

Start crop Point

End crop Point

MSA length = 5672



Start crop Point

End crop Point

0

10

0

1000

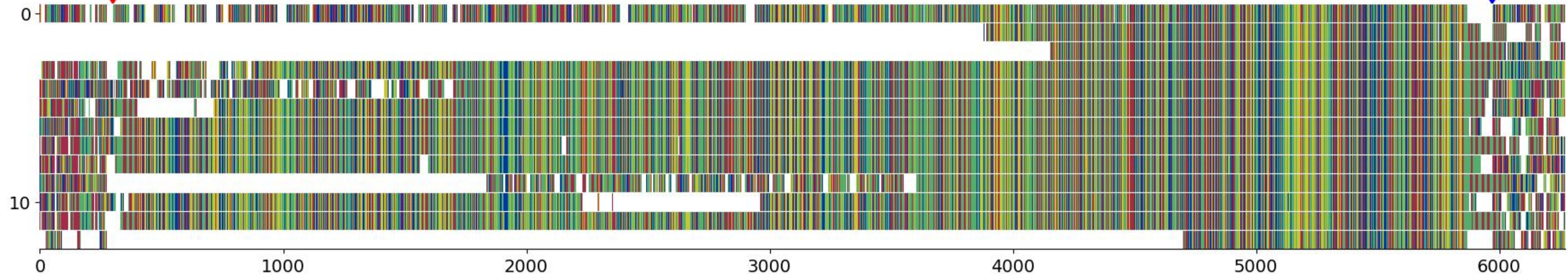
2000

3000

4000

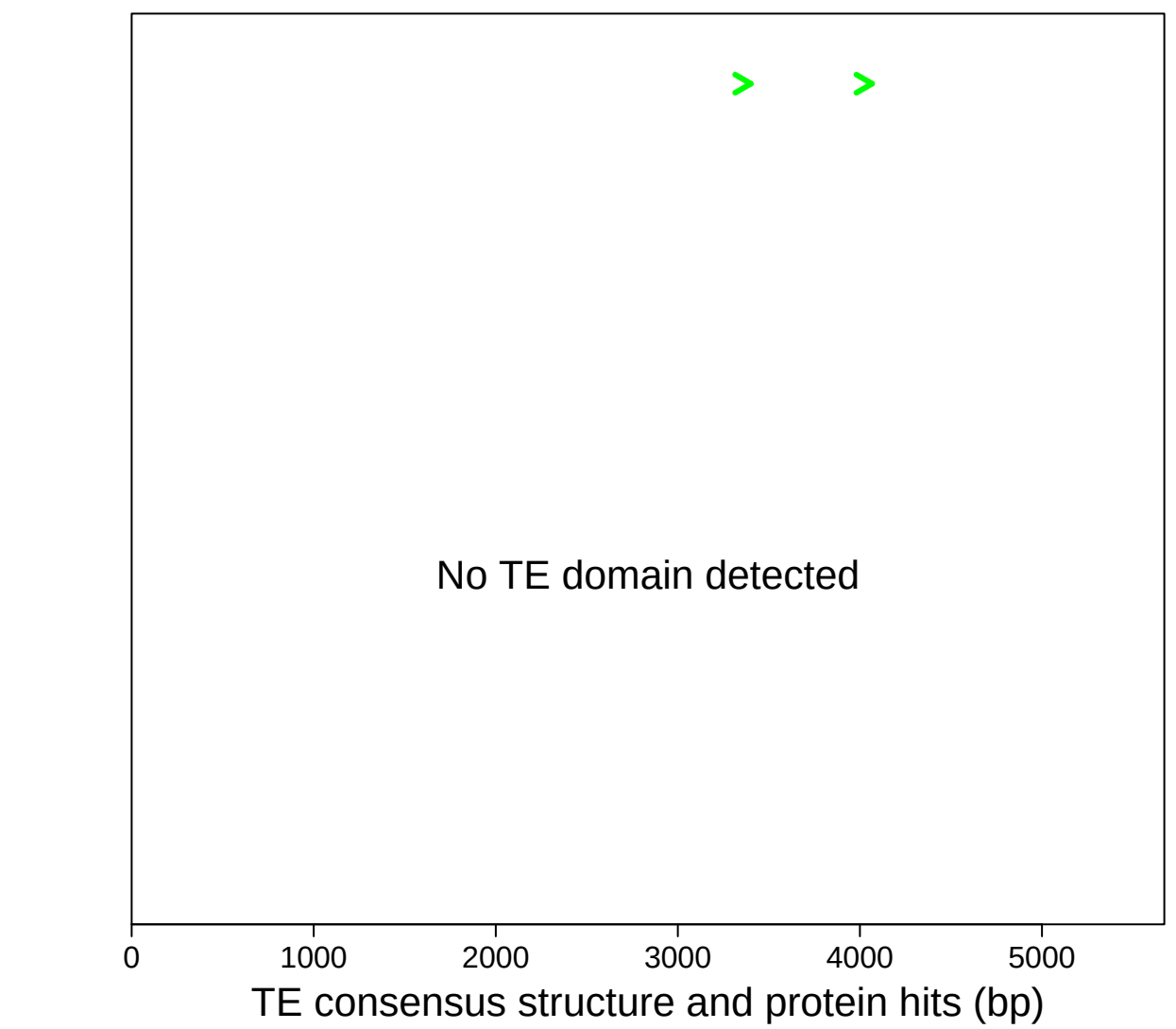
