

MSA length = 6118

[illegible]

Start crop Point

End crop Point

0

10

0

1000

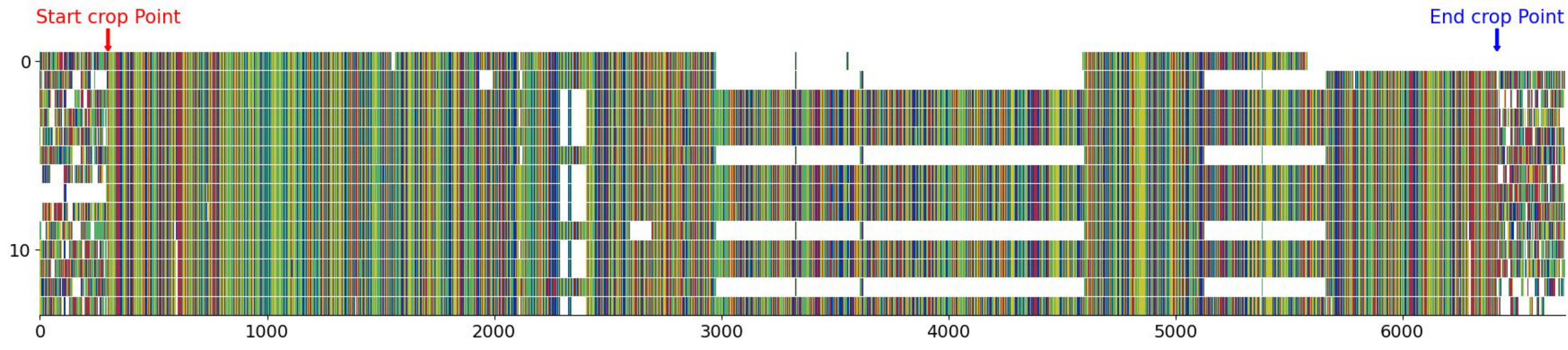
2000

3000

4000

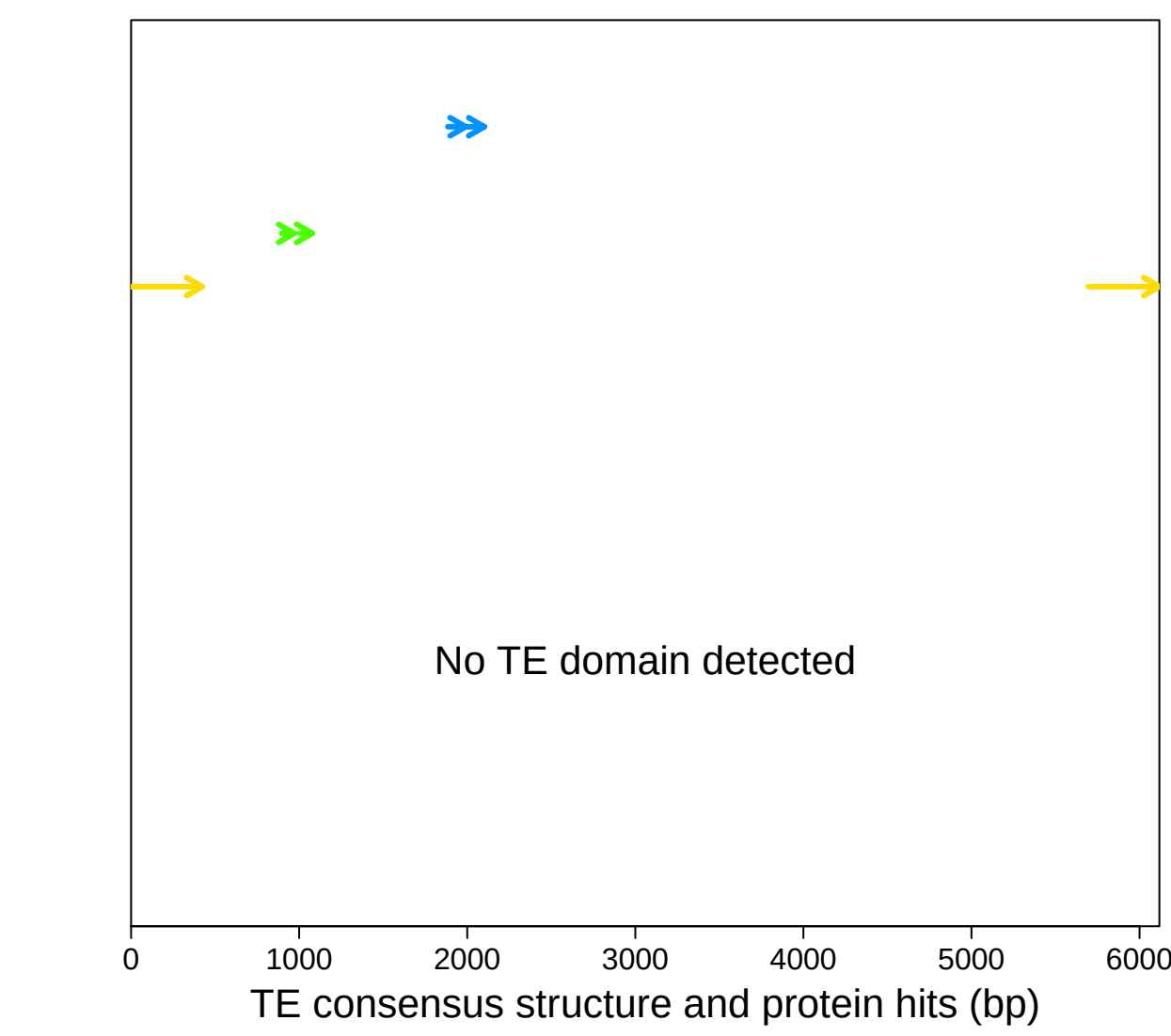
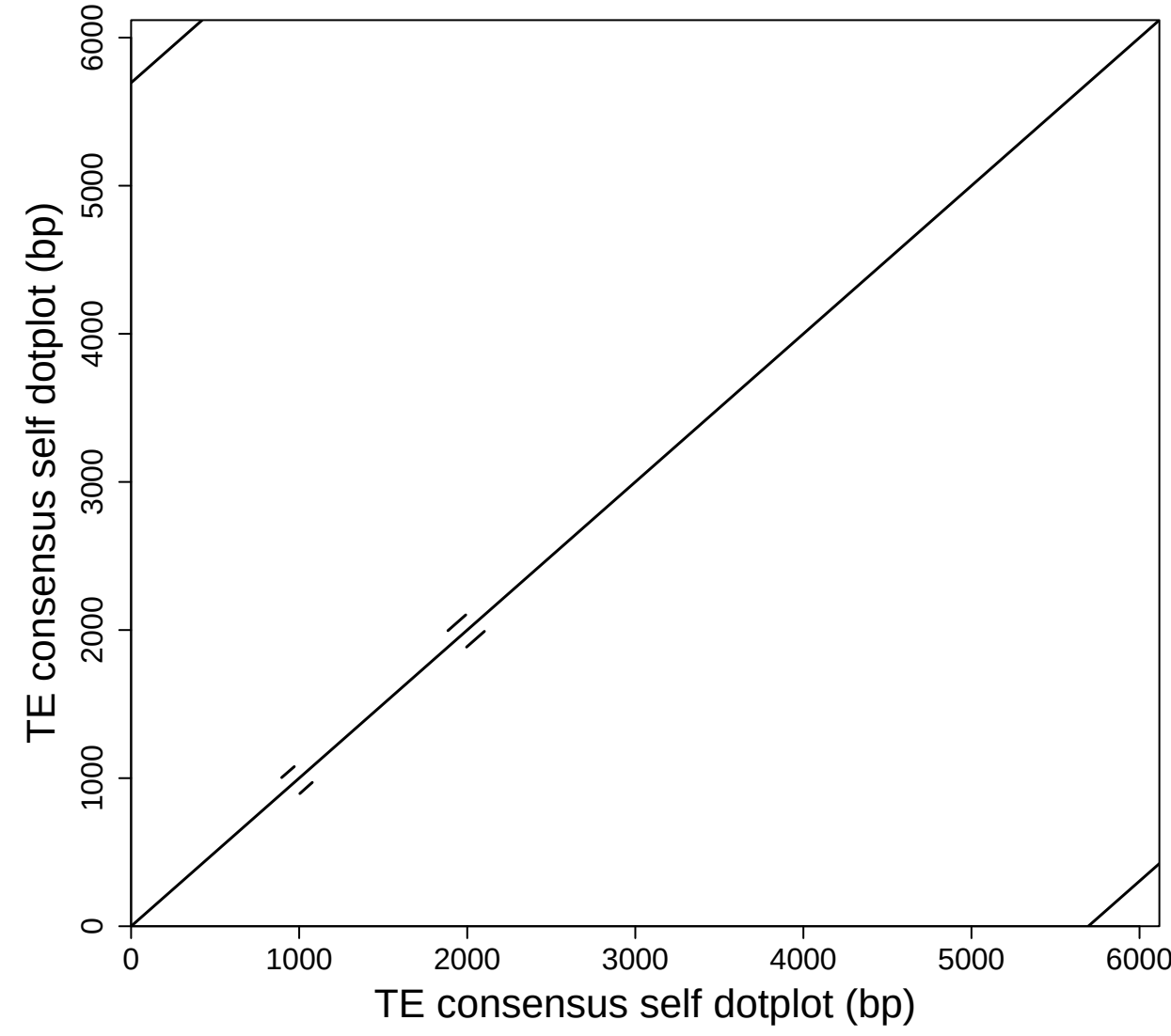
