

MSA length = 3952

Start crop Point

End crop Point



Start crop Point

End crop Point

0

10

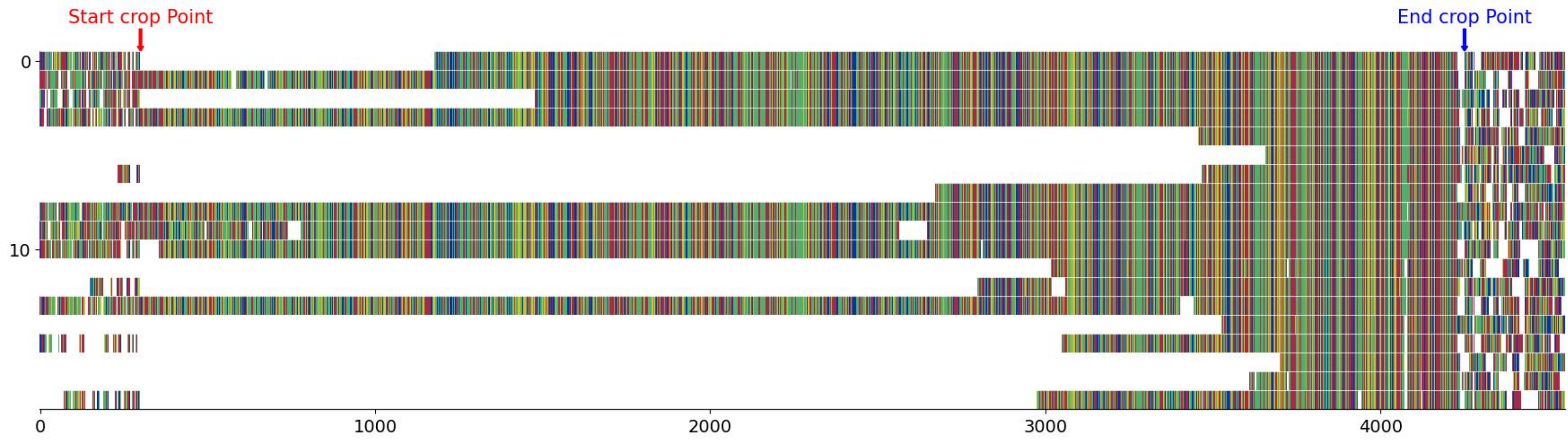
0

1000

2000

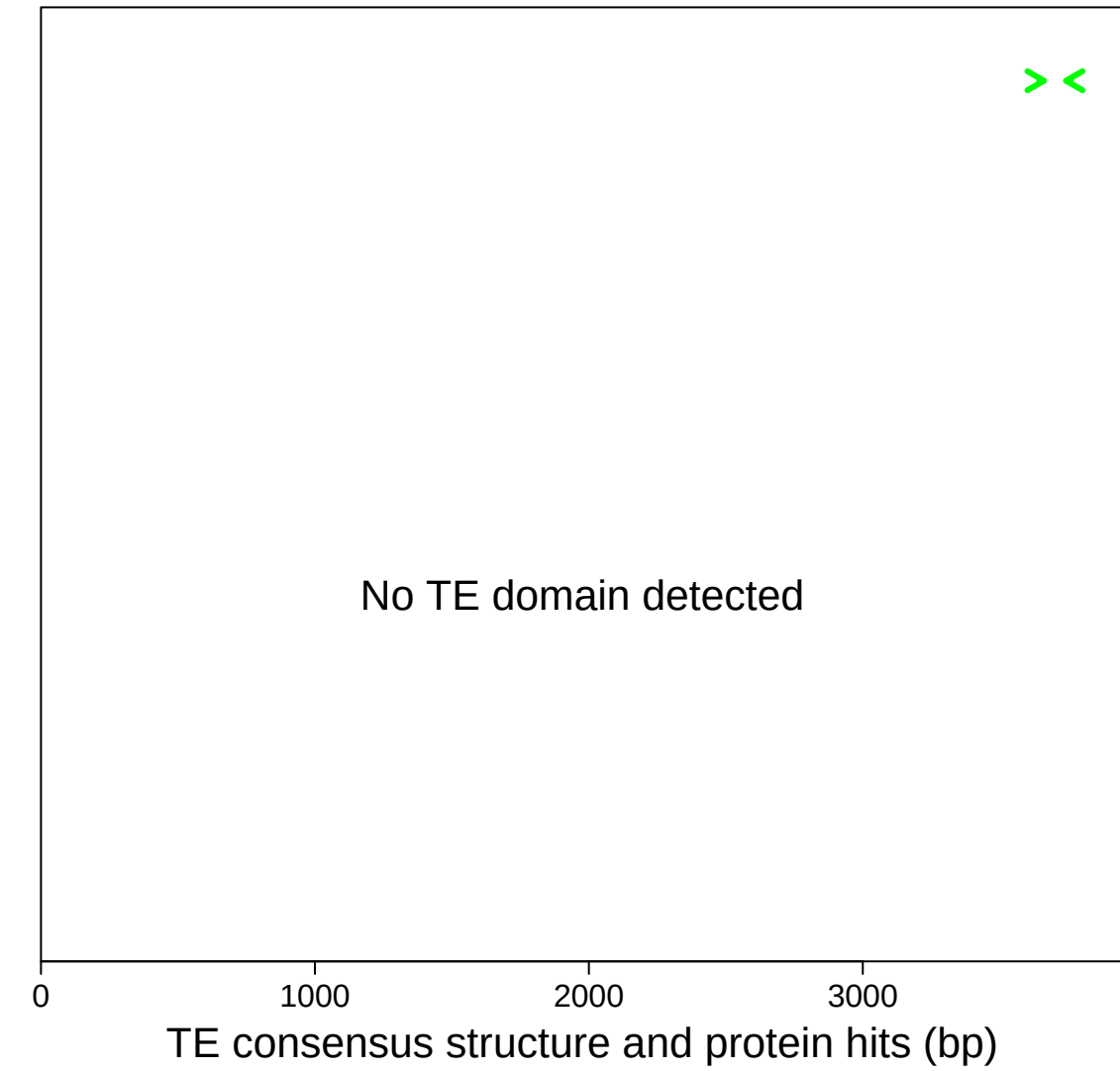
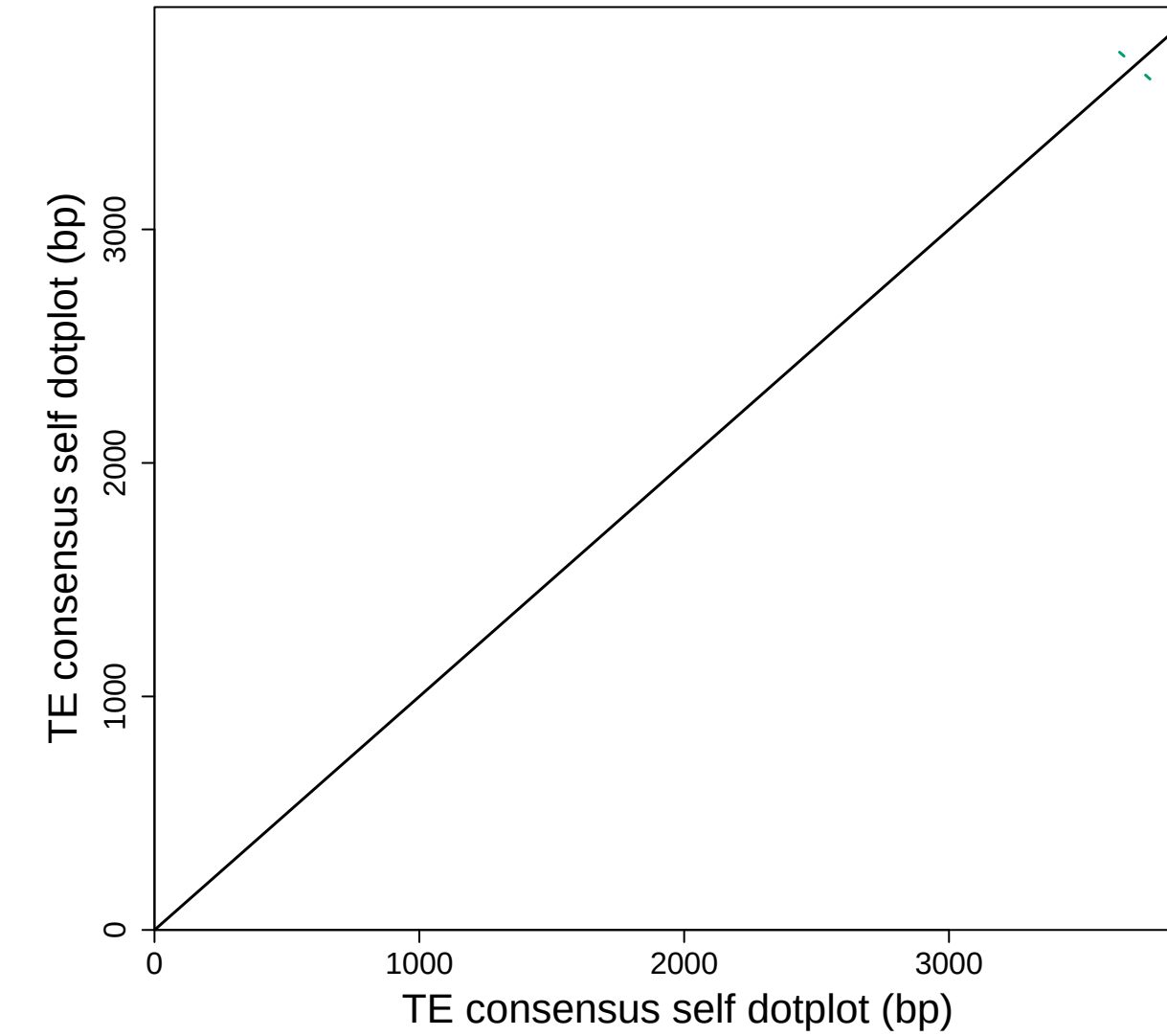
