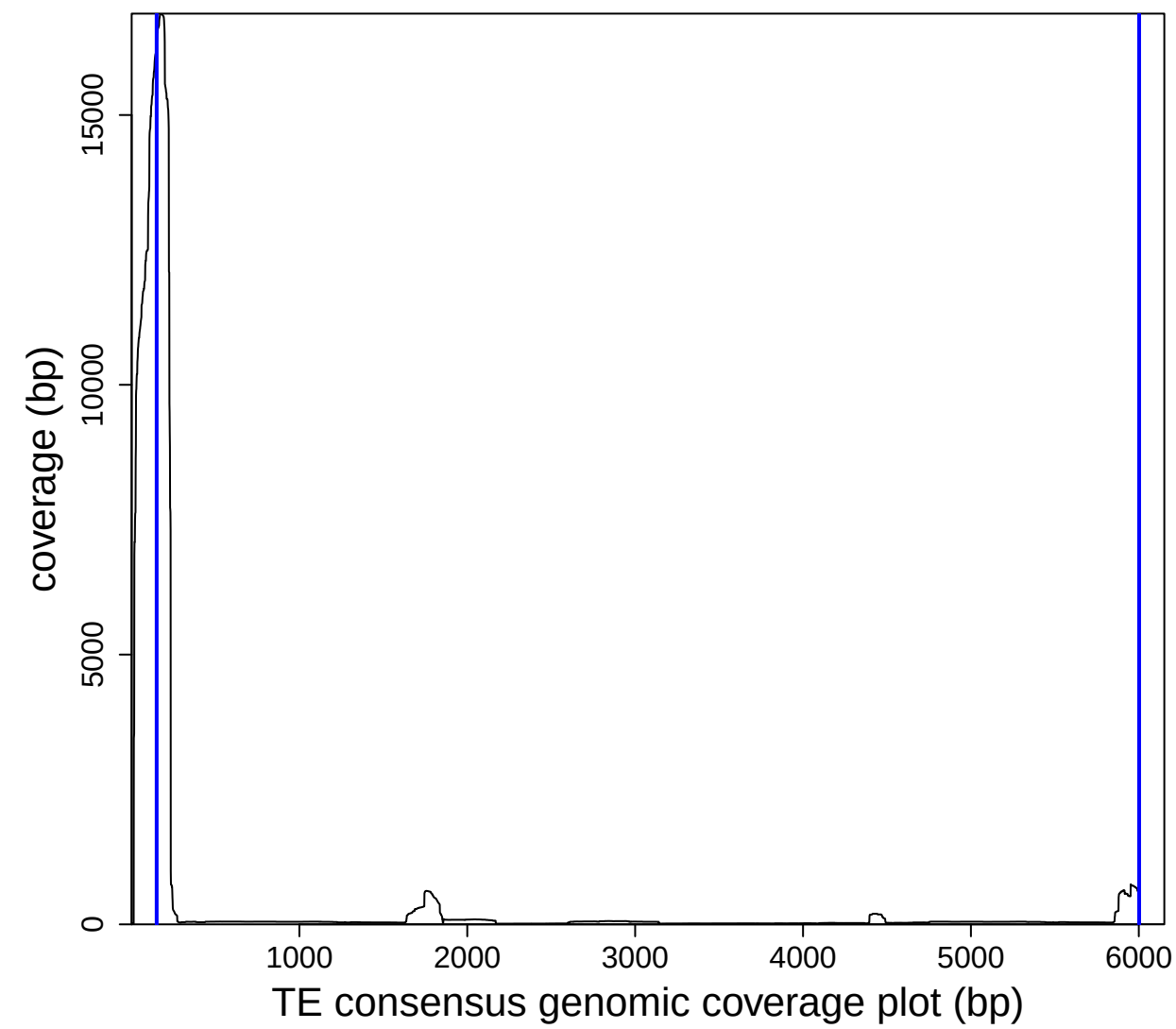
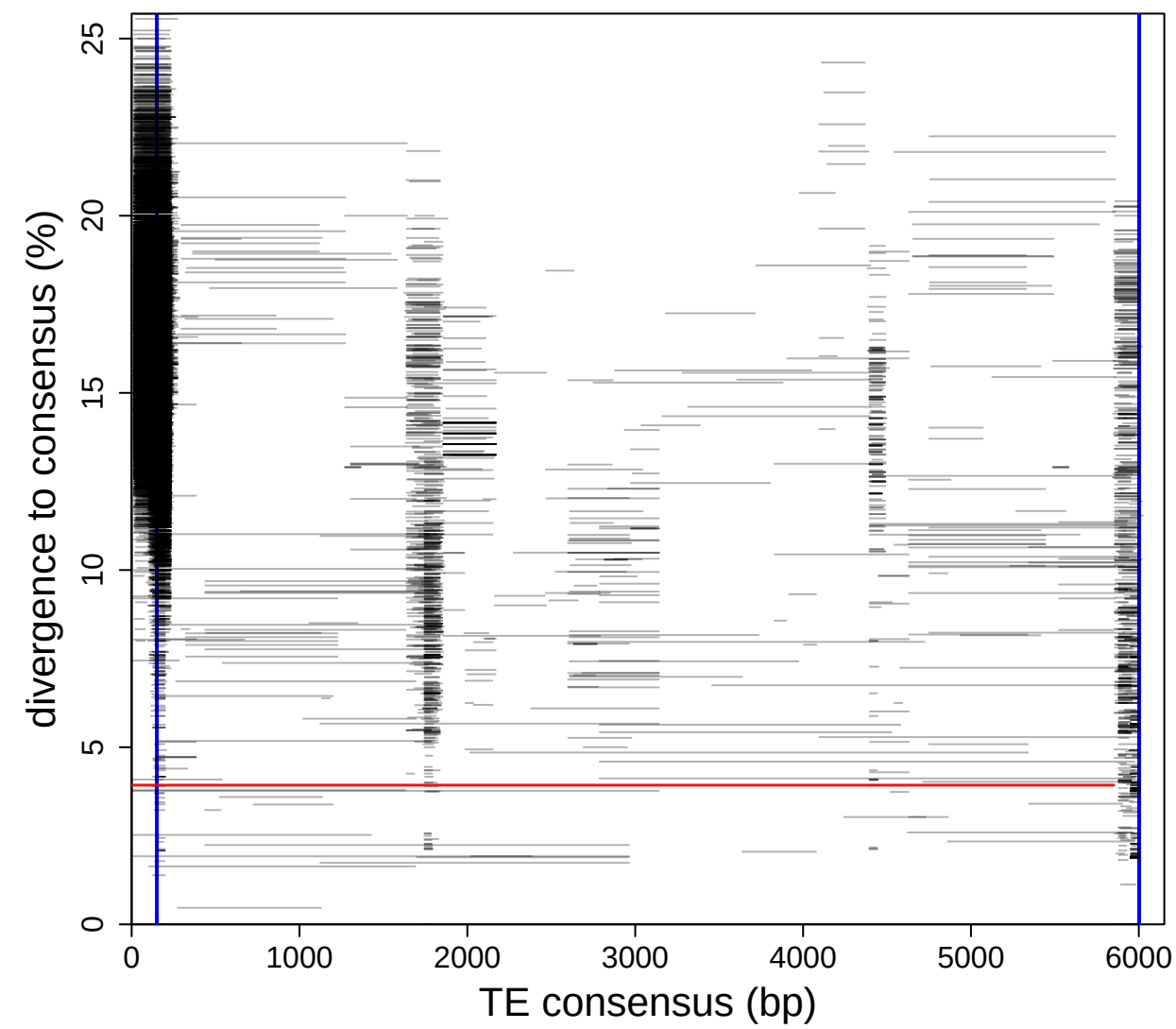
1.bed_fm 1.bed_0_0_n.bed_g 1.bed_fm 2.bed_0_0_bcln.fa aln.