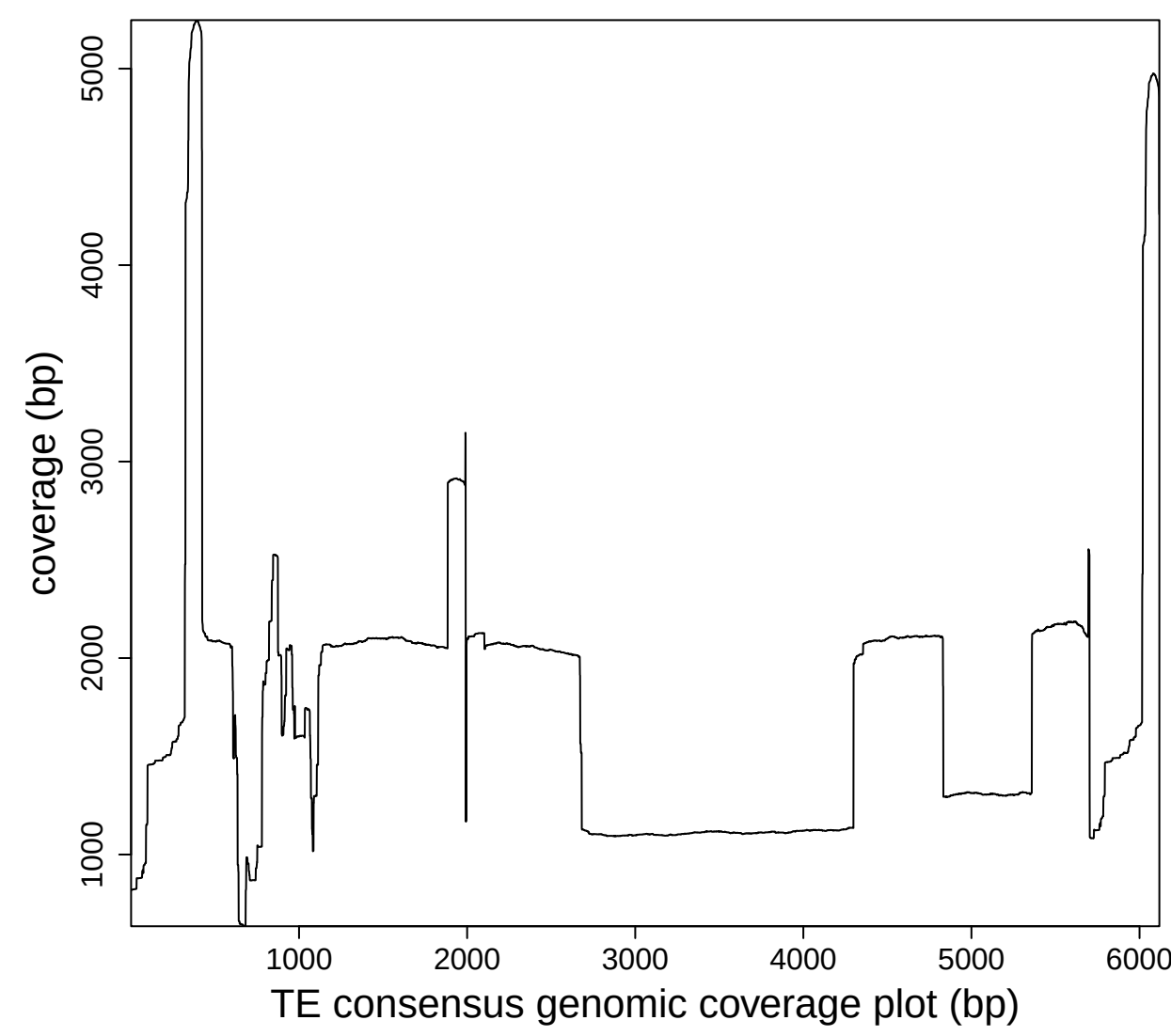
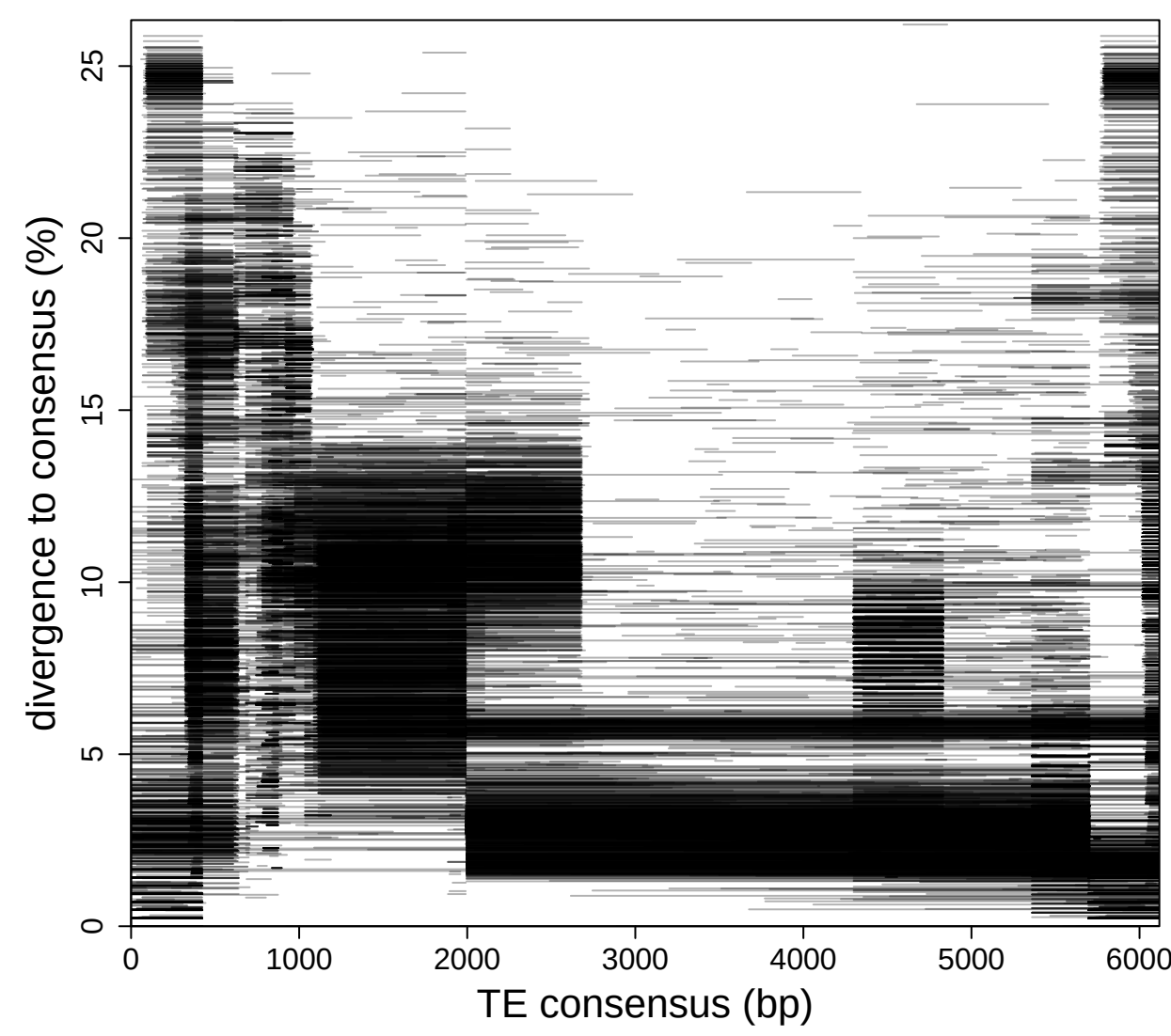
5000