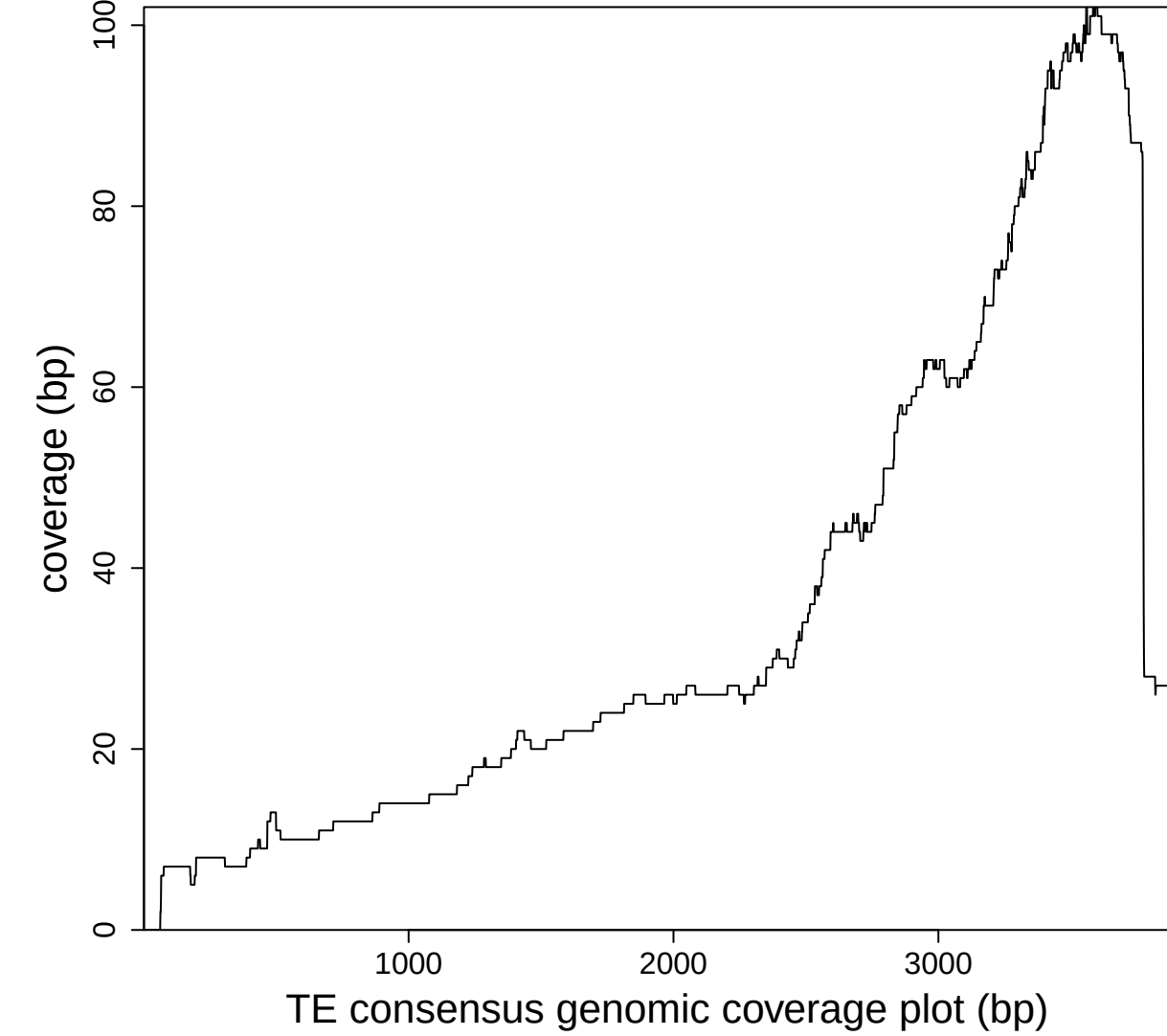
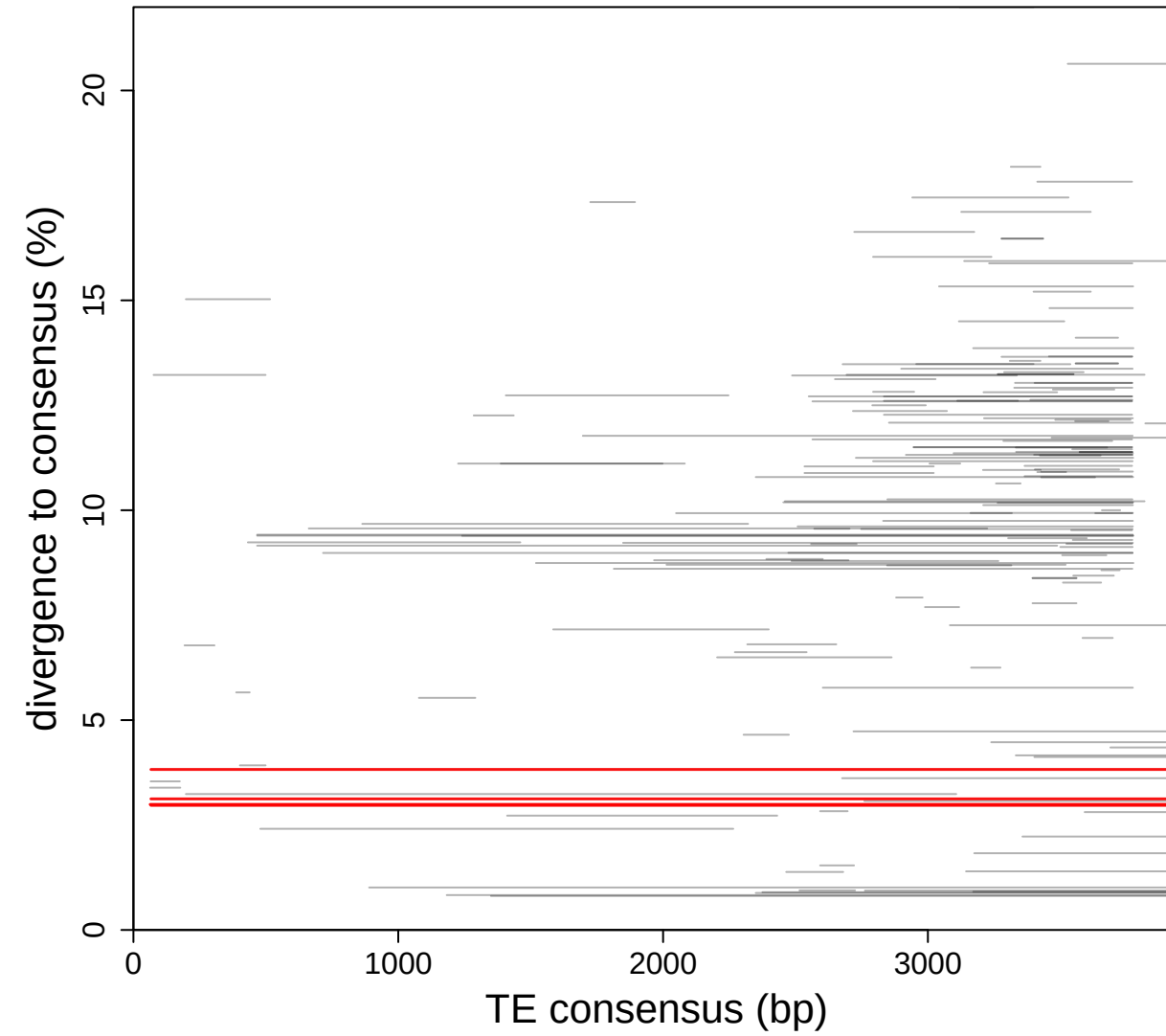
3000