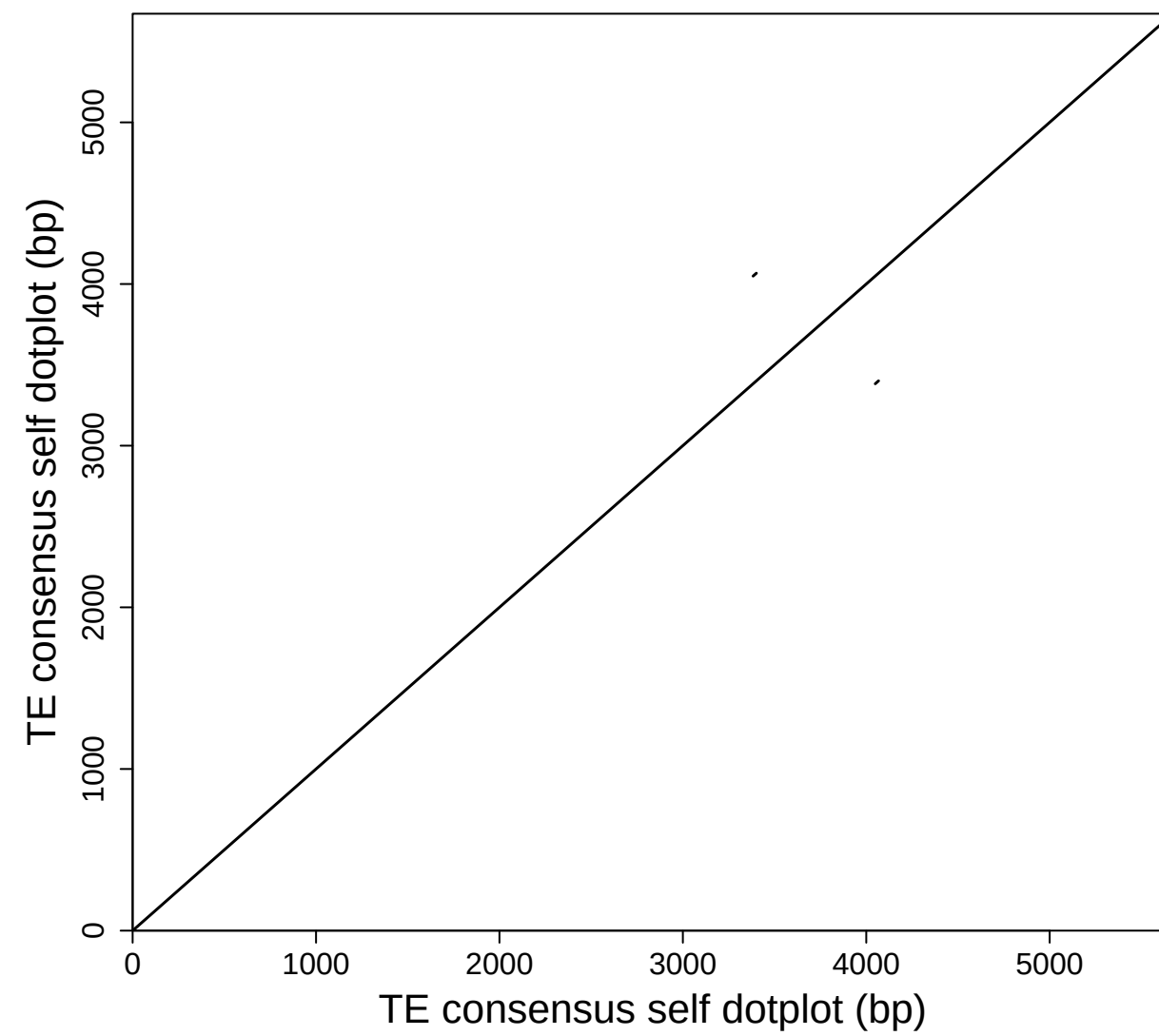
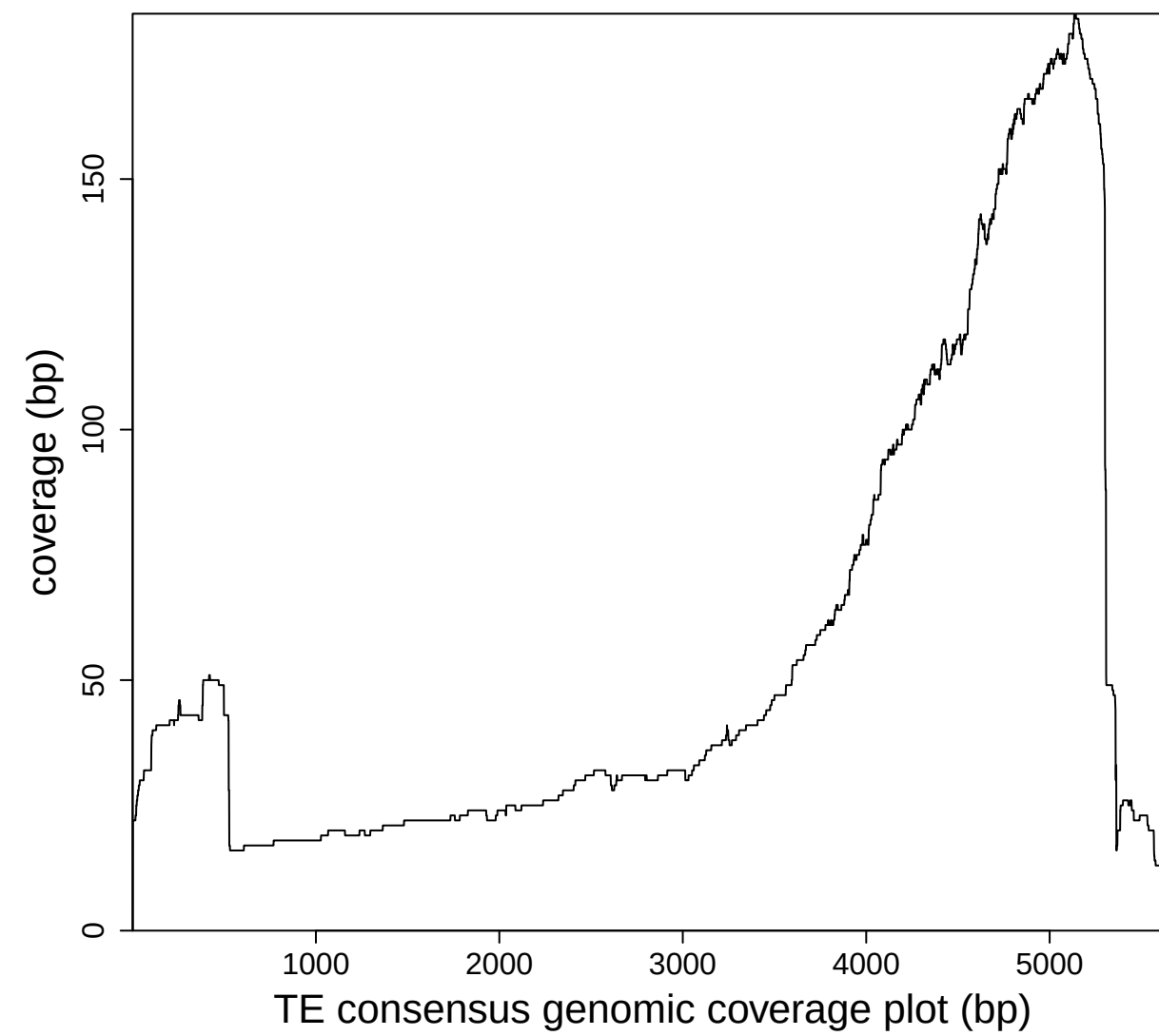
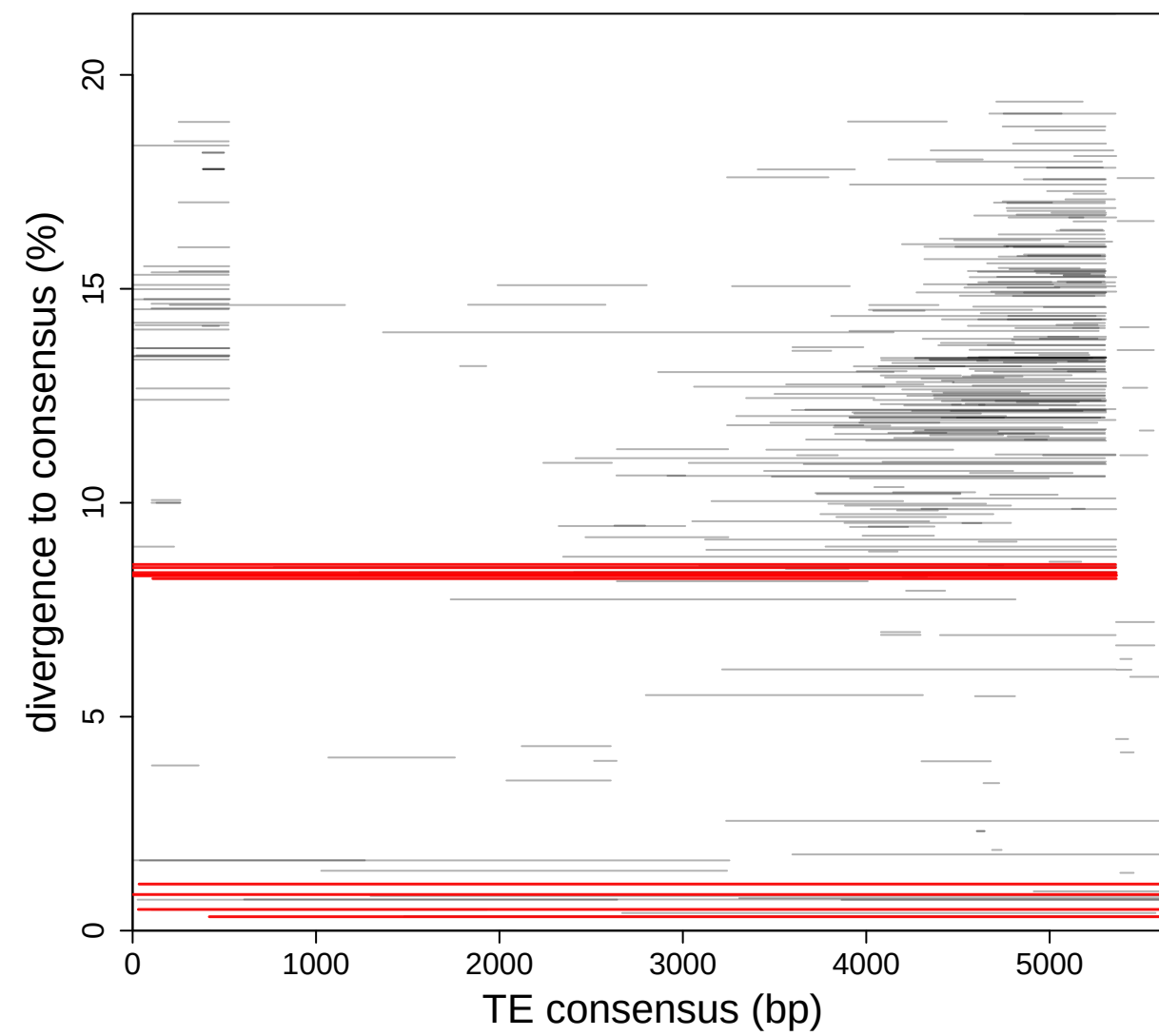
5000