6000



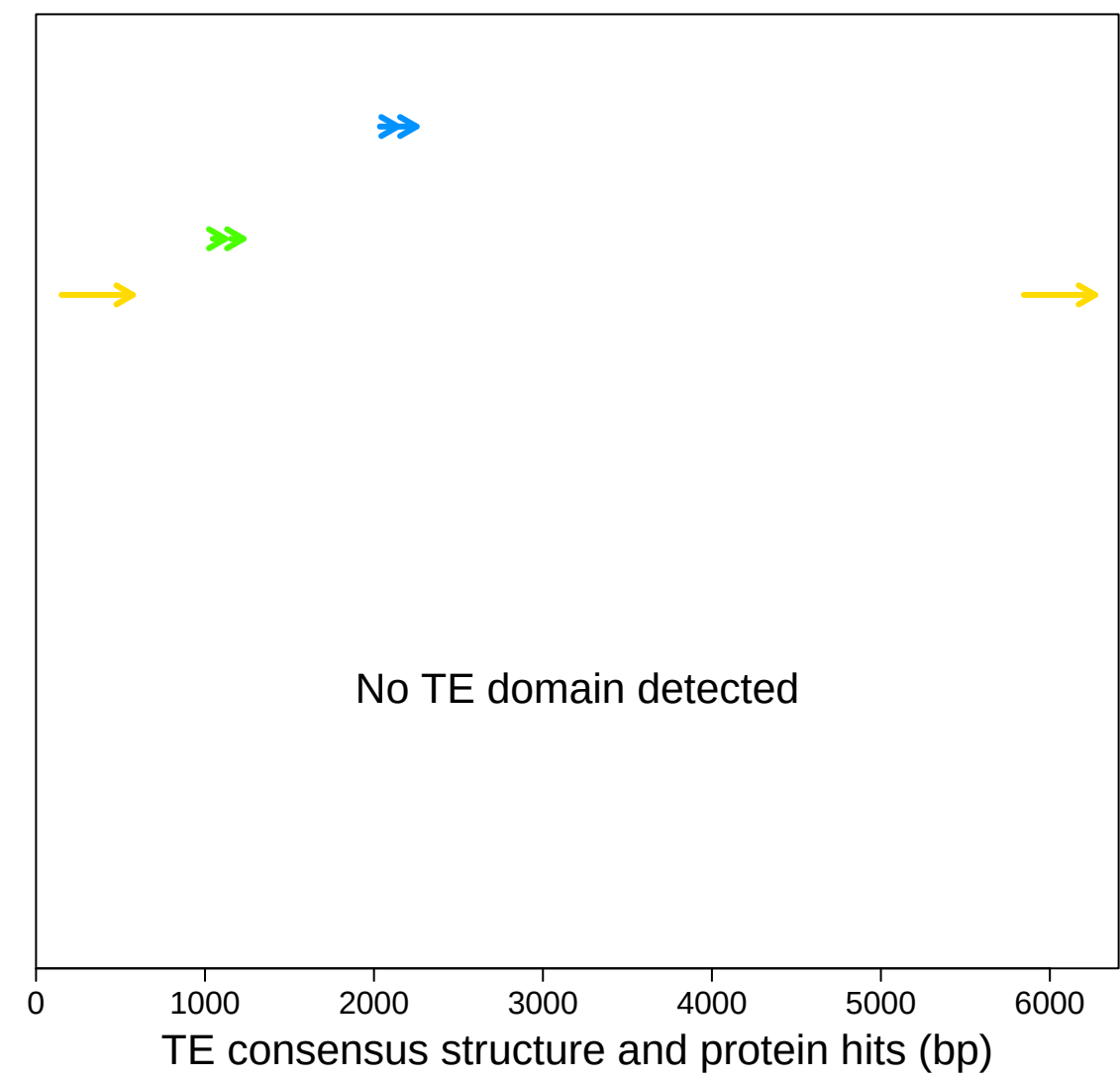
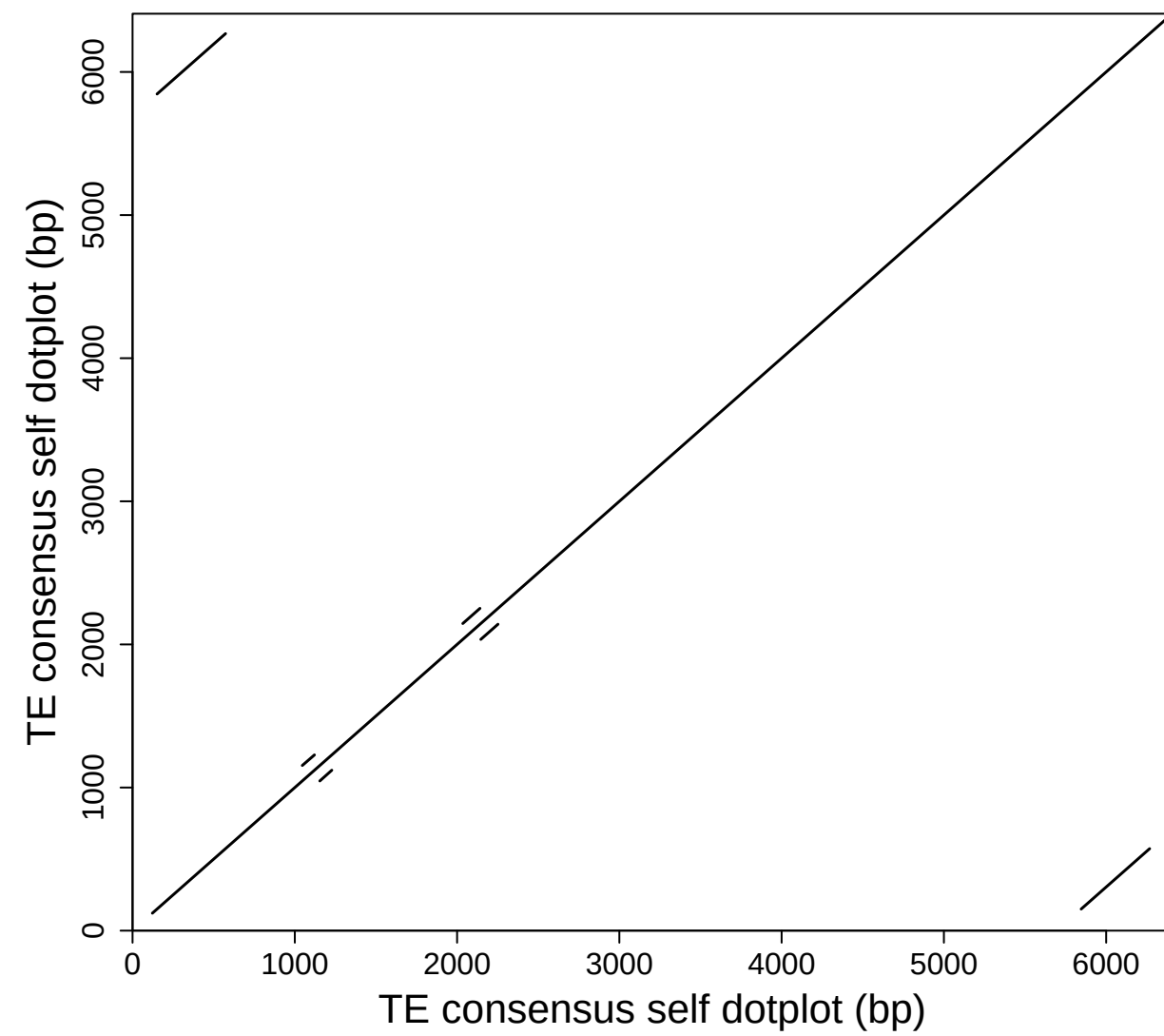
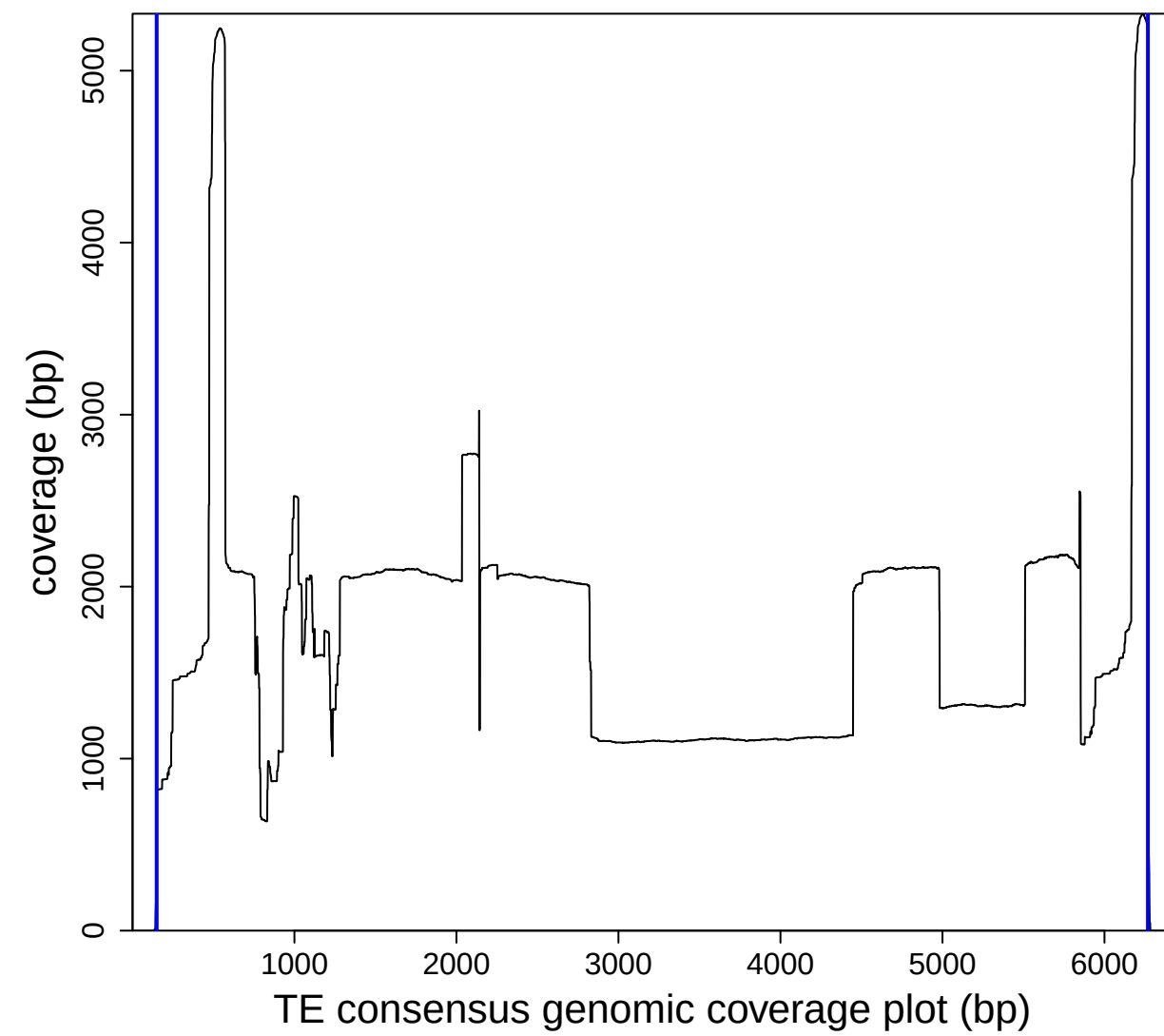