4000



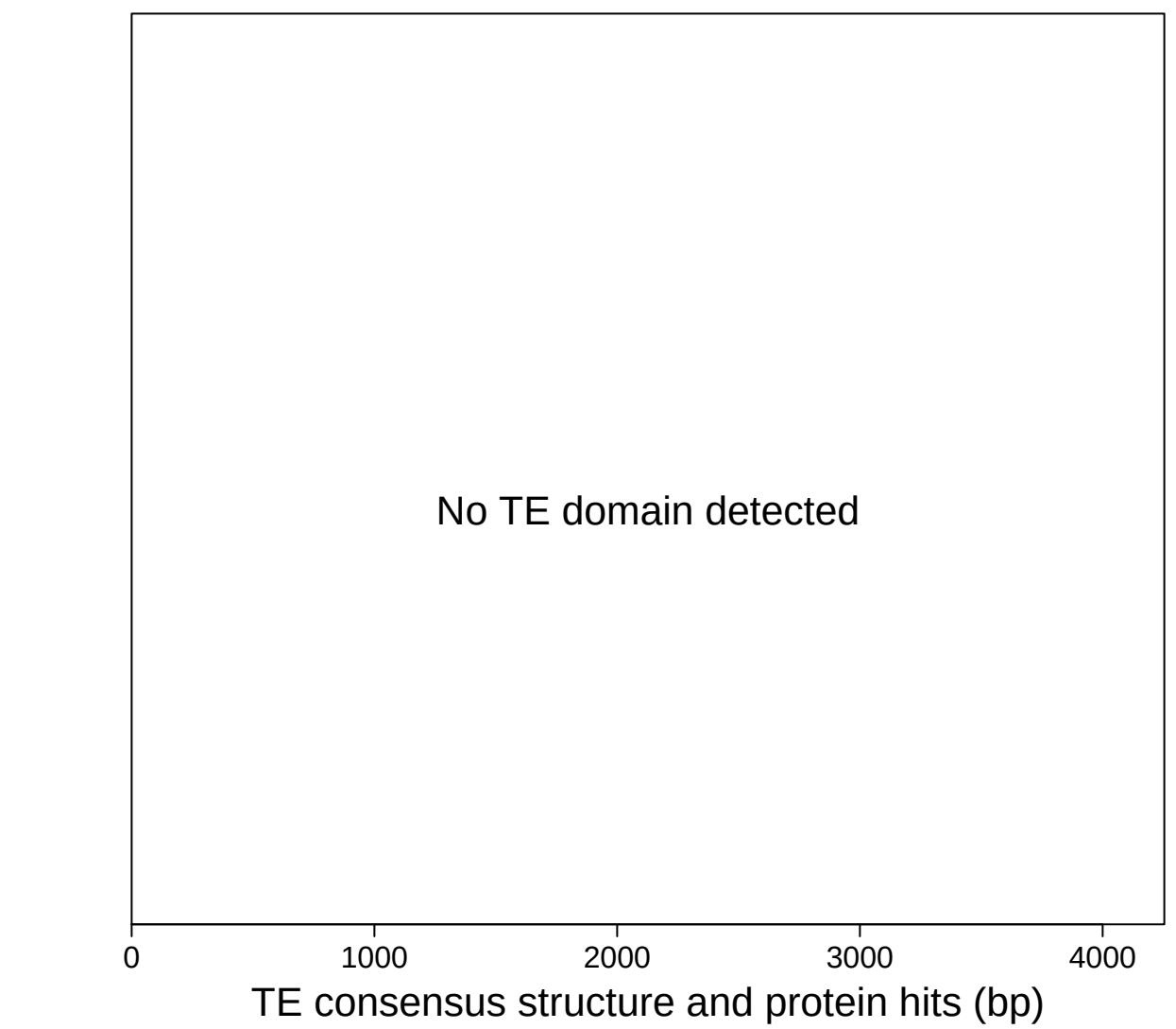
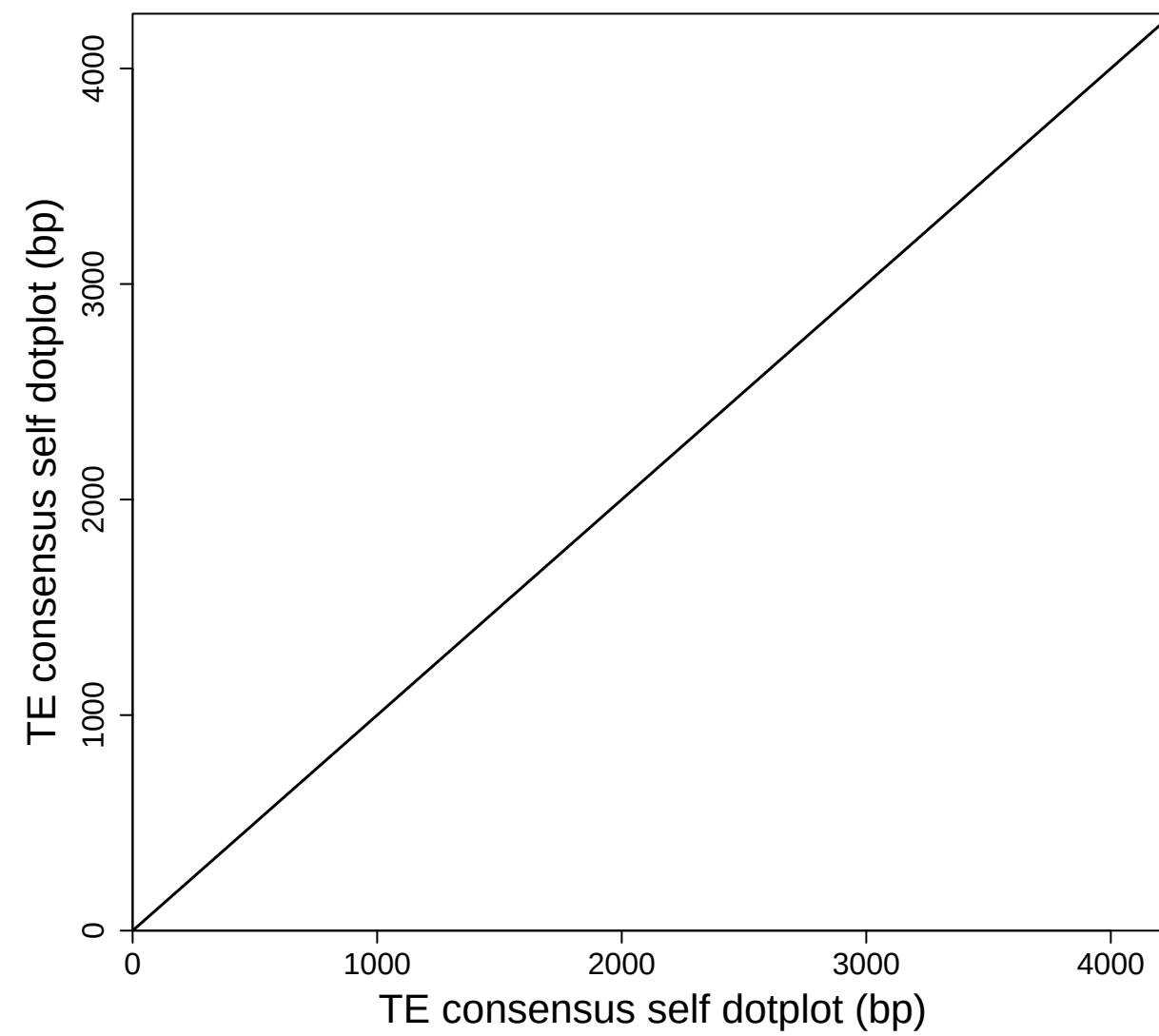