6000



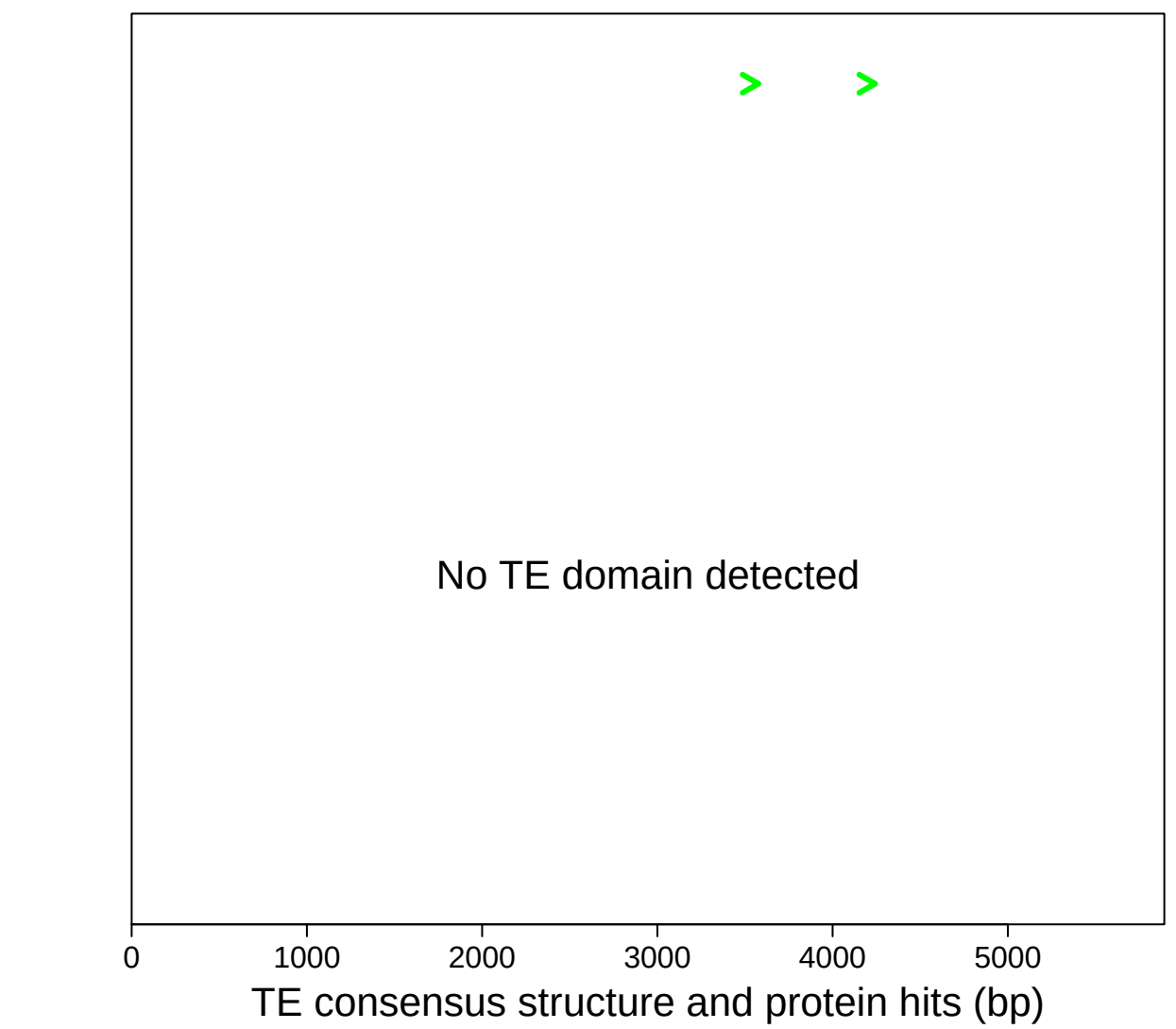
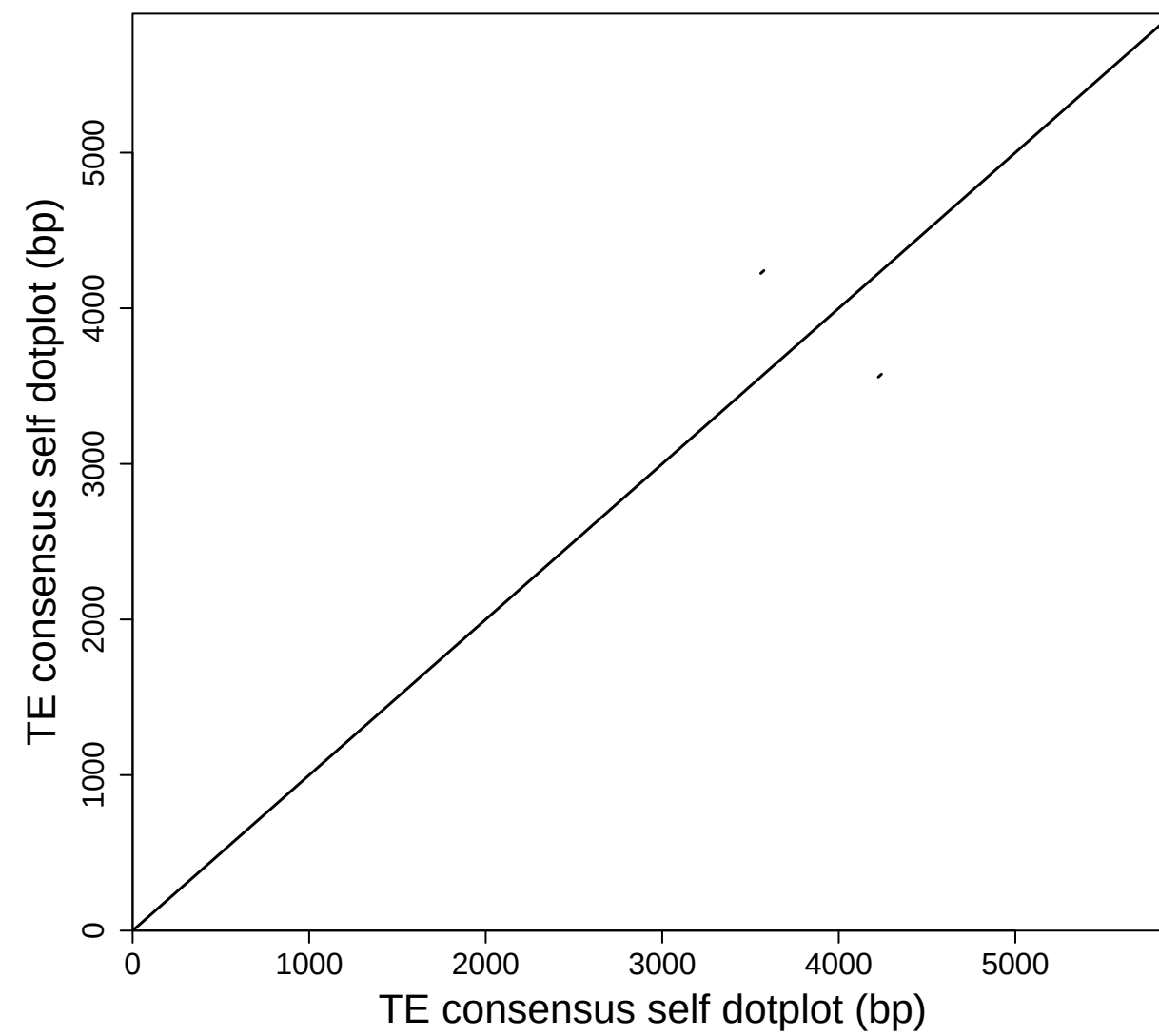
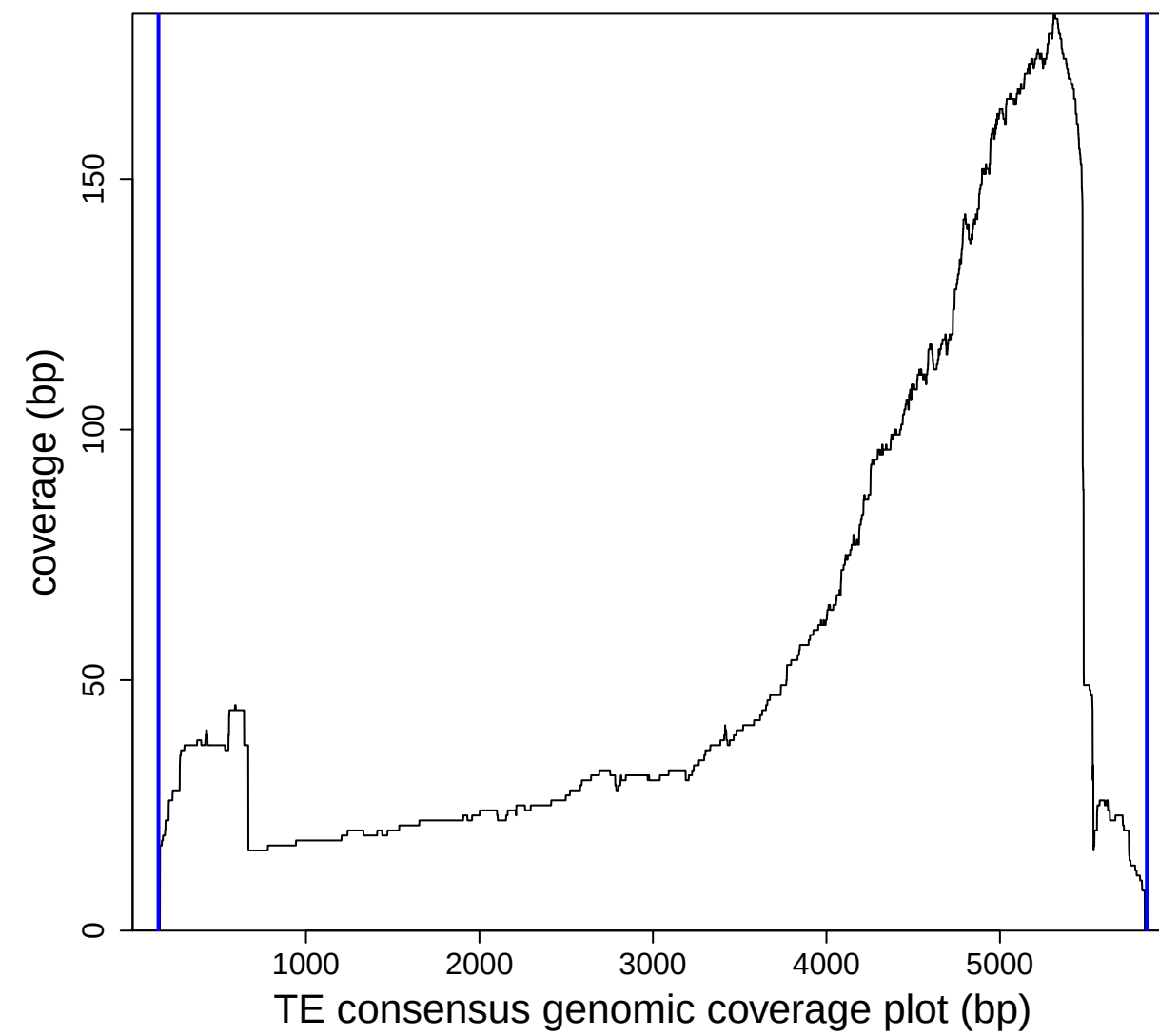