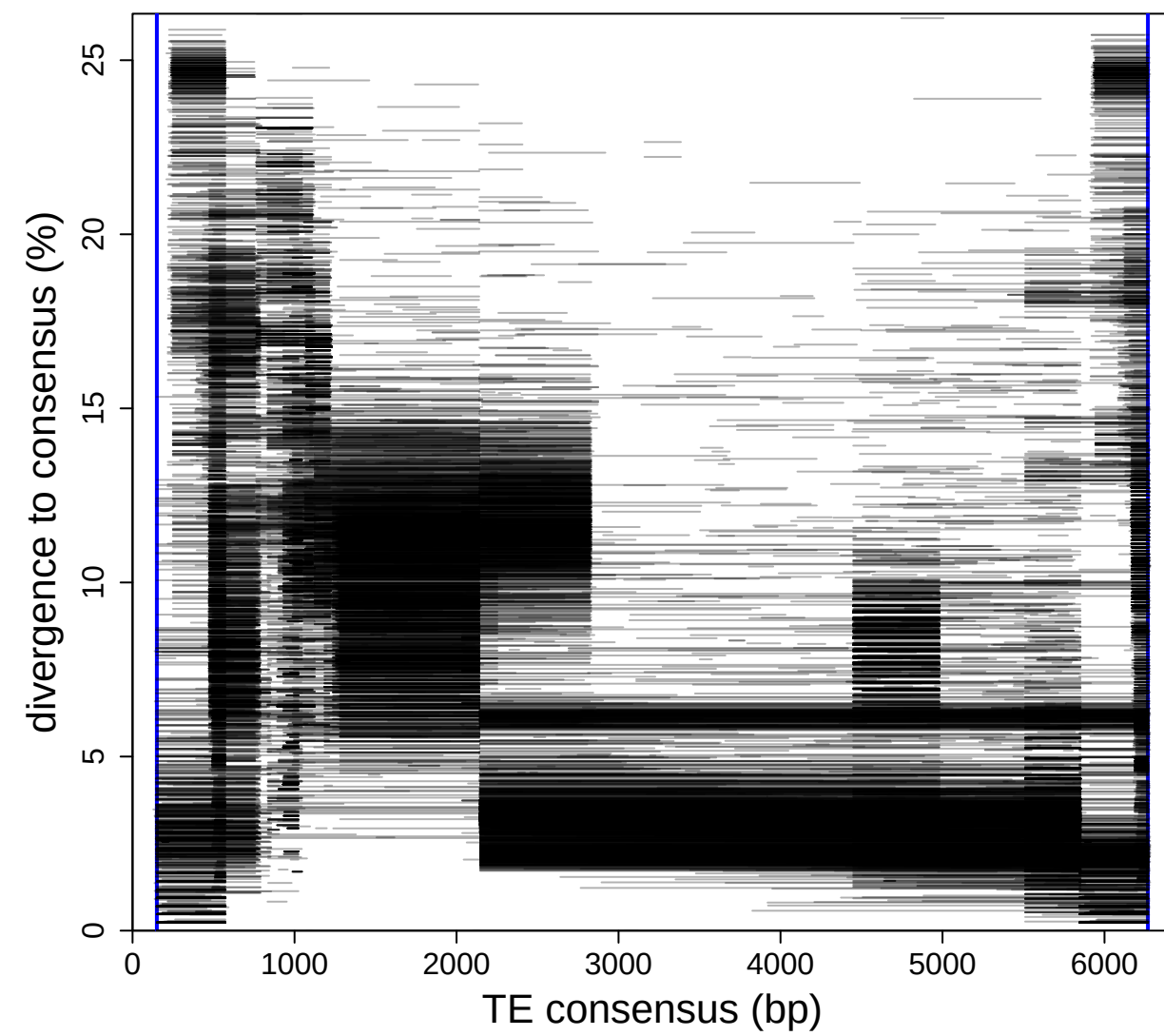
bed_uf.bed g_3.bed fm_1.bed 0_0_bcln.fa aln.fa cl.fa gs.fa c
size: 6118bp; fragments: 18939; full length: 0 (>=5506.2bp)

