

Start crop Point

End crop Point

0

10

0

500

1000

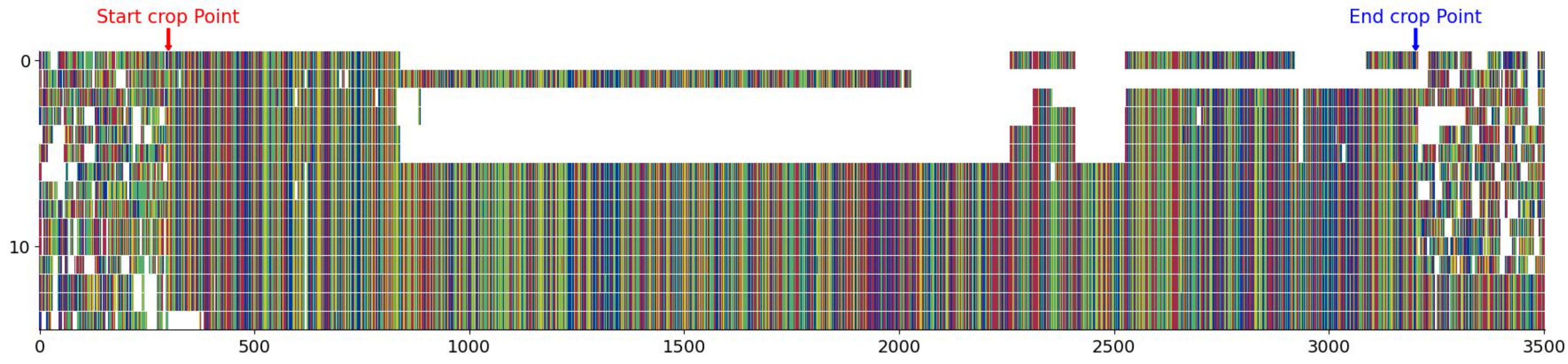