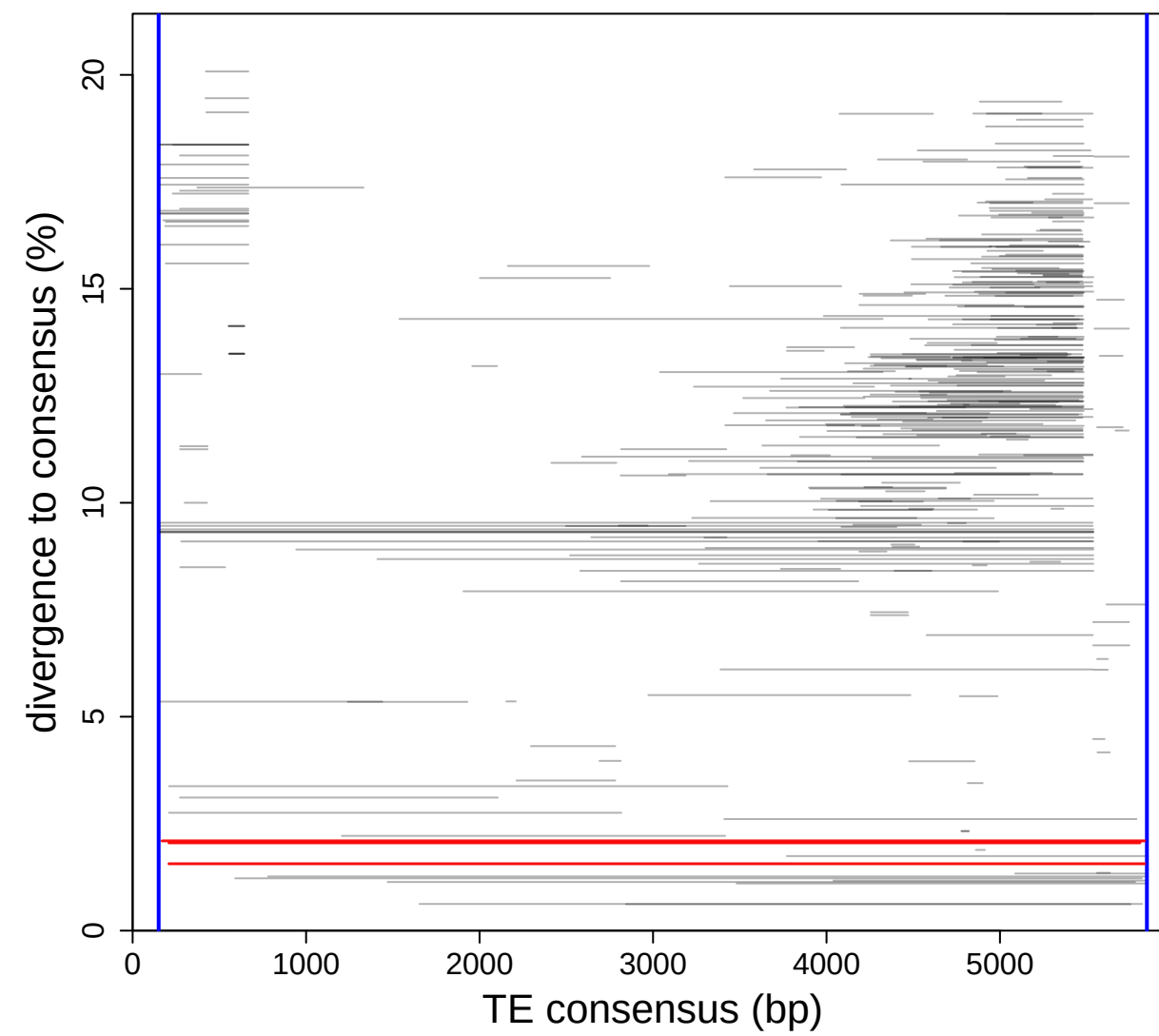
1.bed_fm 1.bed 0 0 n.bed_g 1.bed_fm 2.bed 0 0 bcln.fa aln.f
size: 5672bp; fragments: 354; full length: 11 (>=5104.8bp)