After TEtrimmer 6118 bp



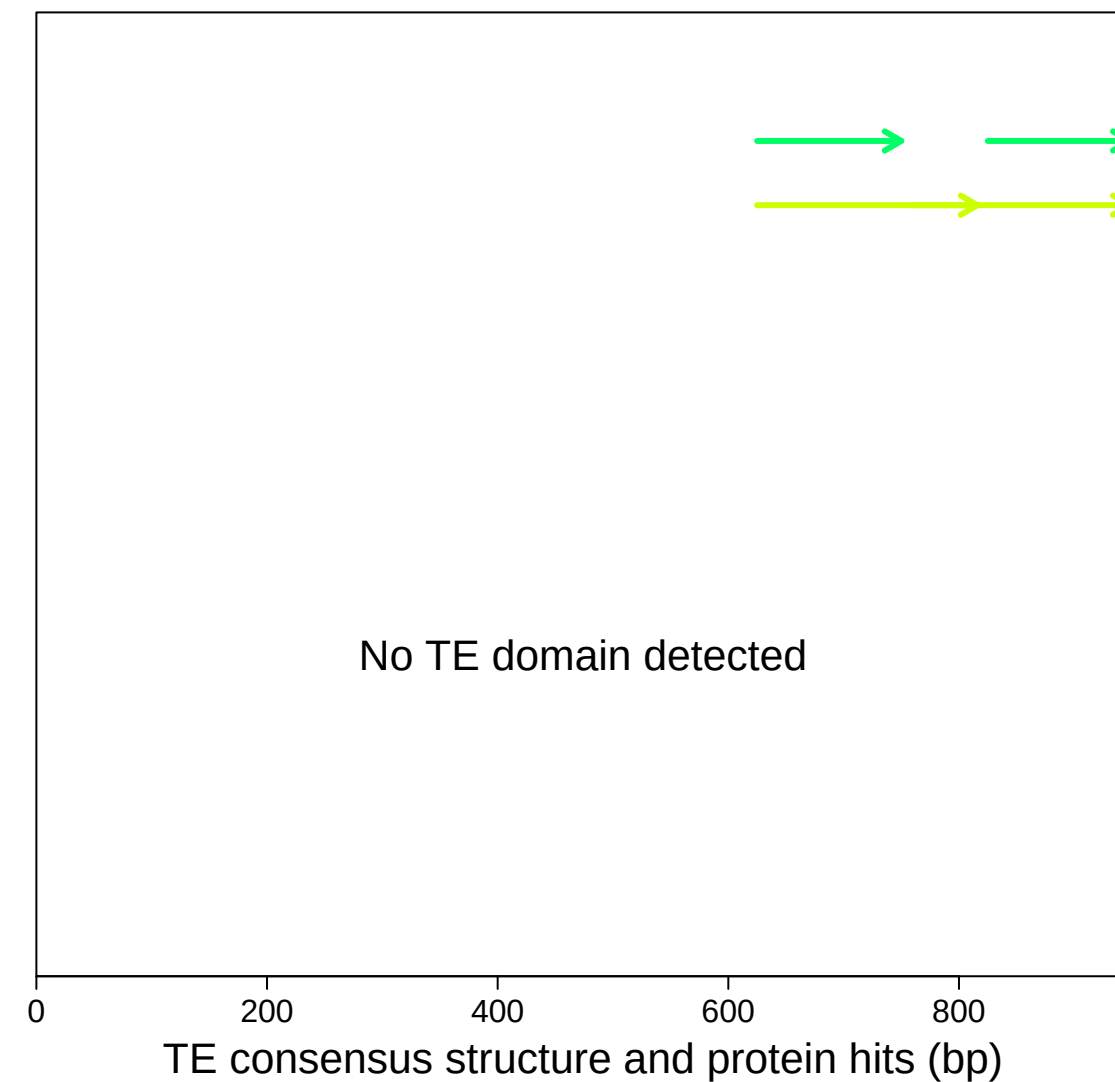
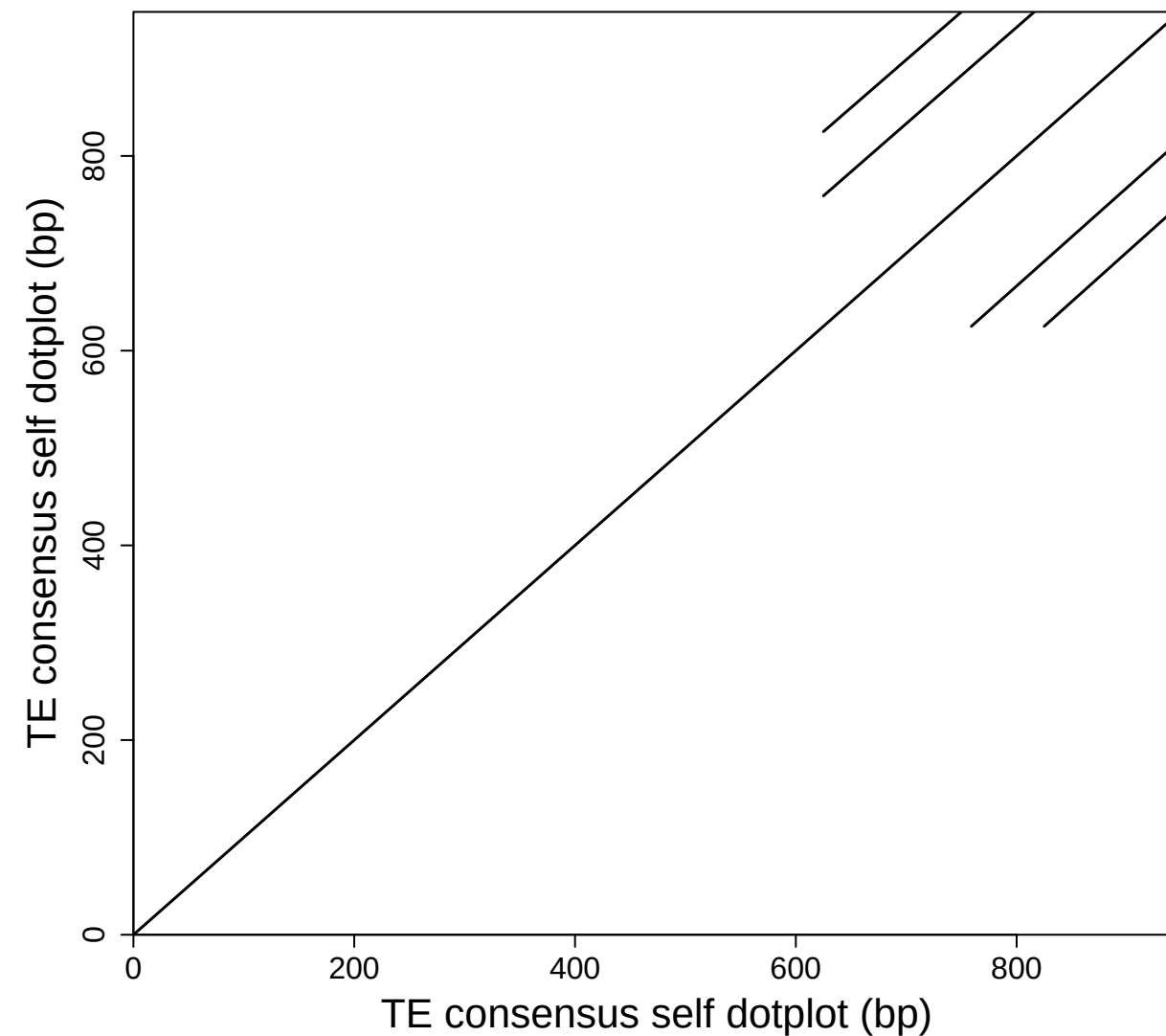
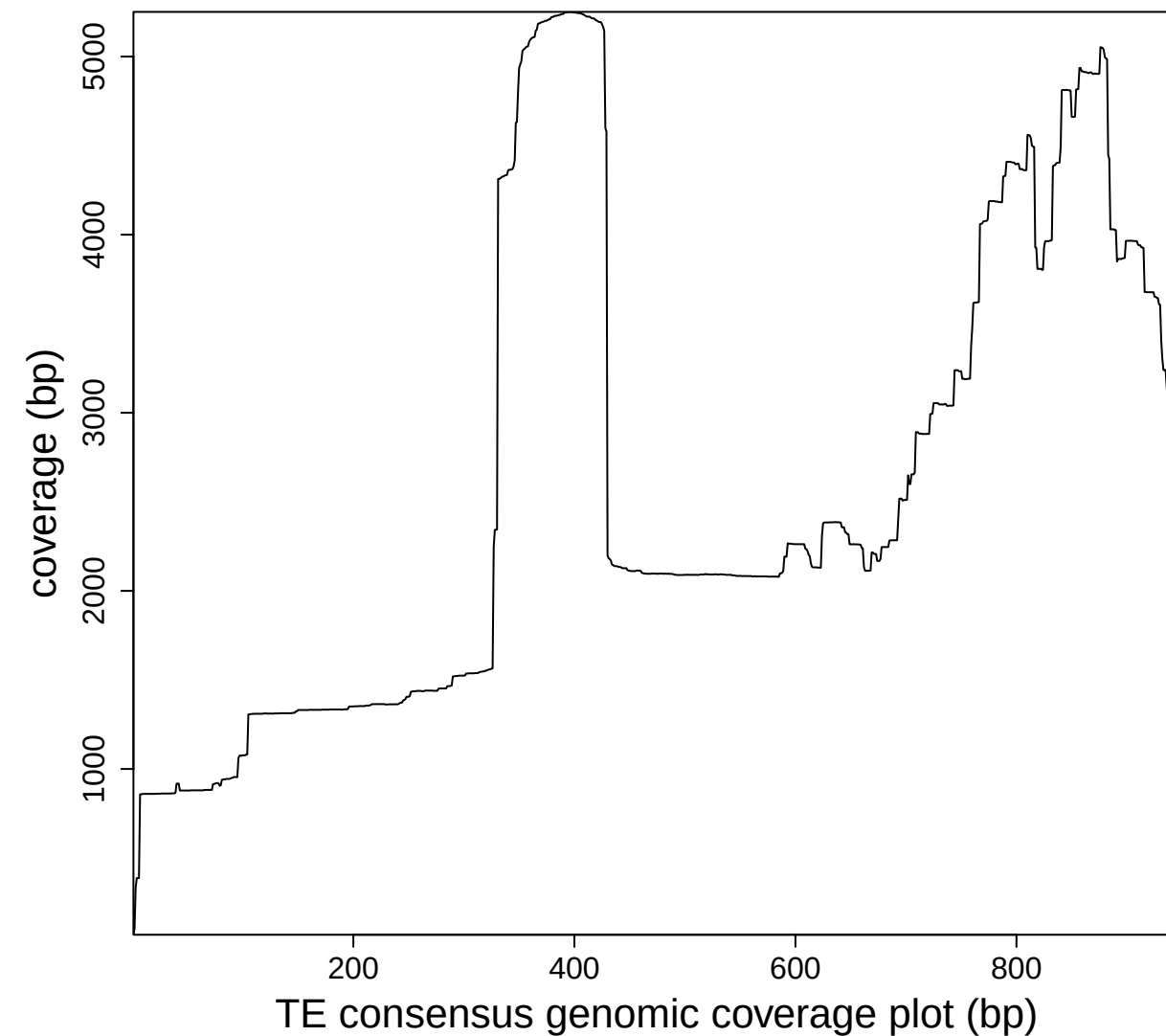