size: 5847bp; fragments: 21449; full length: 1 (>=5262.3bp)

After TEtrimmer 5847 bp



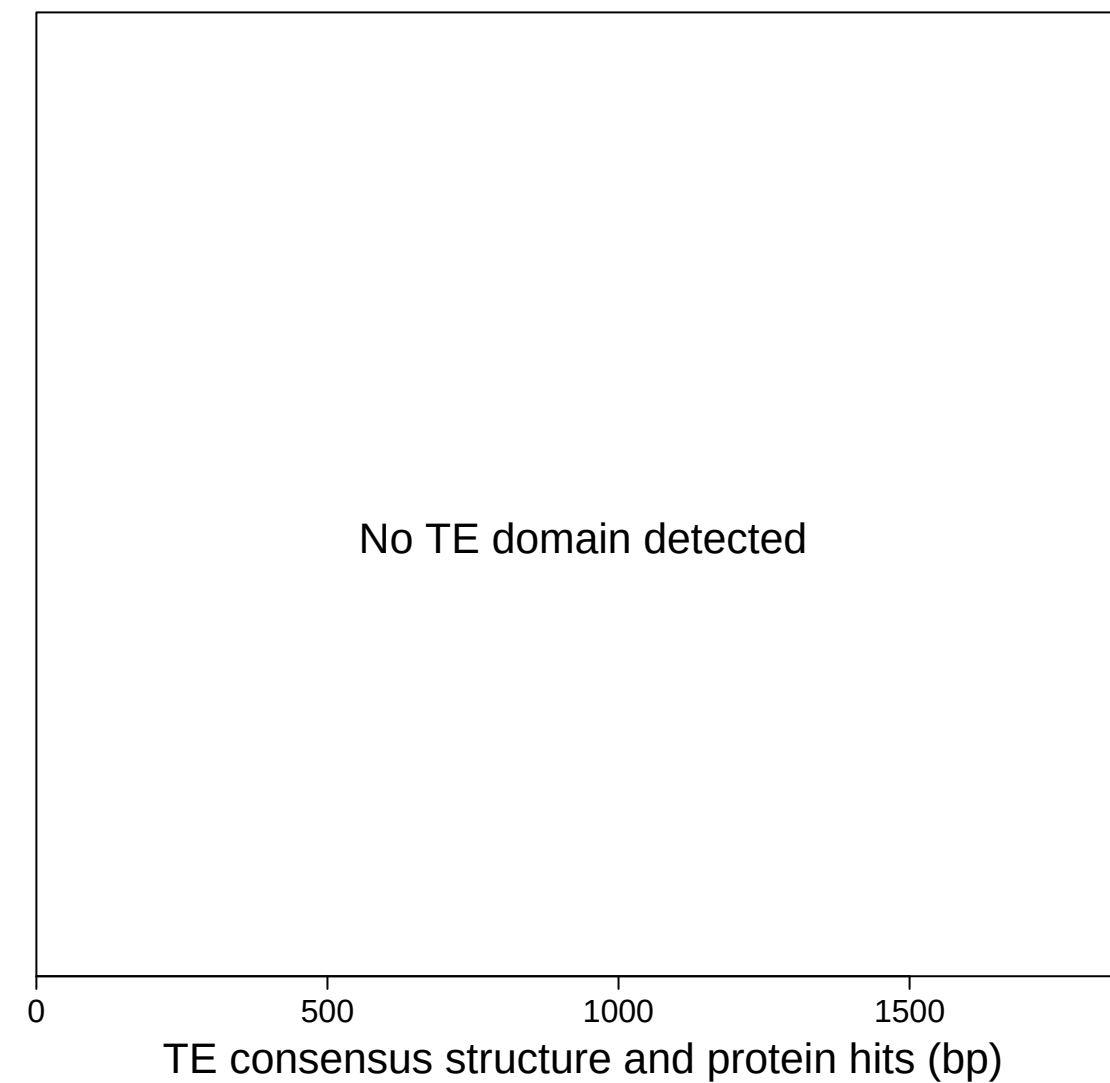