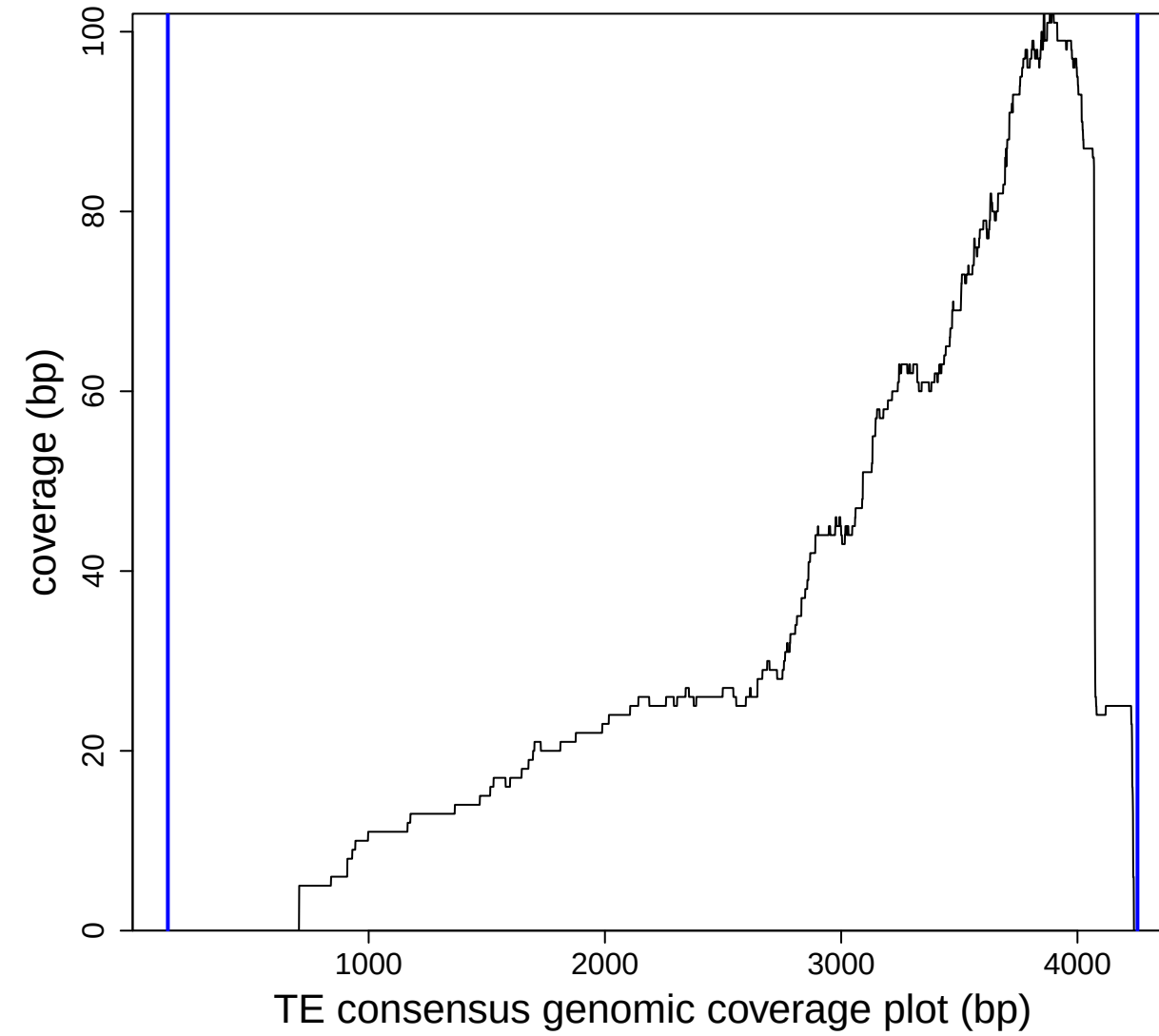
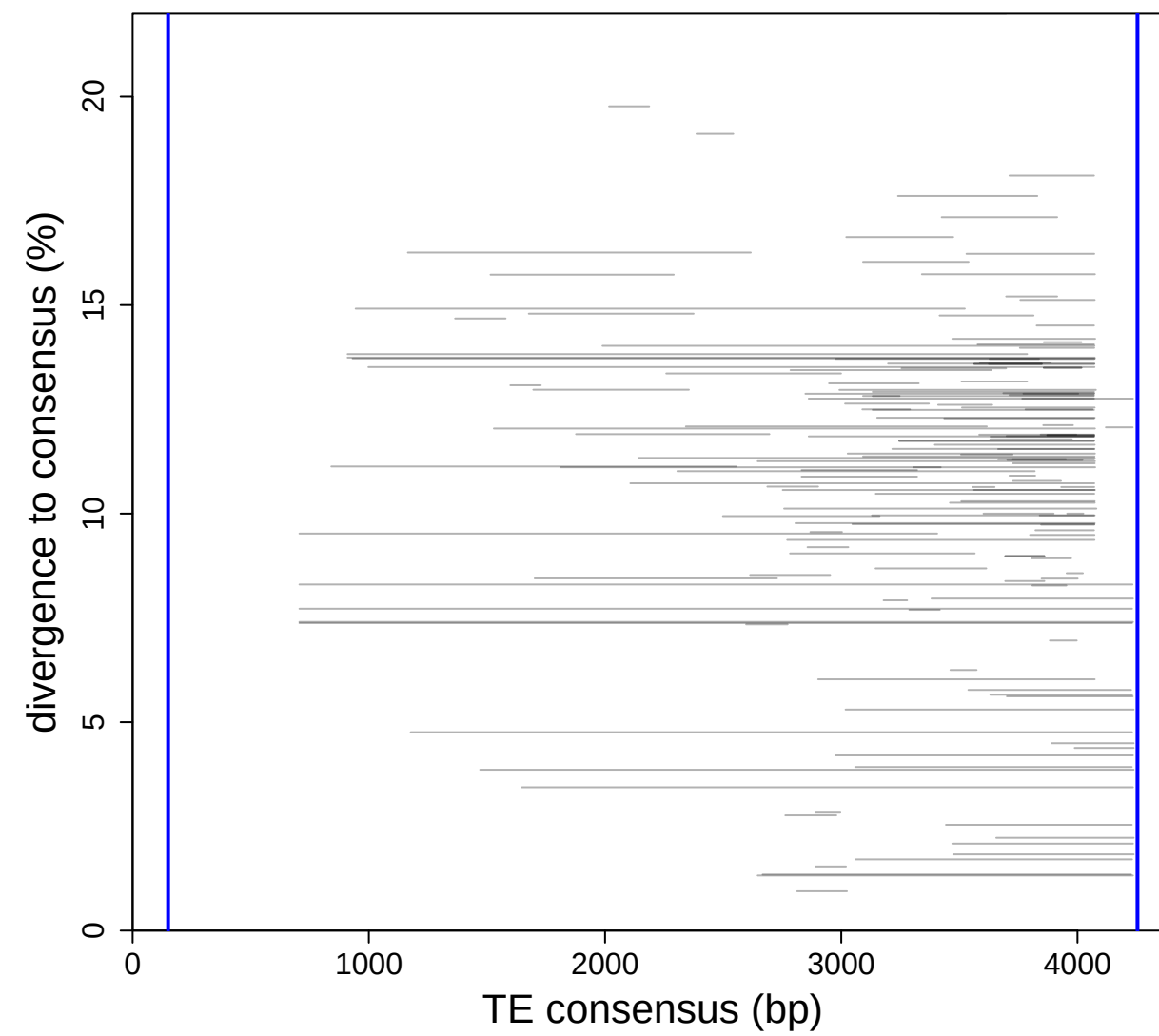
2.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcln.fa_aln.
size: 3952bp; fragments: 177; full length: 4 (>=3556.8bp)