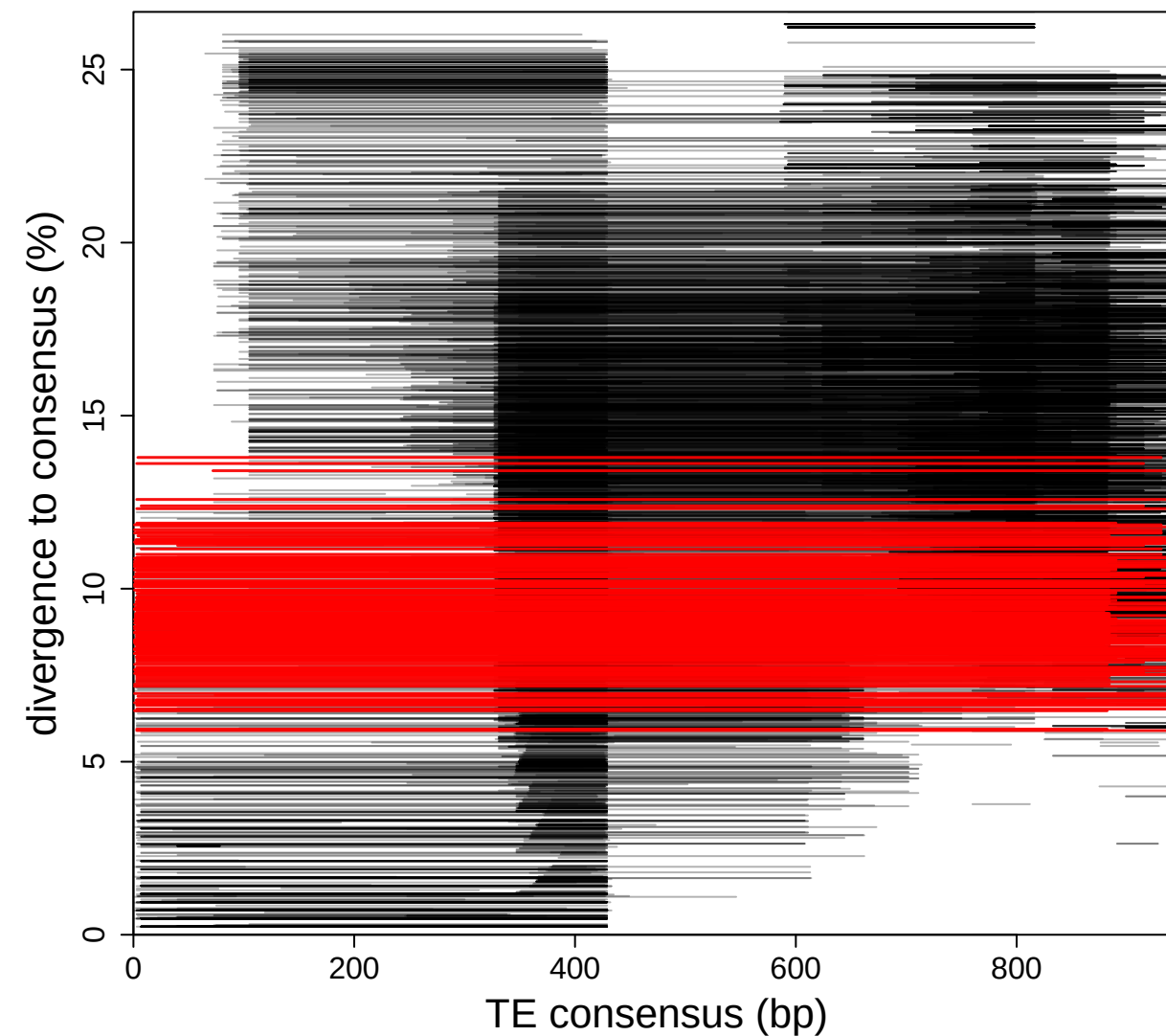
ly_1353.fasta.b.bed uf.bed g_3.bed fm_1.bed_0_0 bcln.fa aln.f
size: 6418bp; fragments: 19144; full length: 0 (>=5776.2bp)