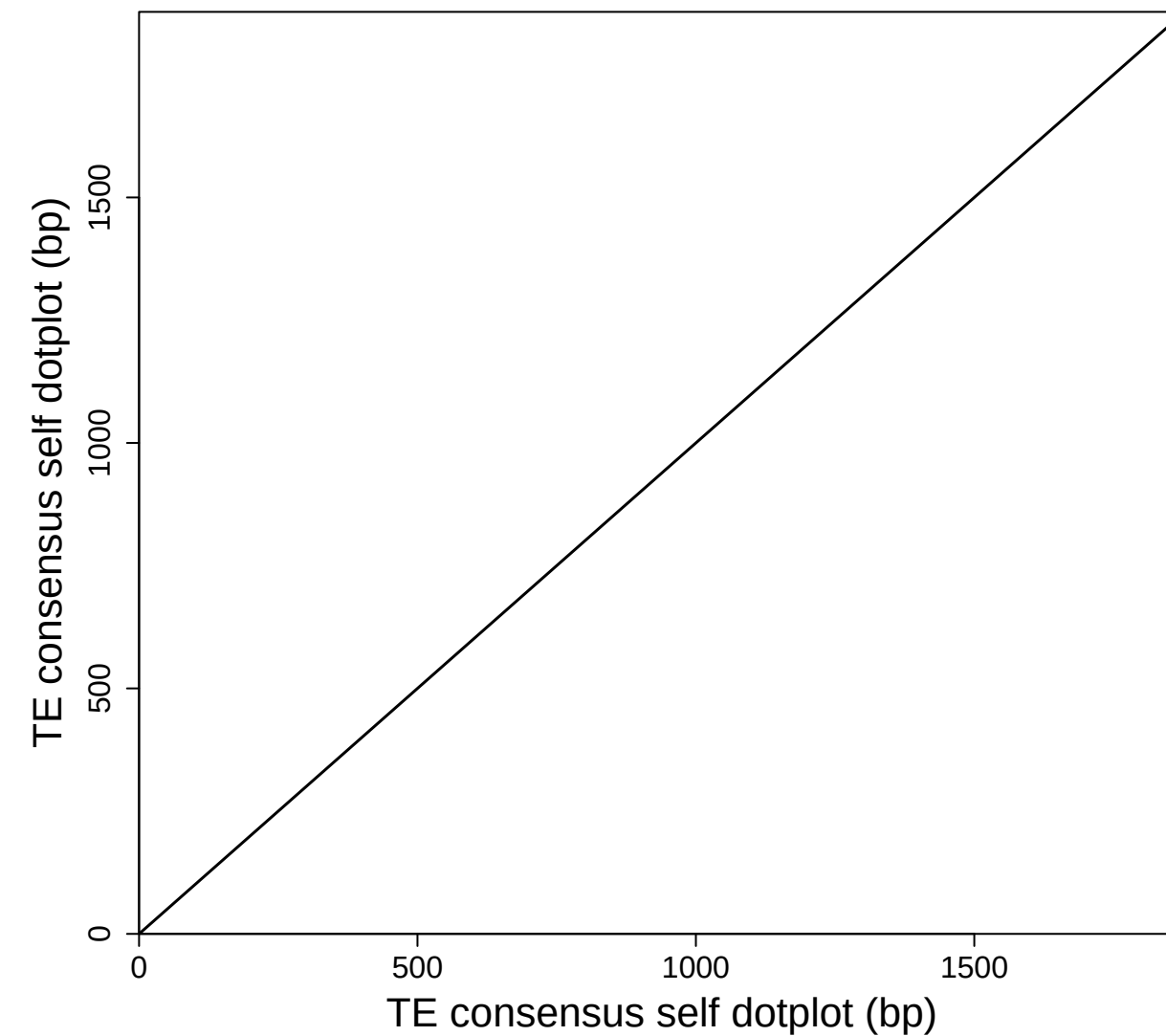
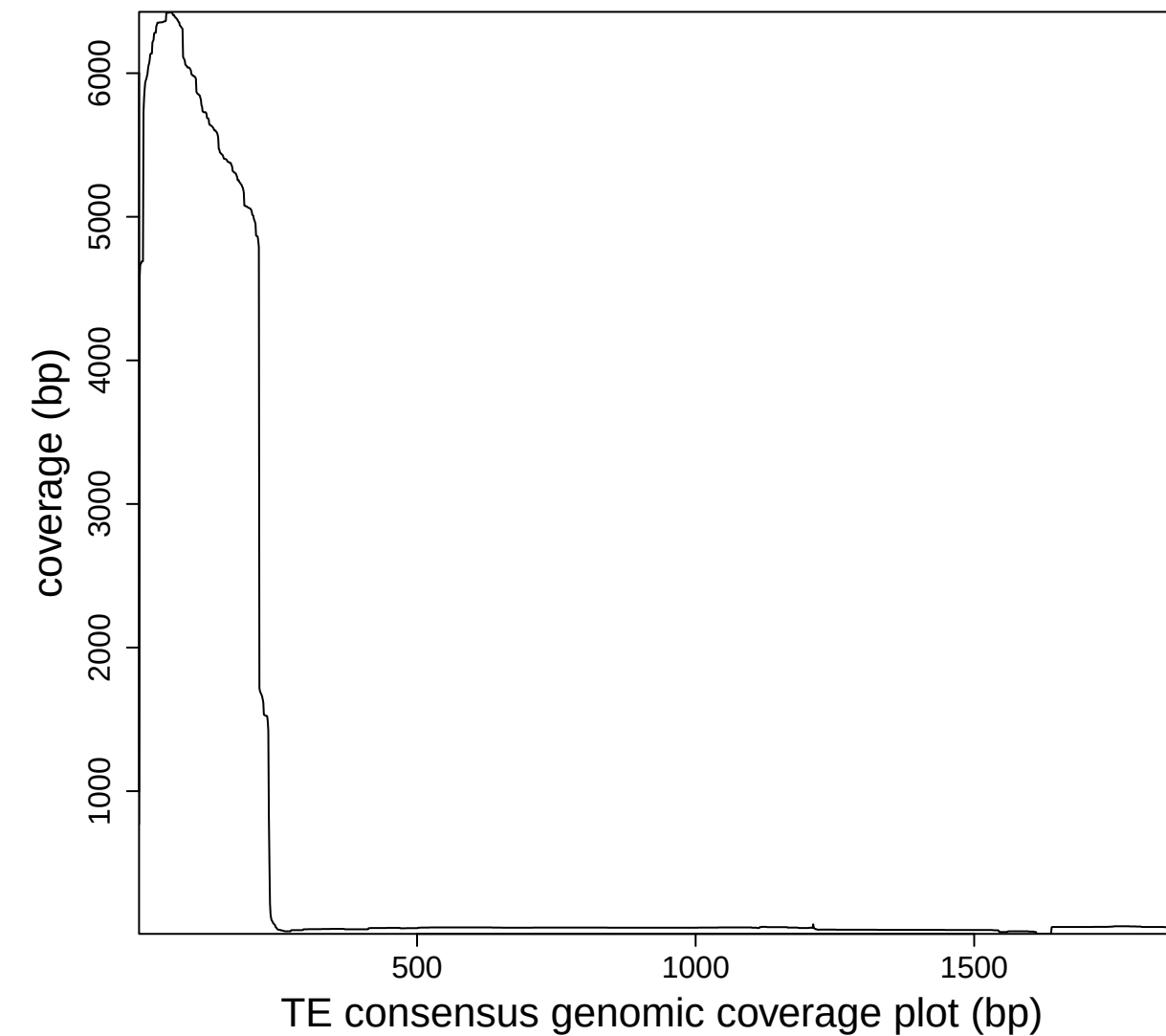
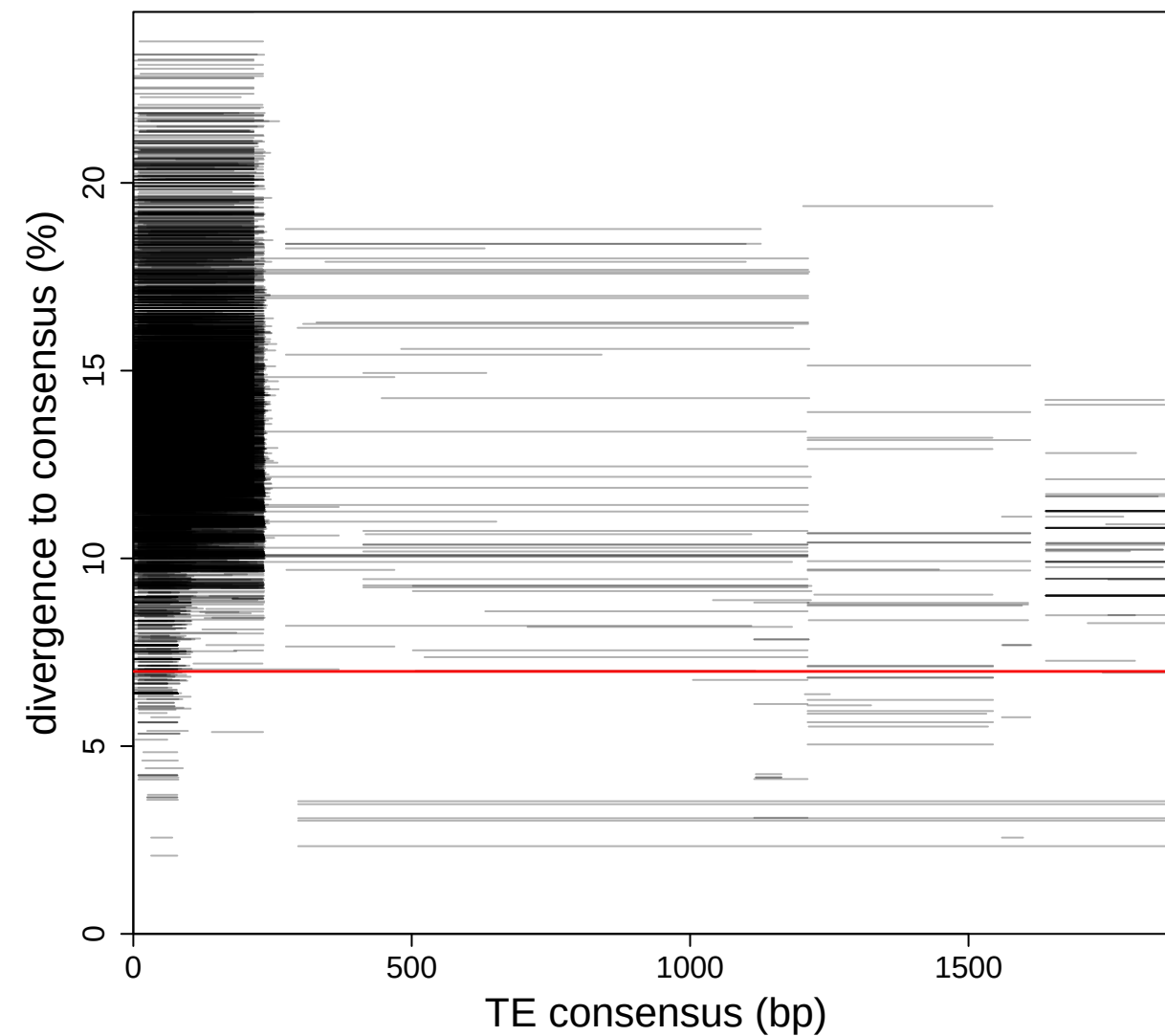