1500

2000

2500

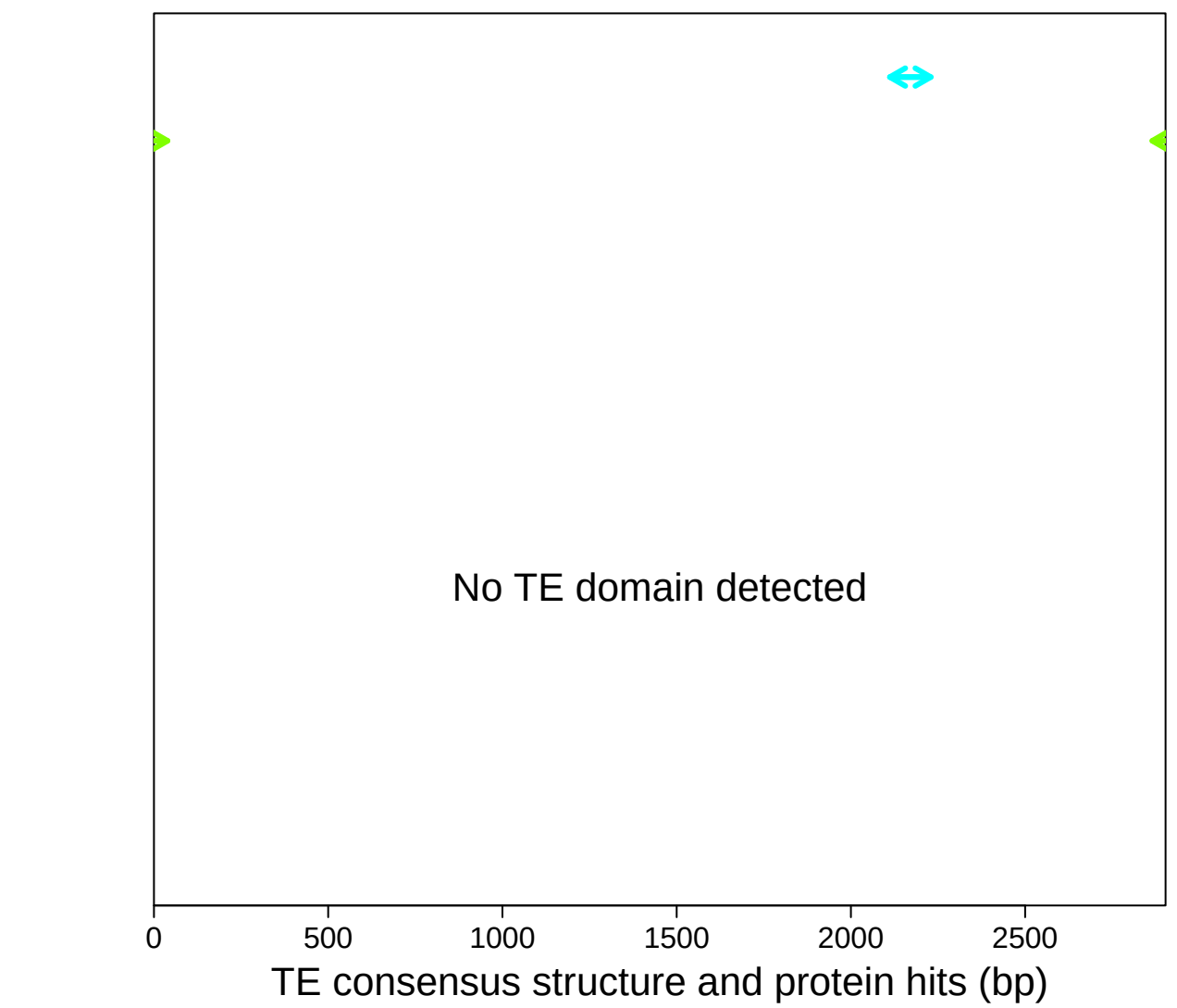
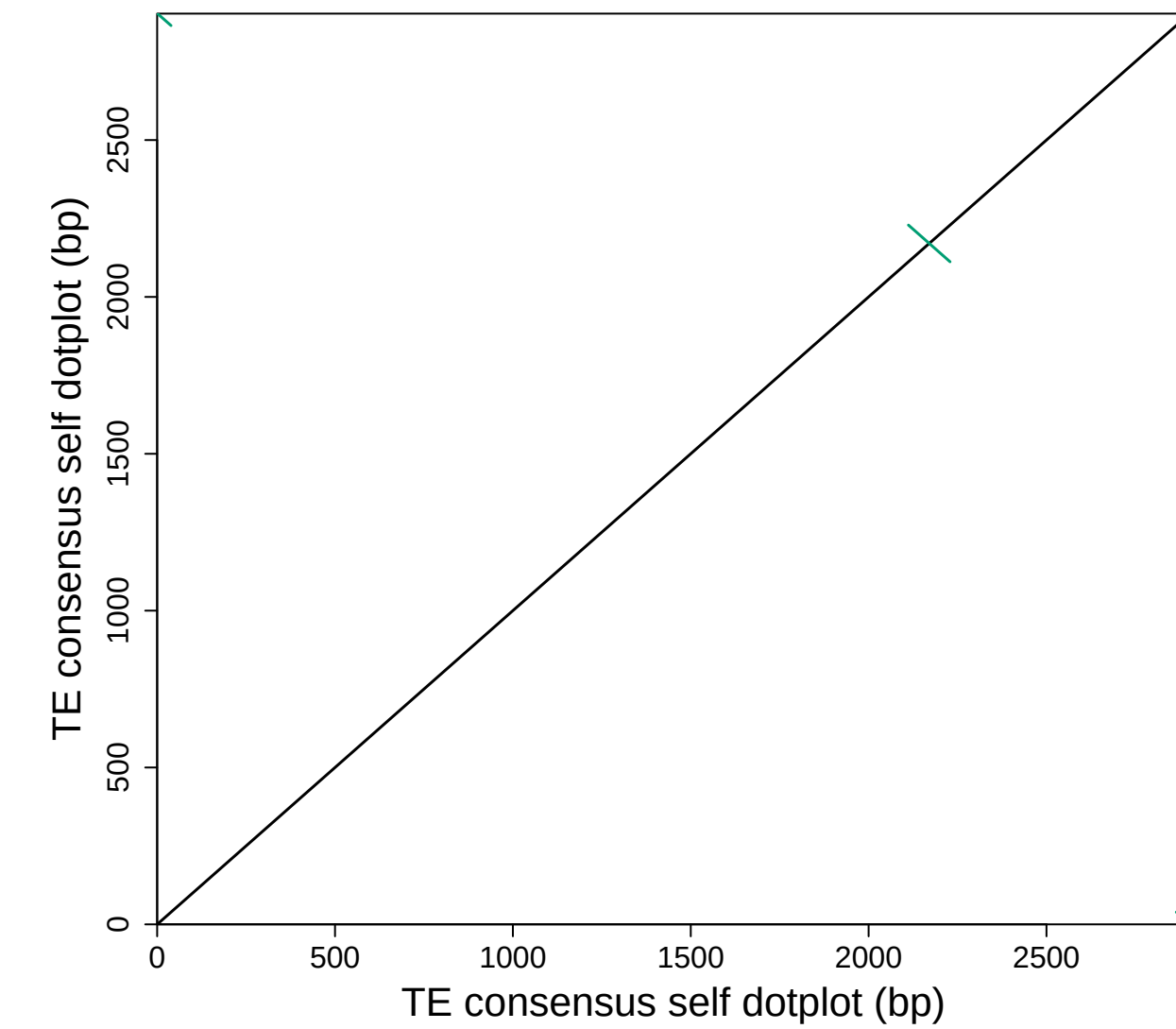