After TEtrimmer 5672 bp

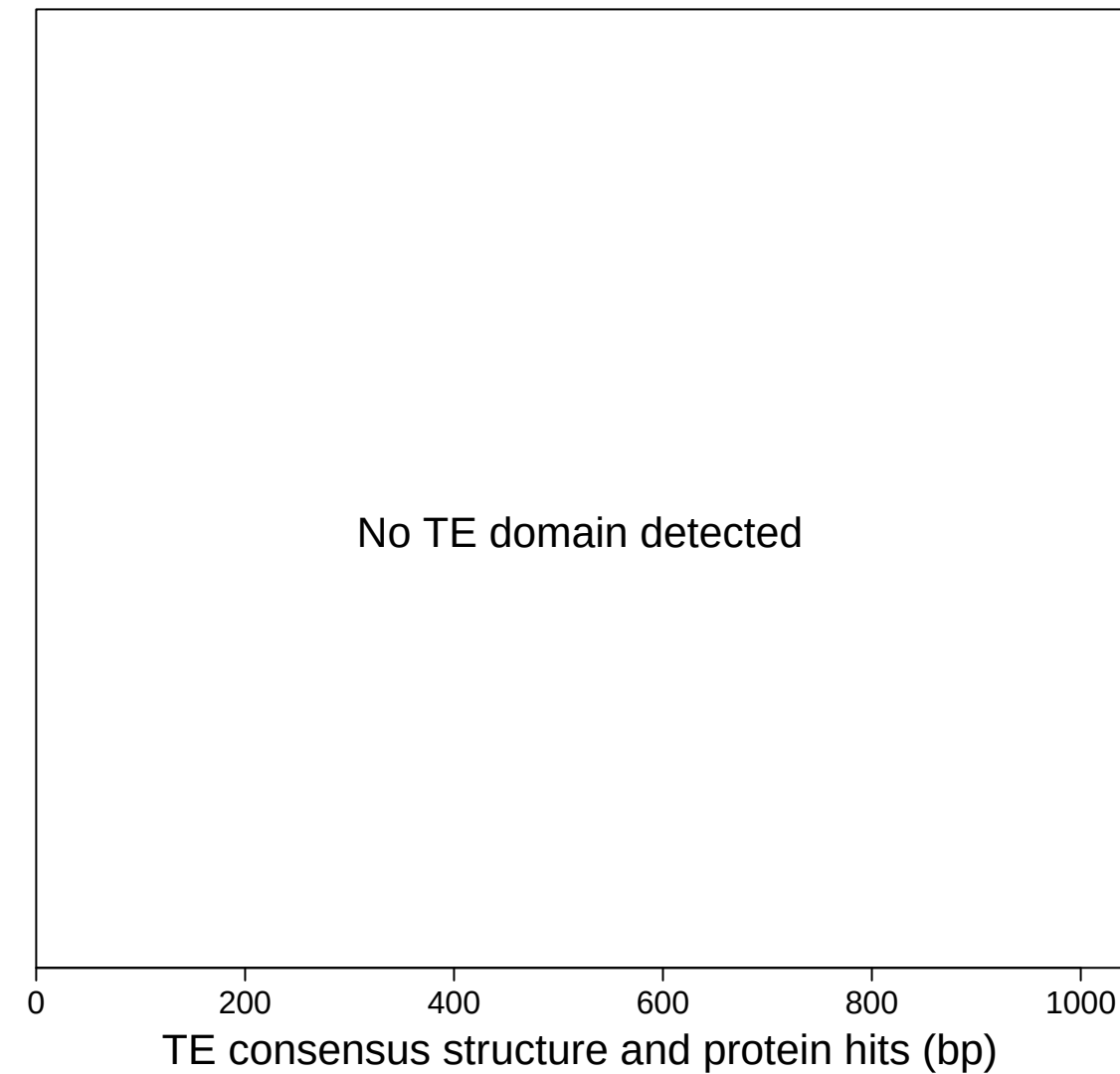
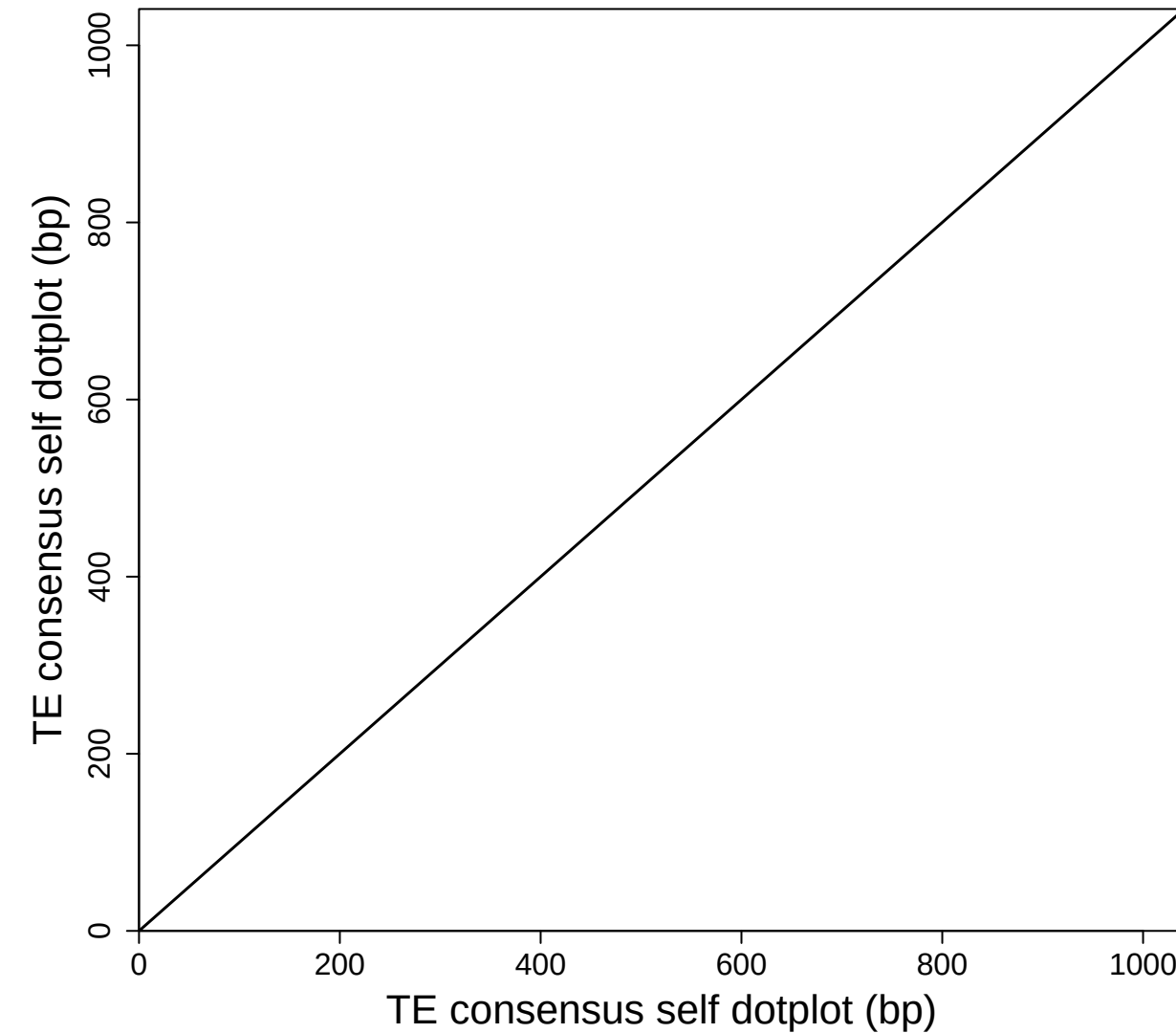