.bed uf.bed g 1.bed fm 1.bed 0 0 n.bed g 1.bed fm 2.bed (
size: 6152bp; fragments: 18978; full length: 1 (>=5536.8bp)

After TEtrimmer Extended plot Blue lines are boundaries

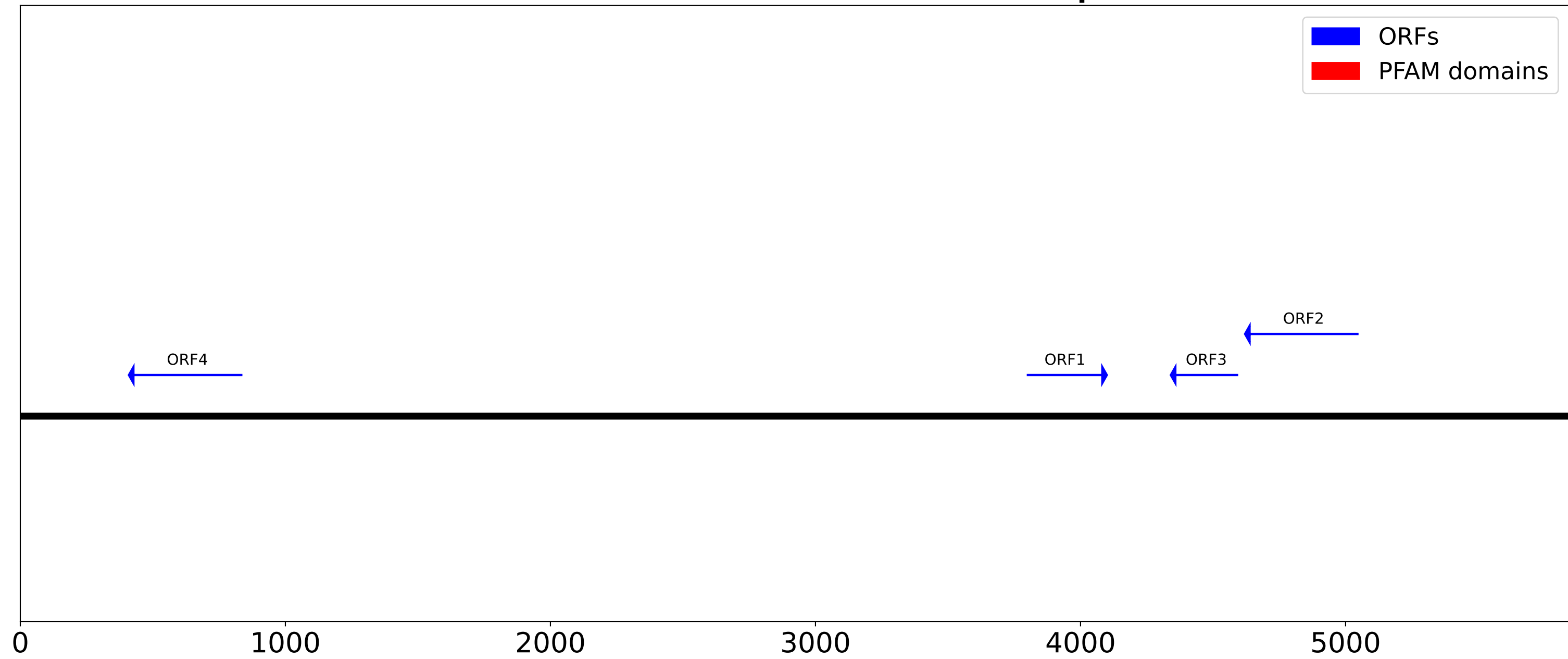


TE: rnd_5_family_15568