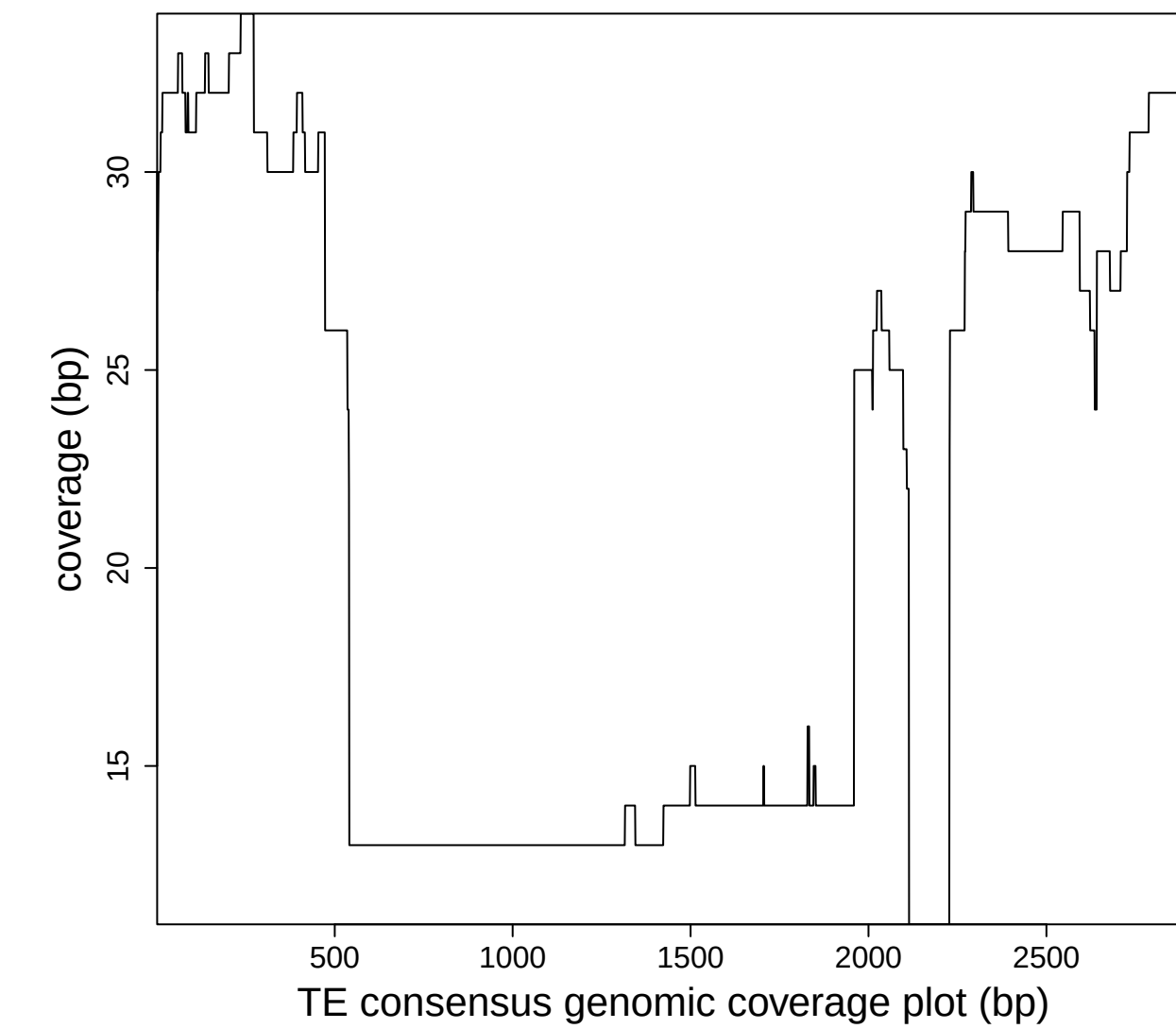
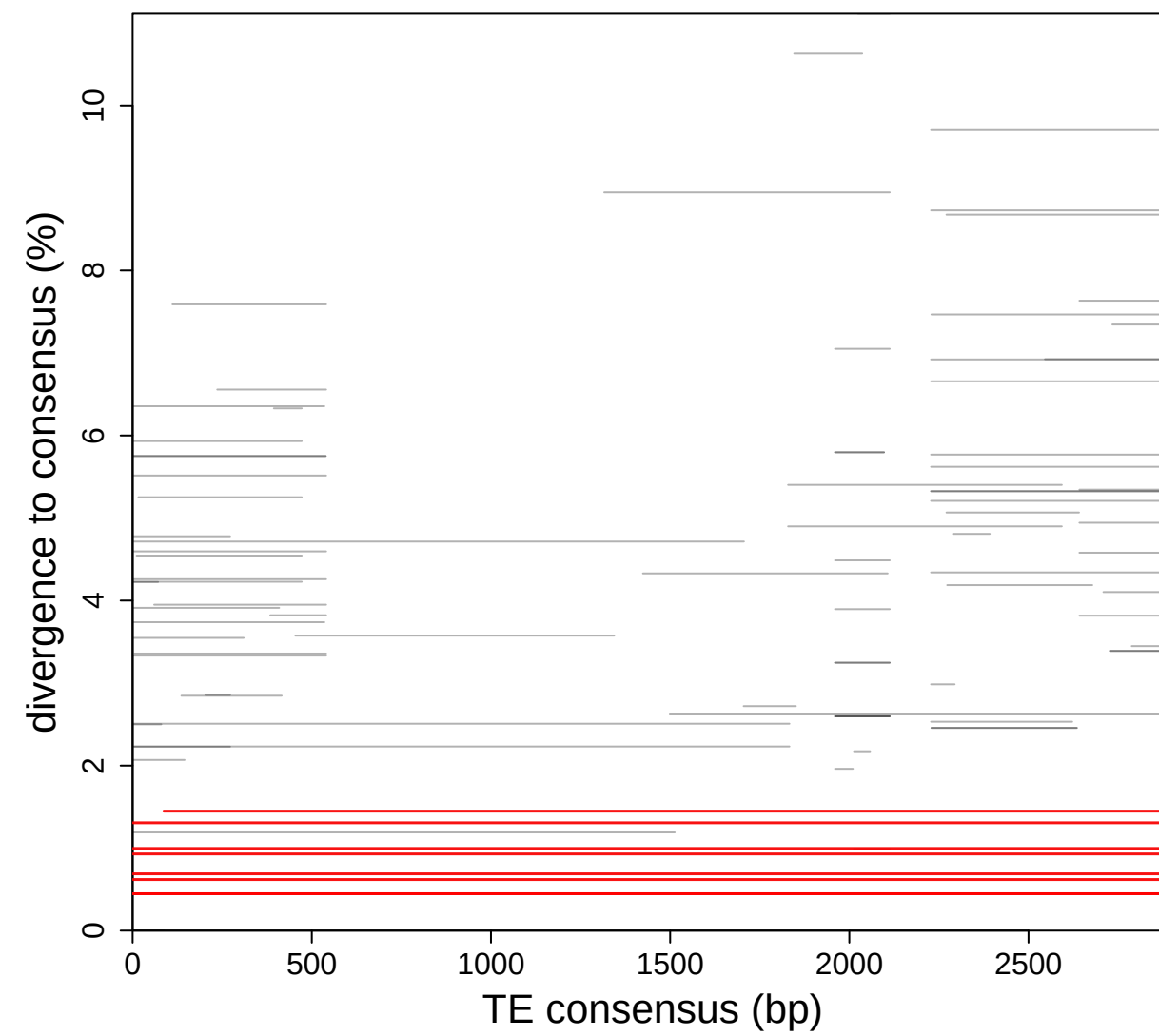
3000