After TEtrimmer 3952 bp

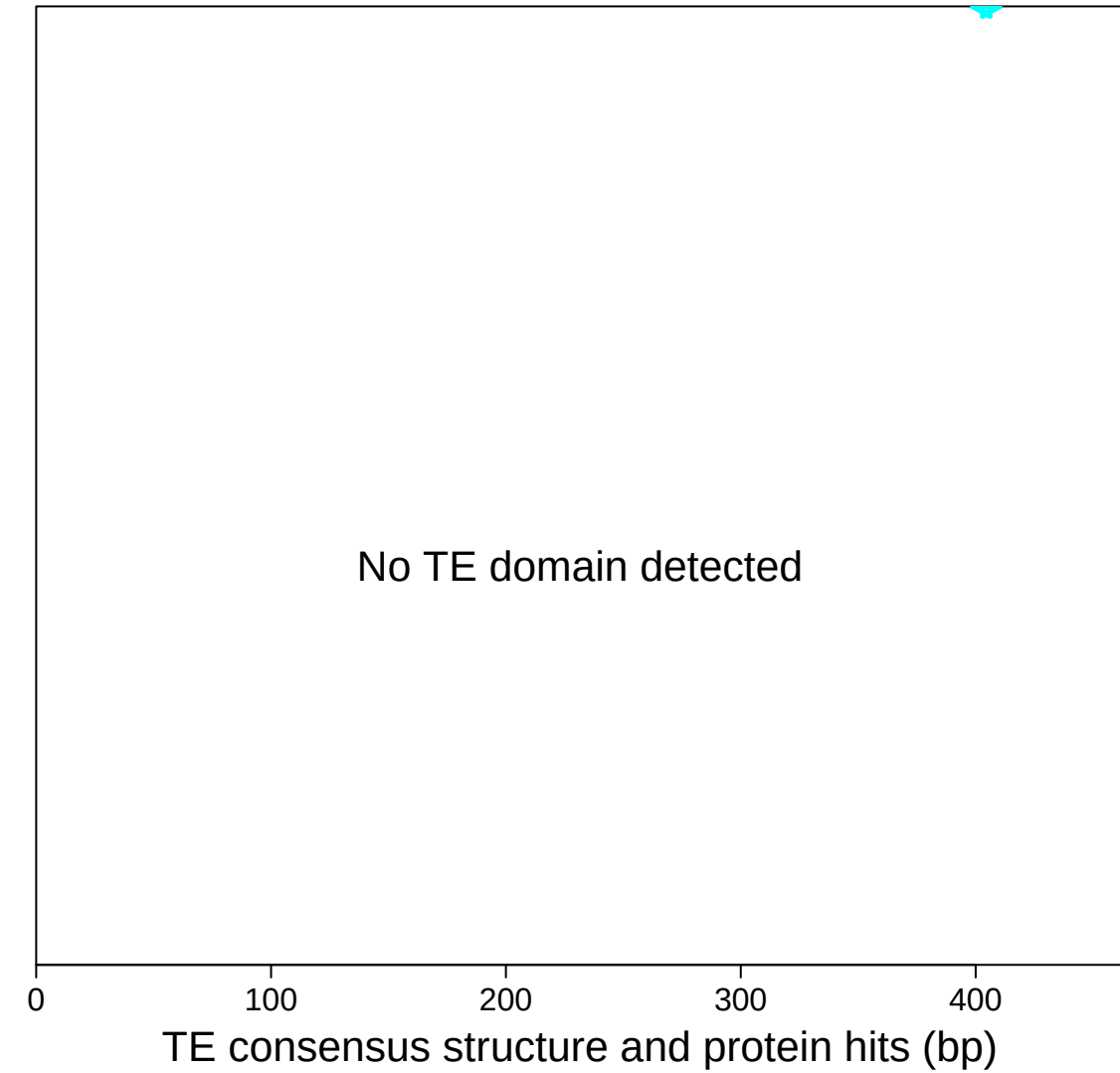