After TEtrimmer Extended plot Blue lines are boundaries

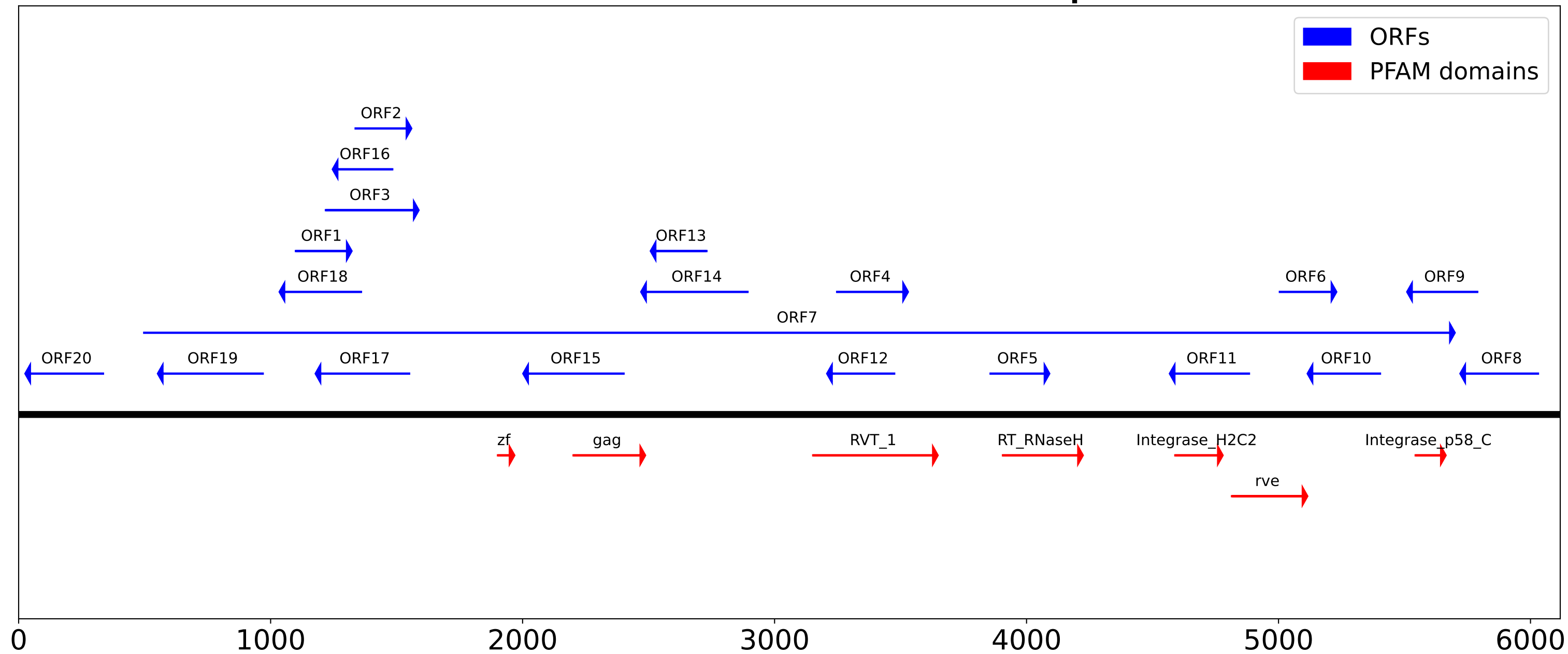


TE: rnd_5_family_1353
size: 948bp; fragments: 10200; full length: 222 (≥ 853.2 bp)

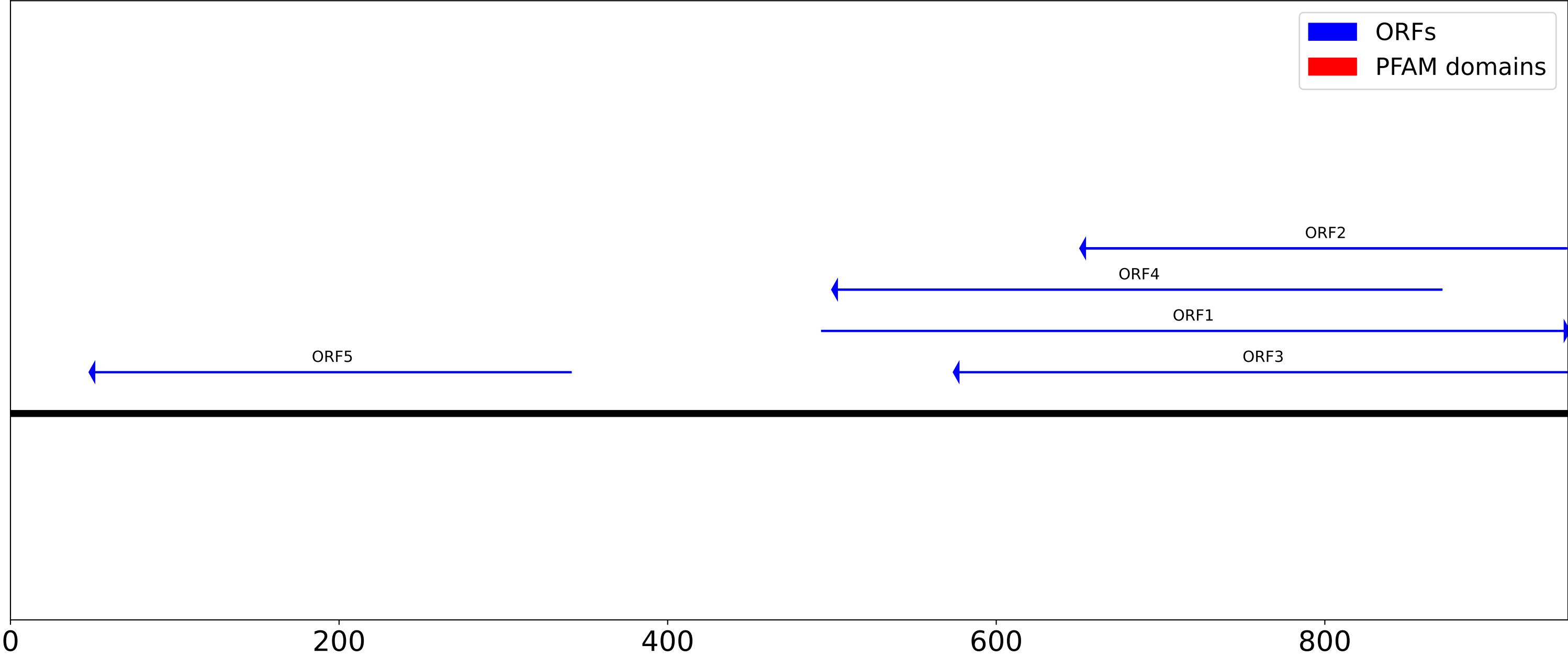
Before TEtrimmer 948 bp



After TETrimmer ORF and PFAM domain plot



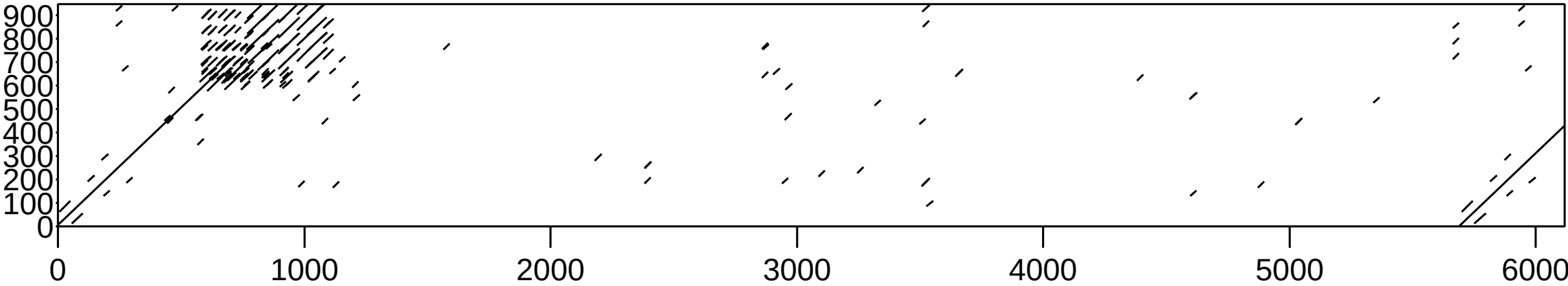
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

nsensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)