3500



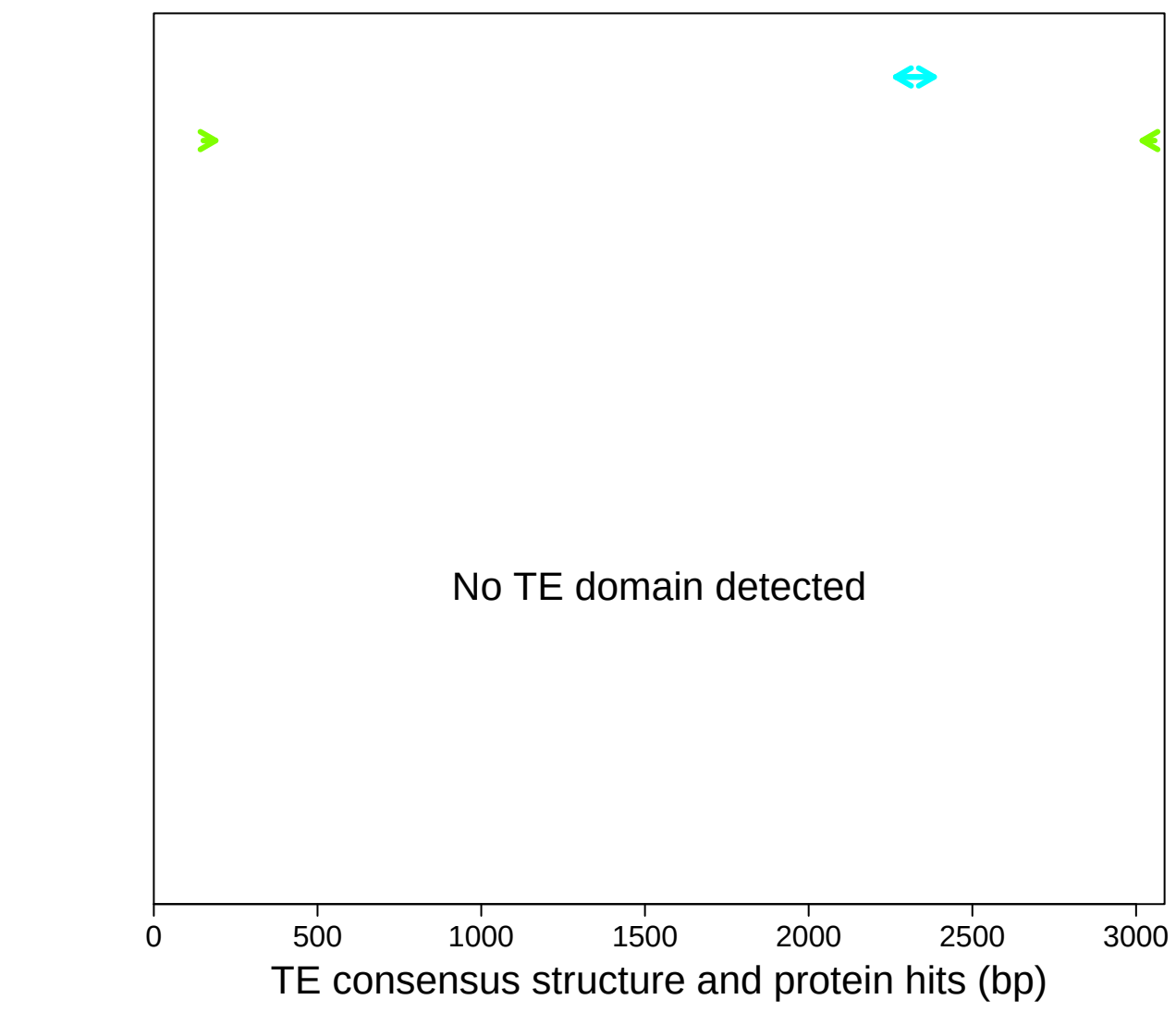
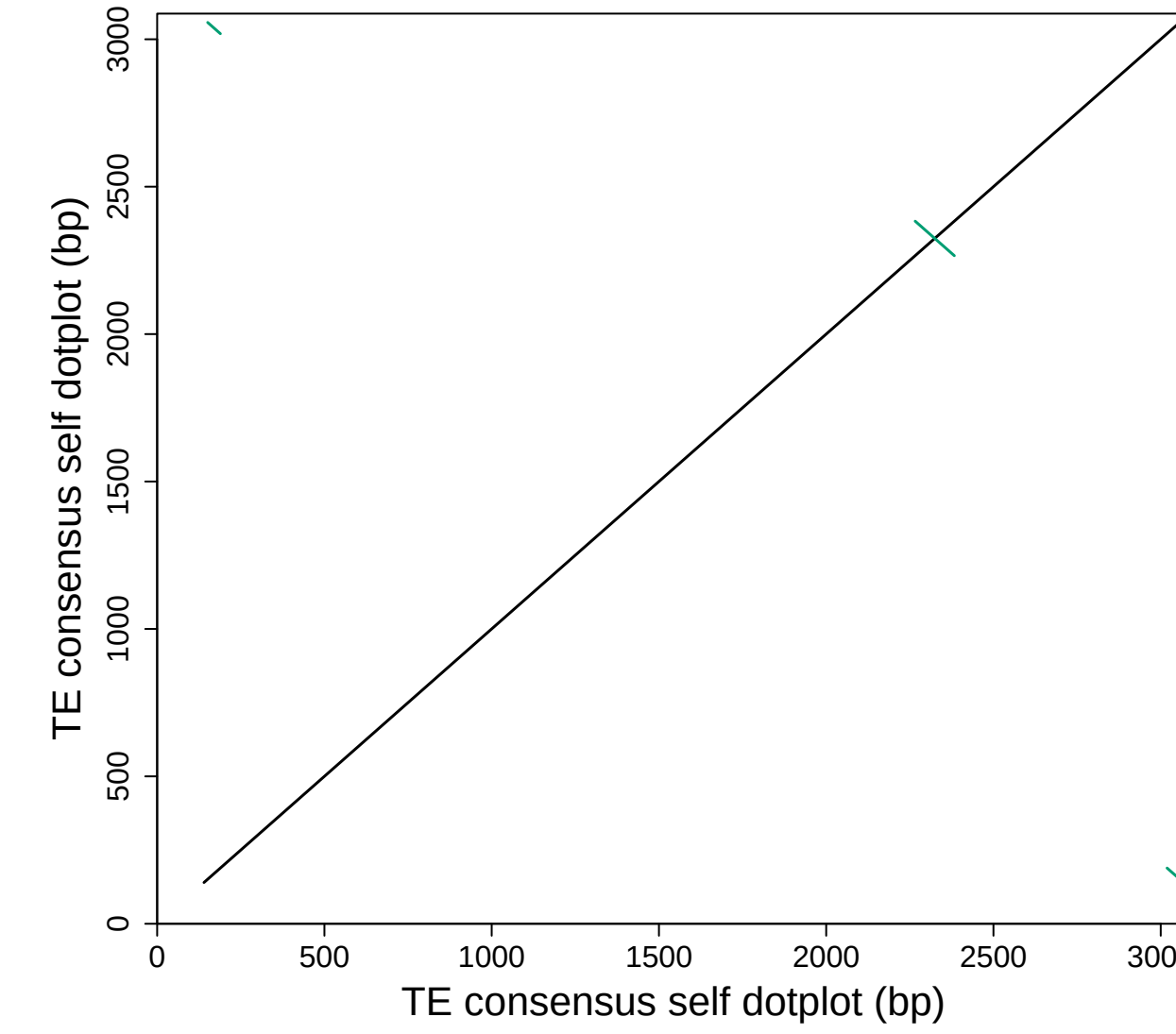
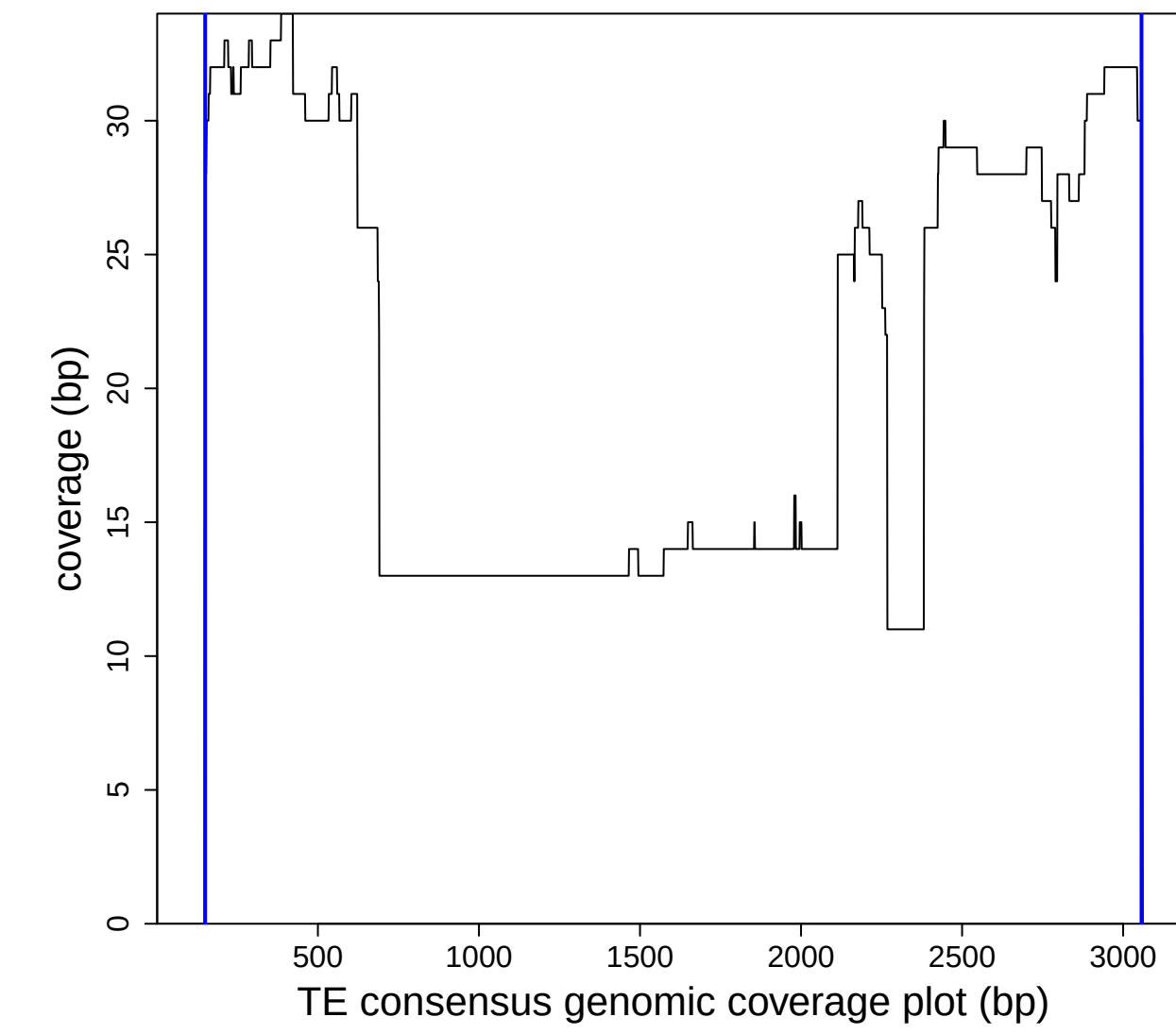