size: 1878bp; fragments: 6632; full length: 1 (≥ 1690.2 bp)

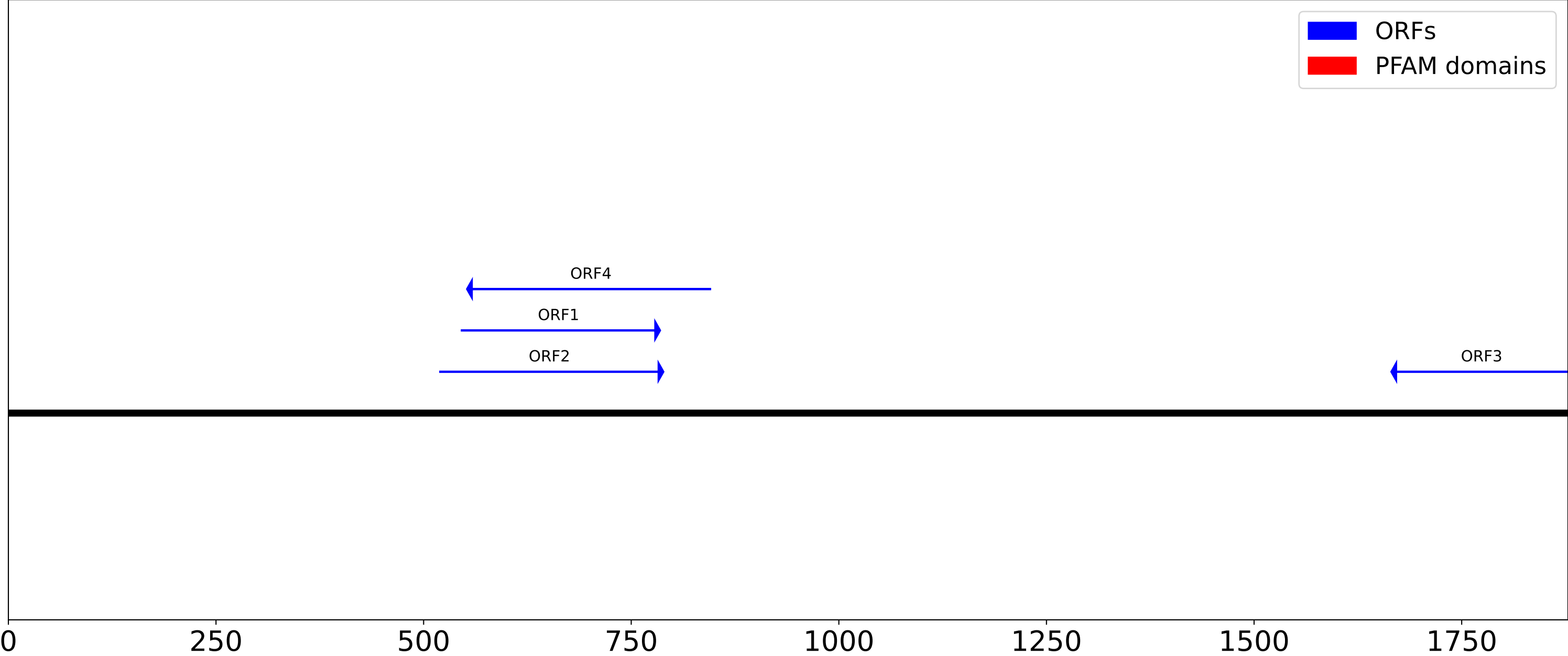
Before TEtrimmer 1878 bp