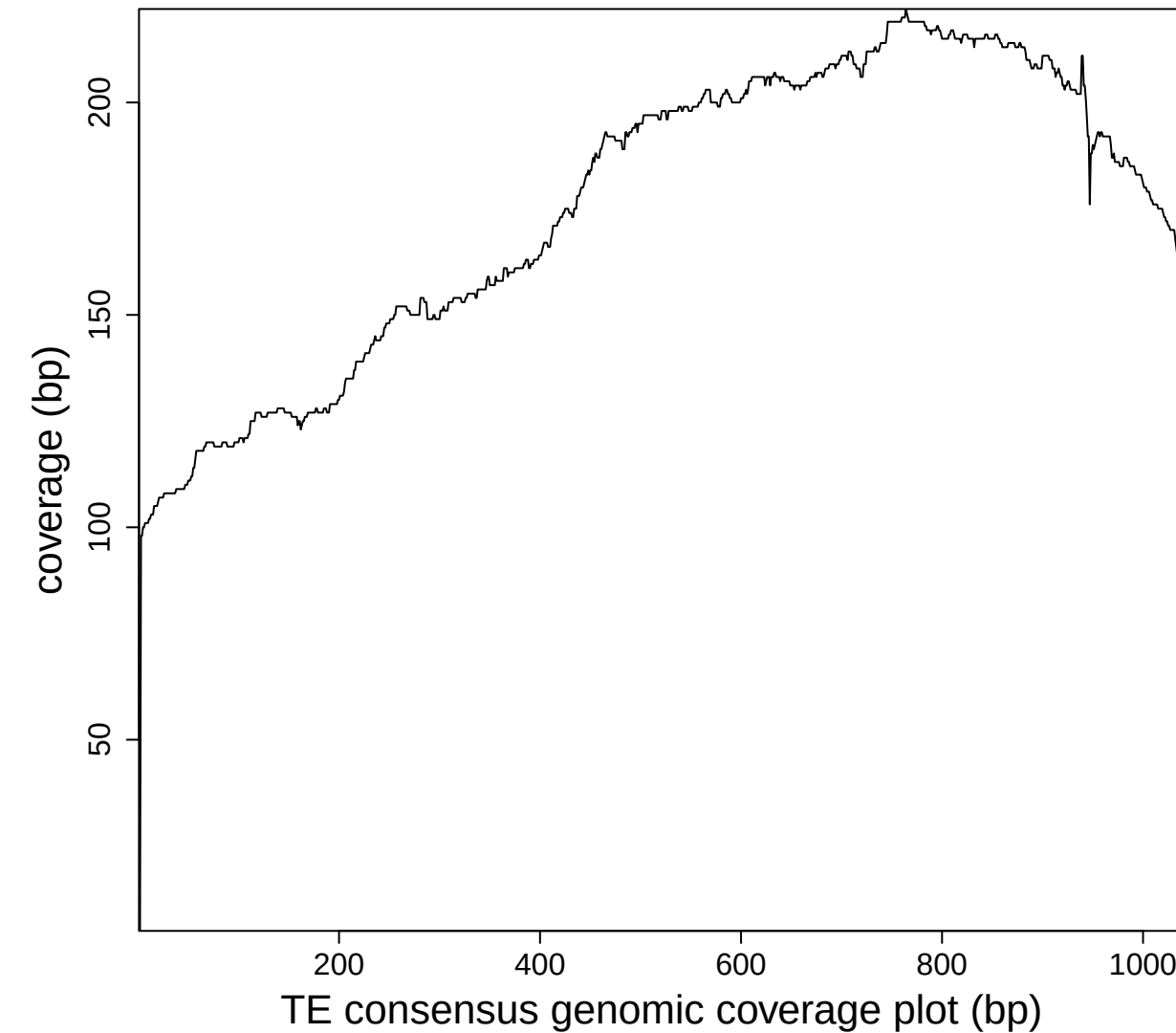
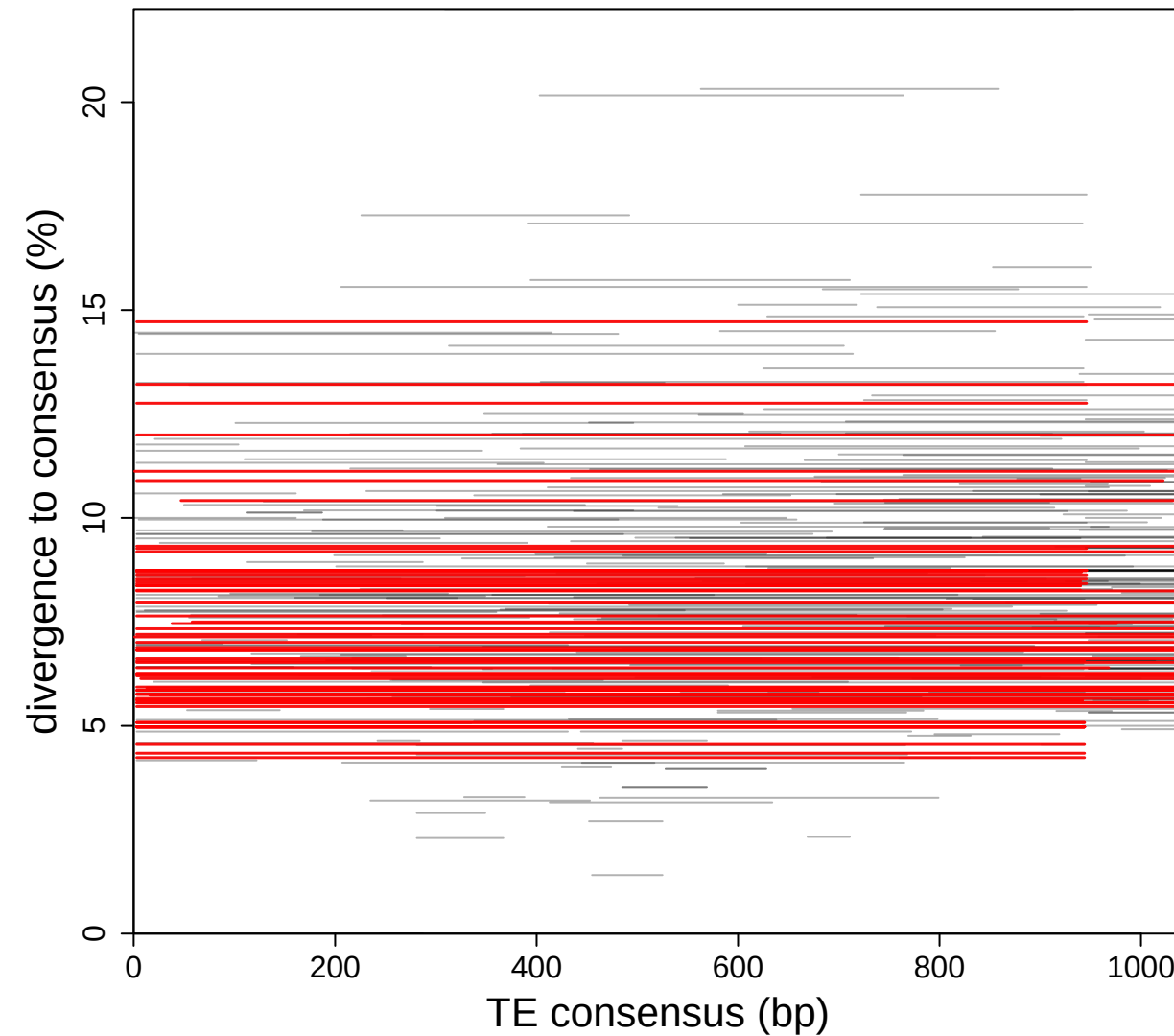