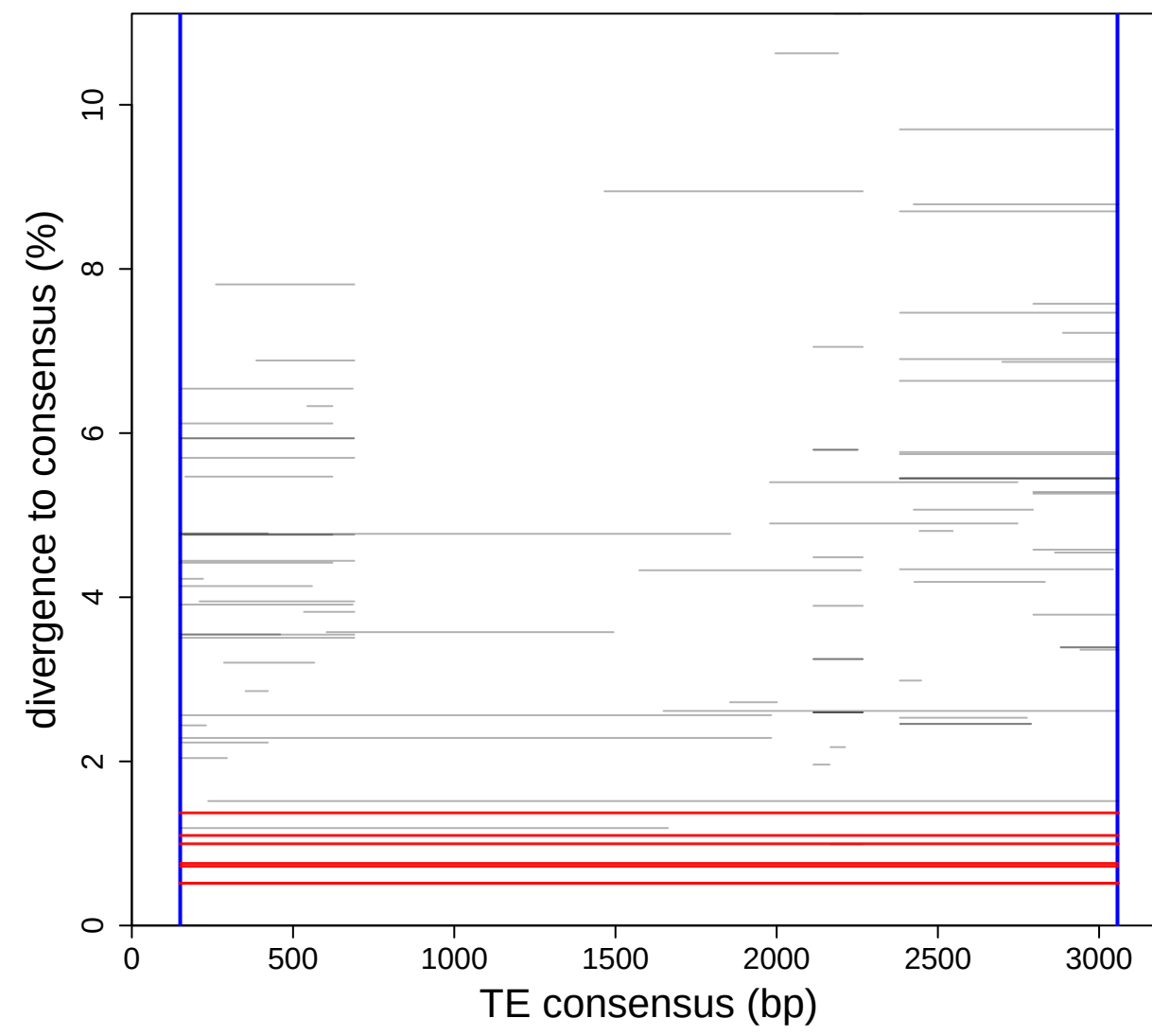
g_2.bed fm_1.bed 0 0 n.bed g_1.bed fm_2.bed 0 0 bcln.fa_ε
size: 2903bp; fragments: 92; full length: 8 (>=2612.7bp)