After TEtrimmer ORF and PFAM domain plot



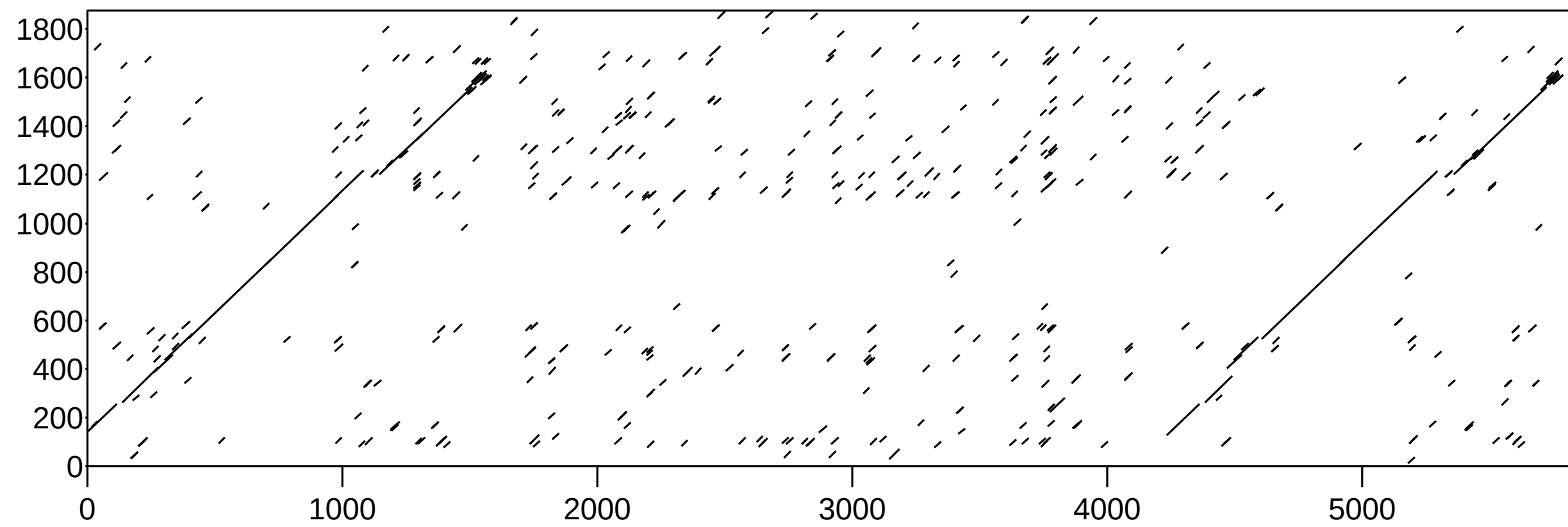
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)