f.bed_g_2.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bc
size: 5997bp; fragments: 347; full length: 3 (>=5397.3bp)

After TEtrimmer Extended plot Blue lines are boundaries

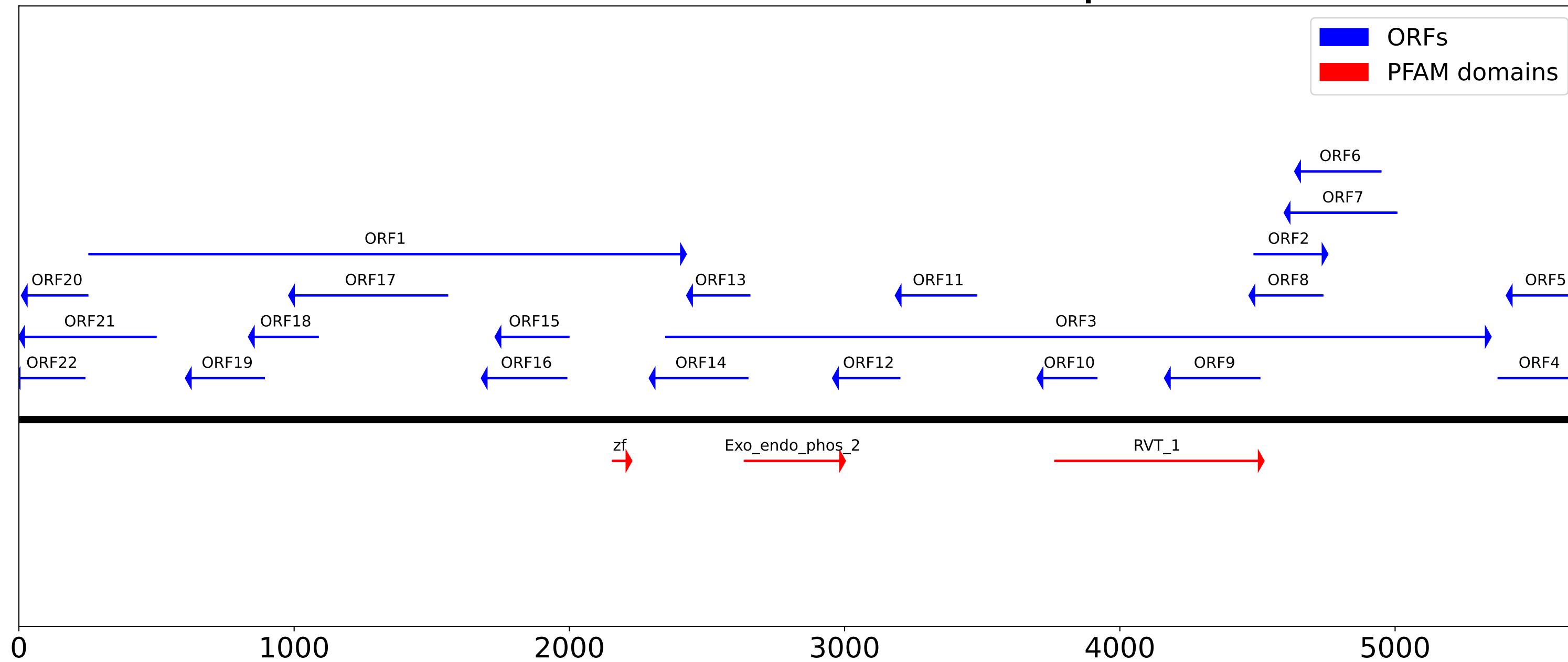


TE: rnd_5_family_7251
size: 1041bp; fragments: 388; full length: 72 (≥ 936.9 bp)

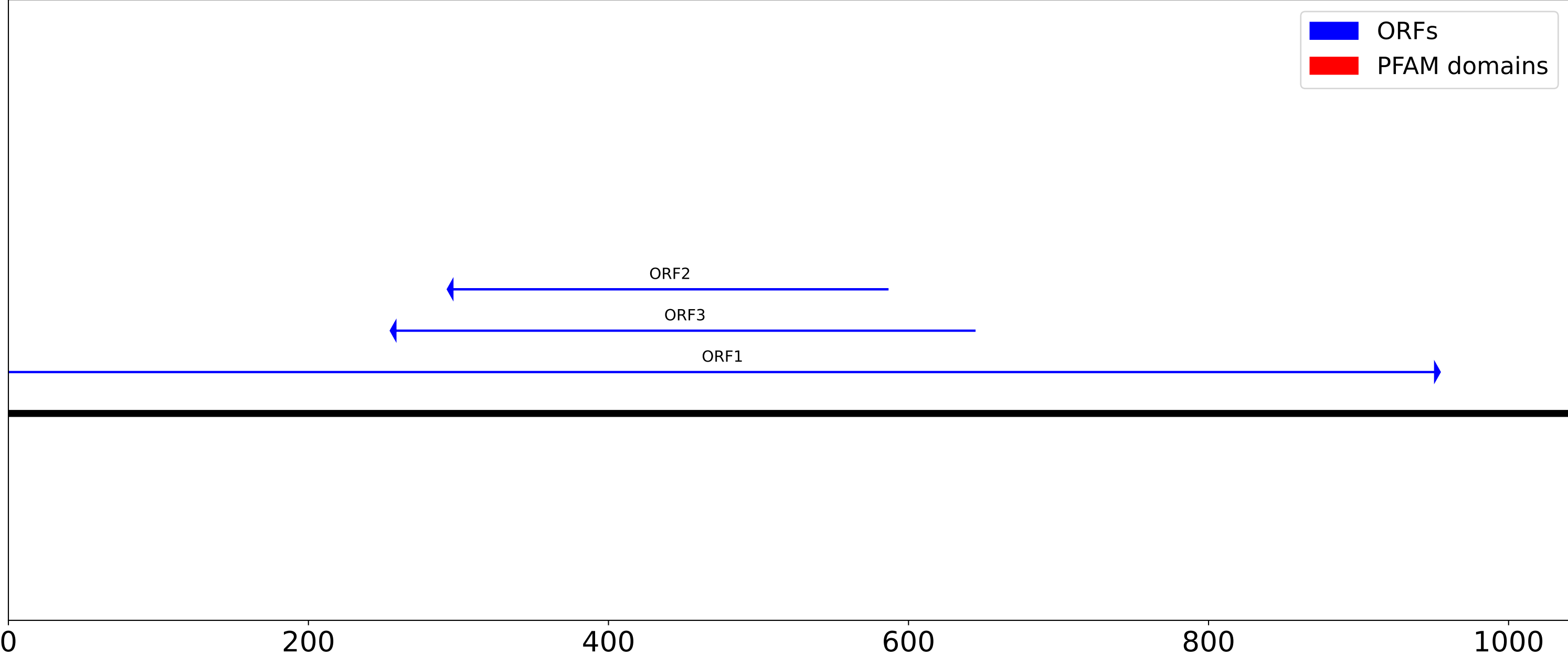


Before TEtrimmer 1041 bp

After TEtrimmer ORF and PFAM domain plot



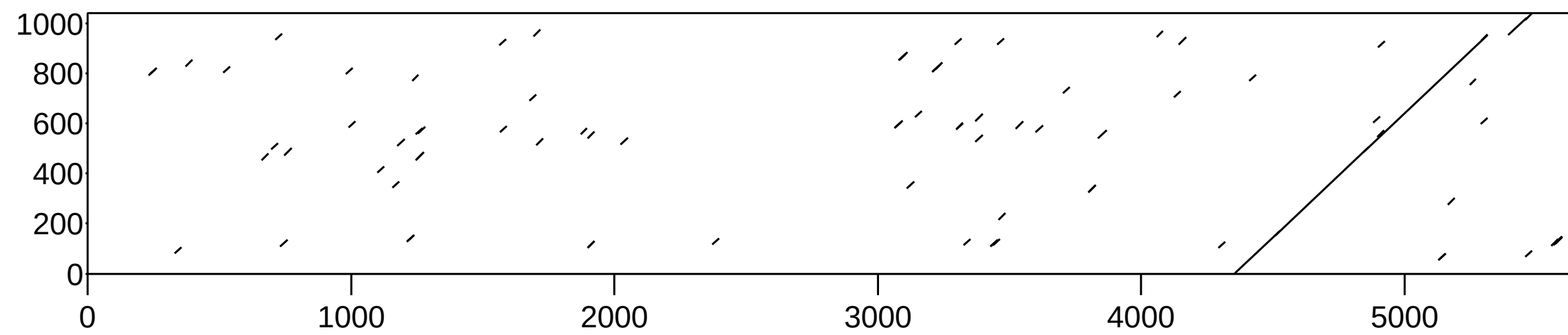
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

nsensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)