After TEtrimmer 2903 bp



f.bed_g_2.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bc
size: 3207bp; fragments: 92; full length: 7 (>=2886.3bp)

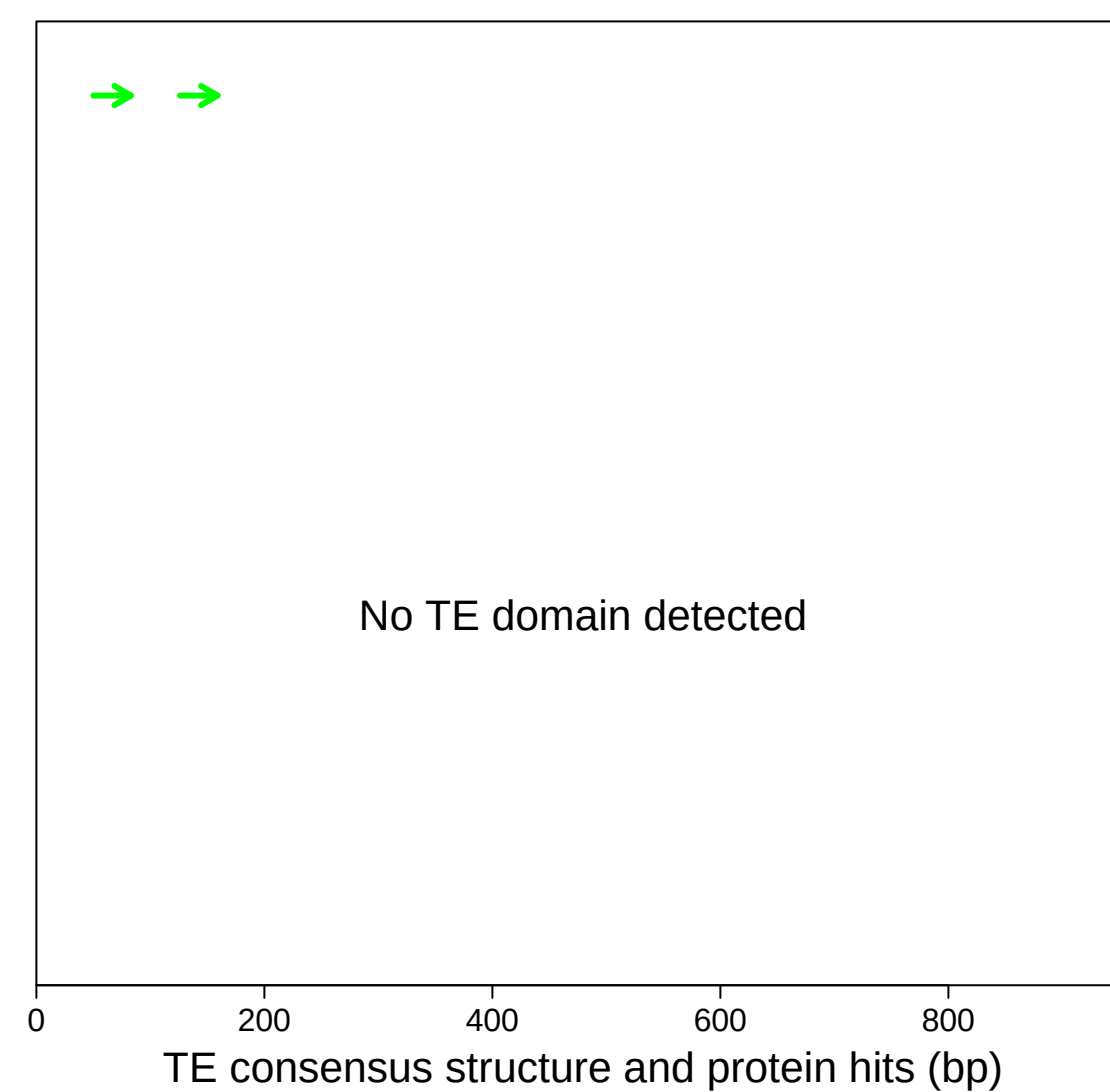
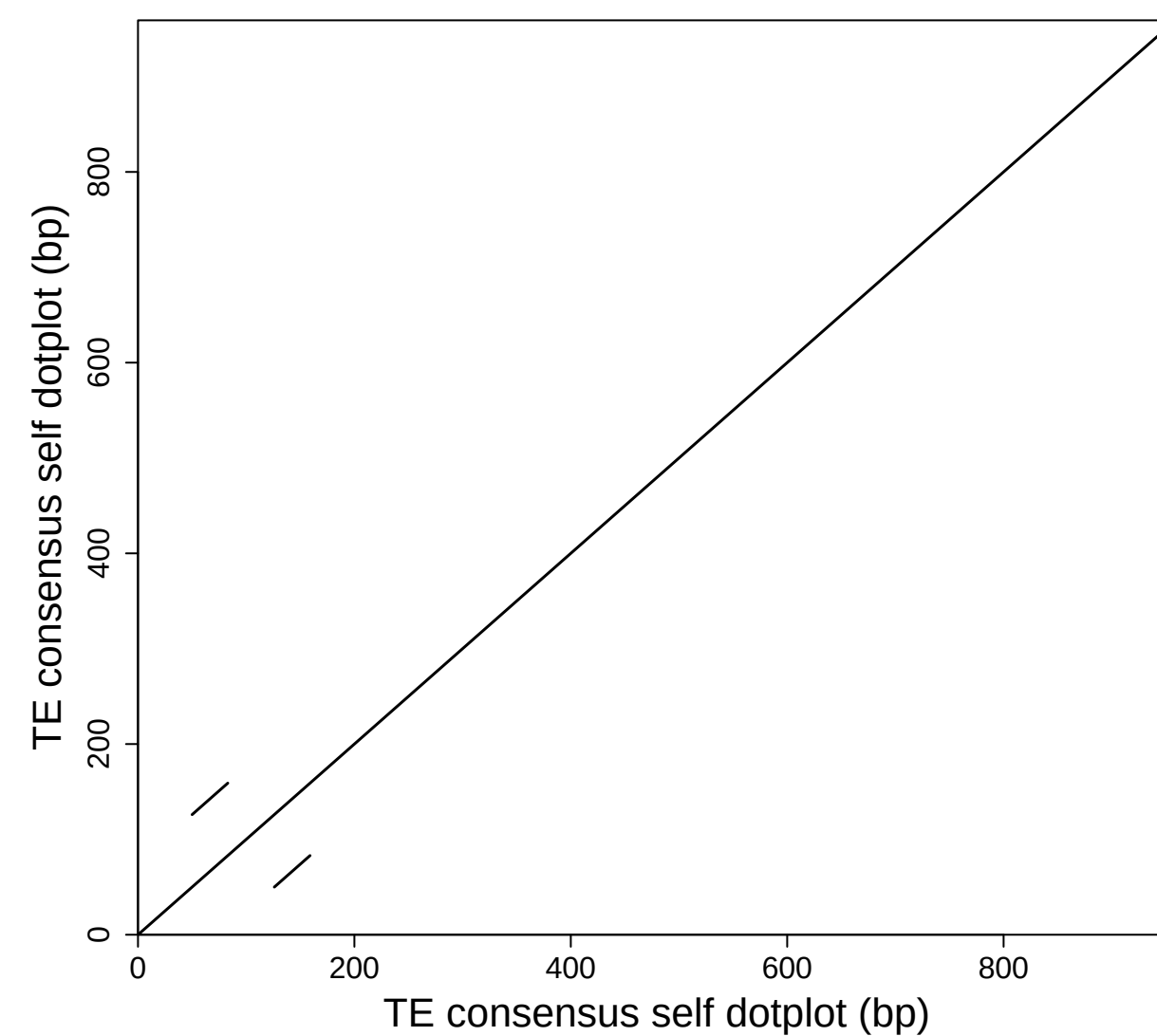
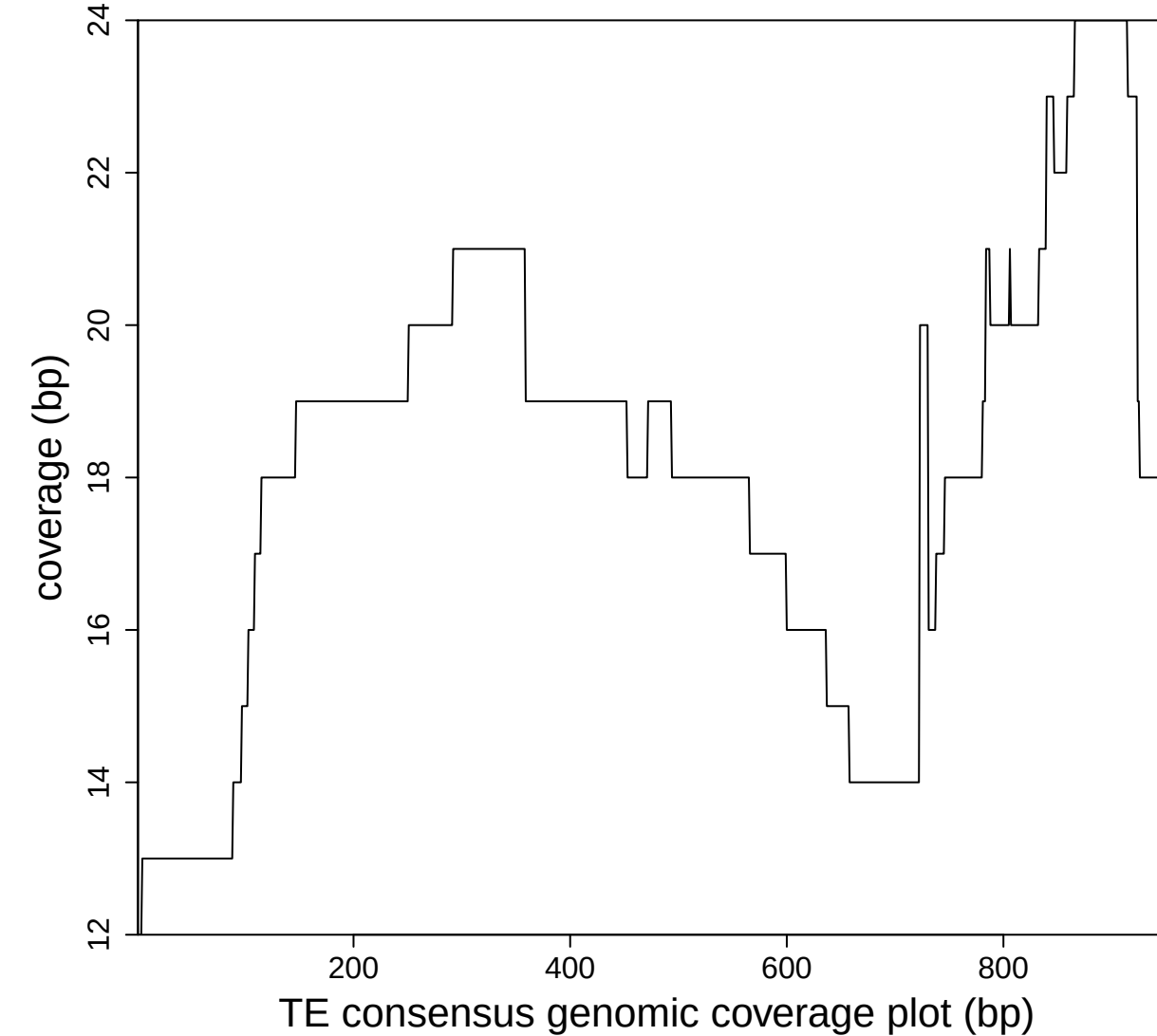
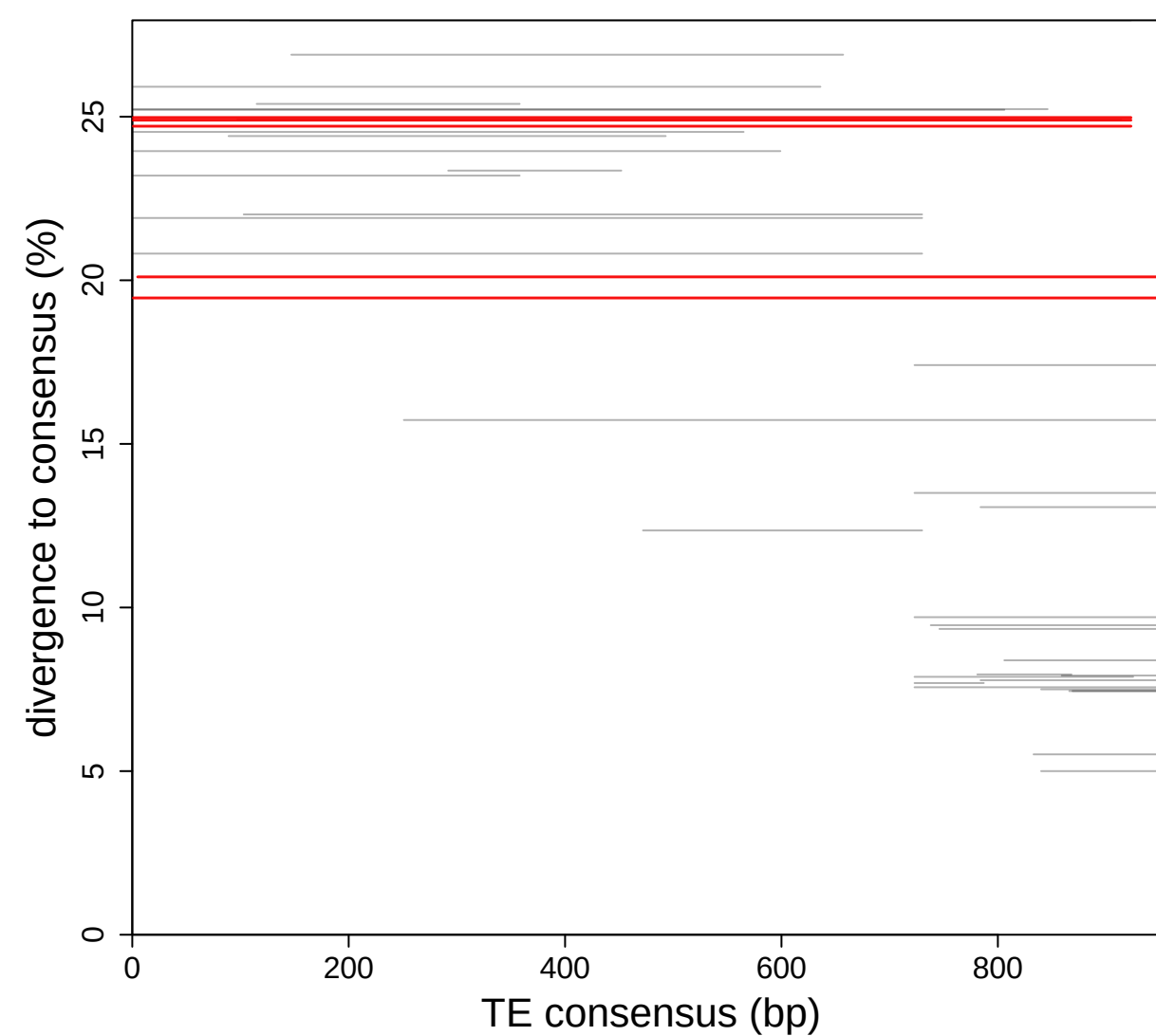
After TEtrimmer Extended plot Blue lines are boundaries



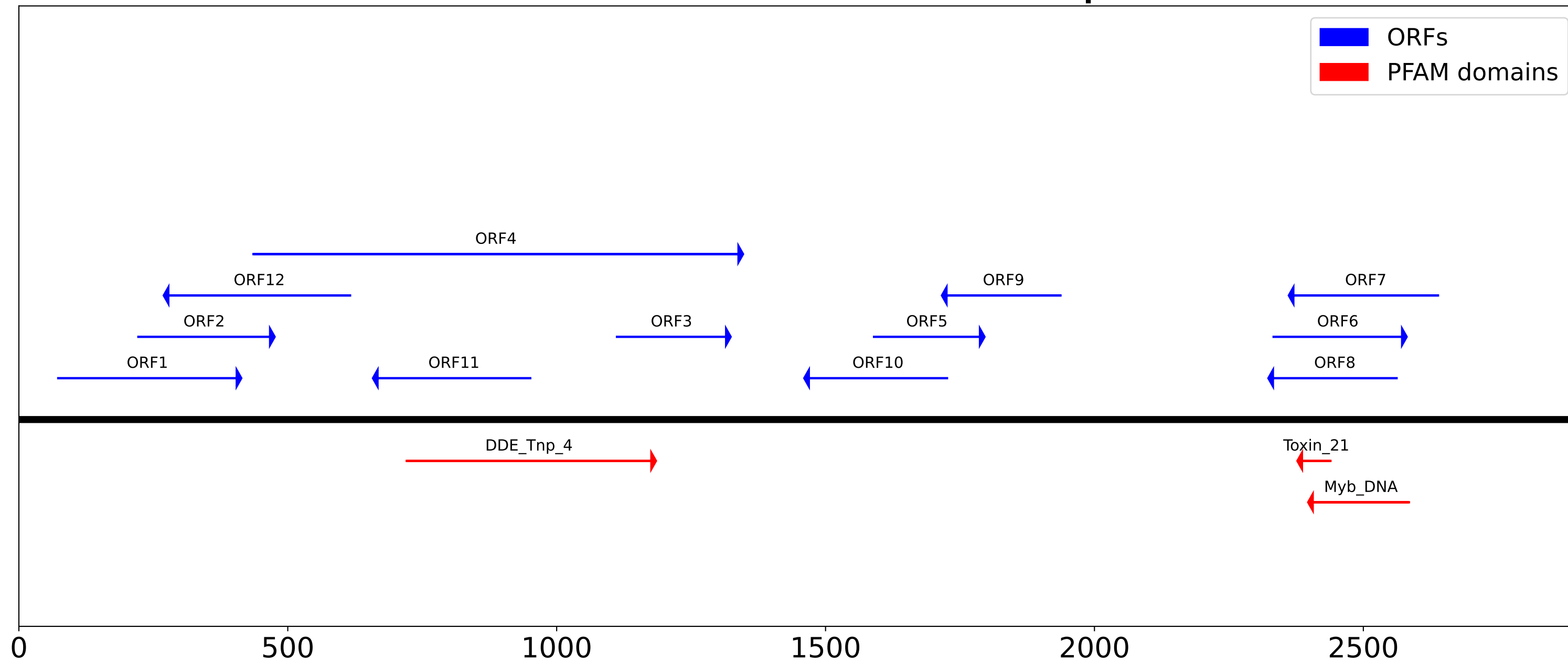
Before TEtrimmer 959 bp

TE: rnd_5_family_6478

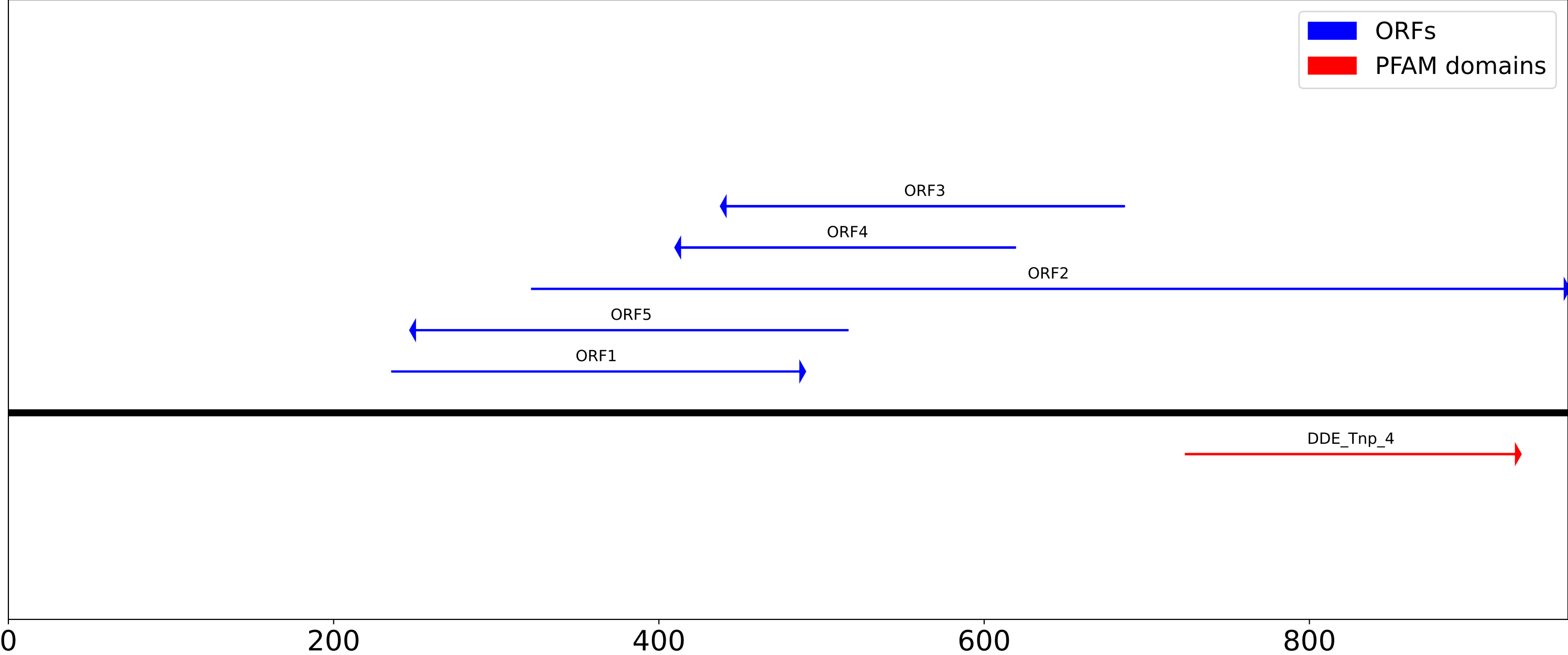
size: 959bp; fragments: 40; full length: 5 (≥ 863.1 bp)



After TEtrimmer ORF and PFAM domain plot



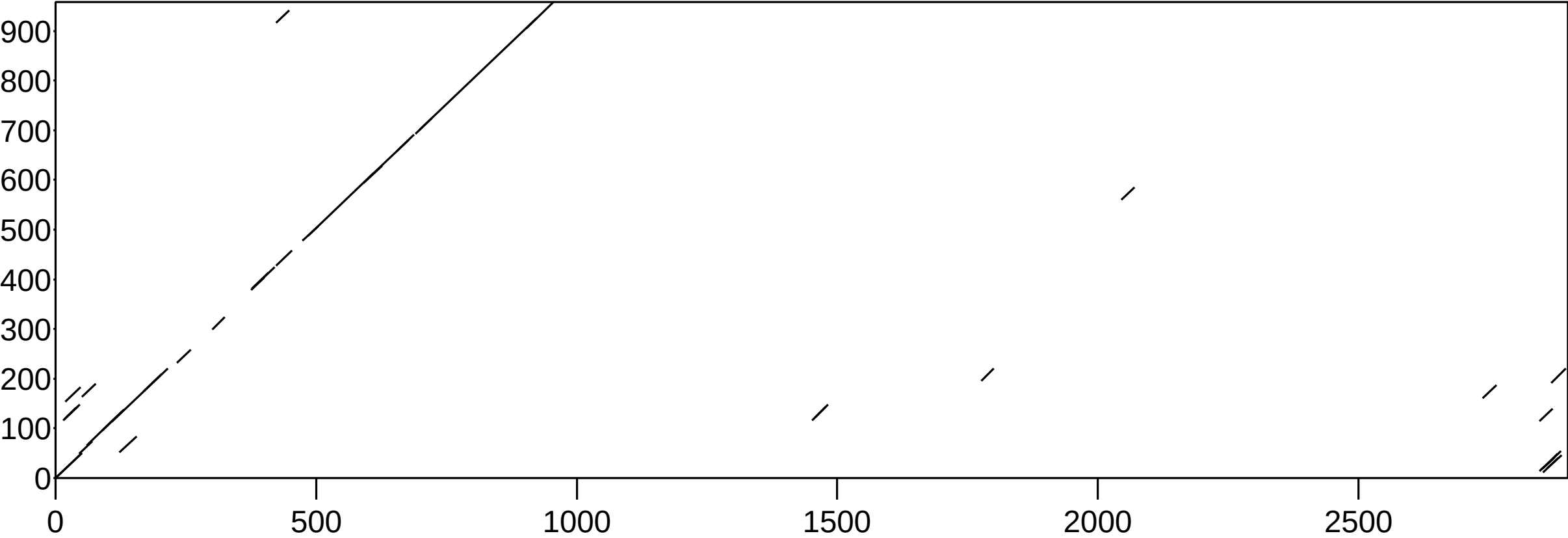
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)