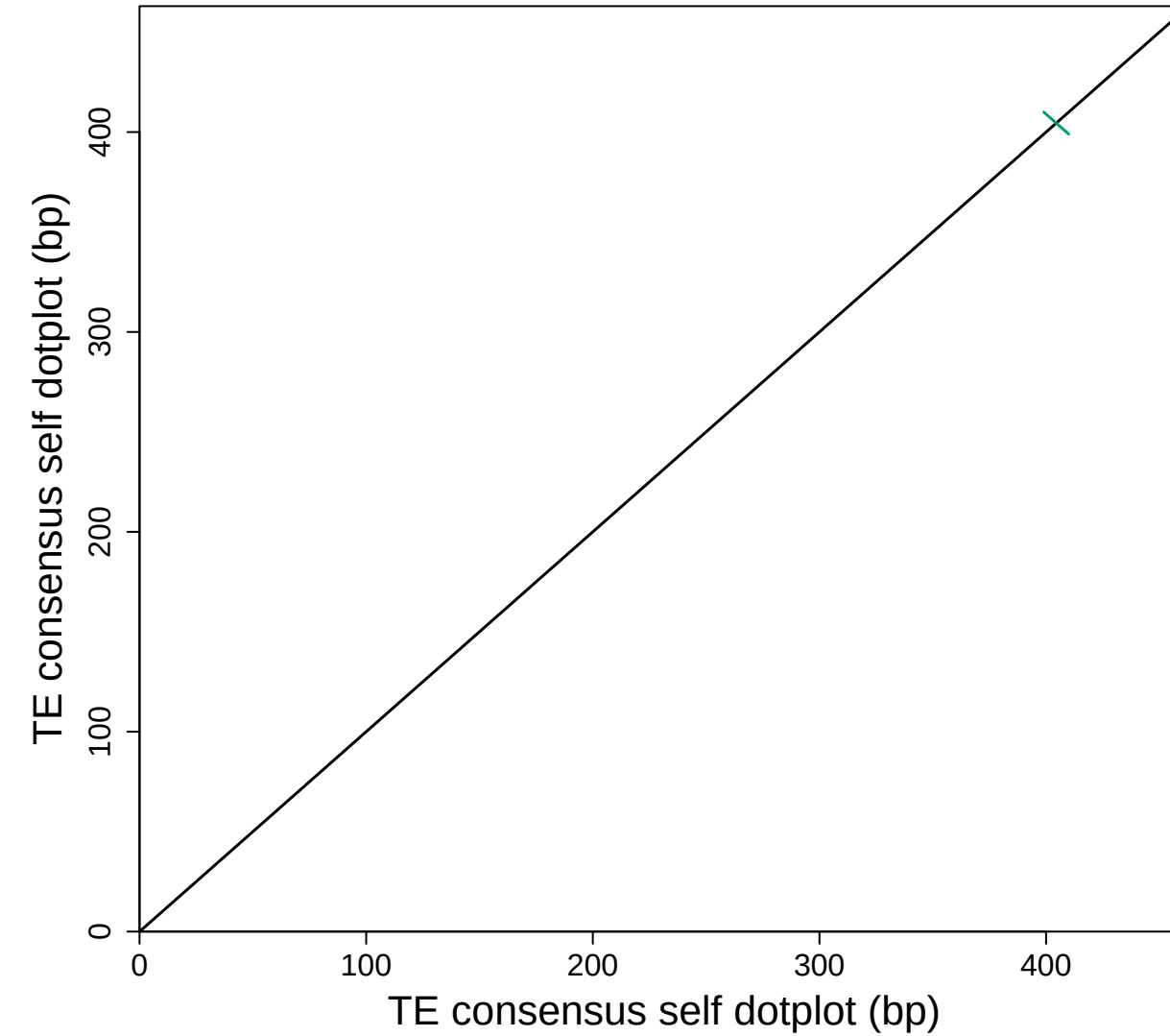
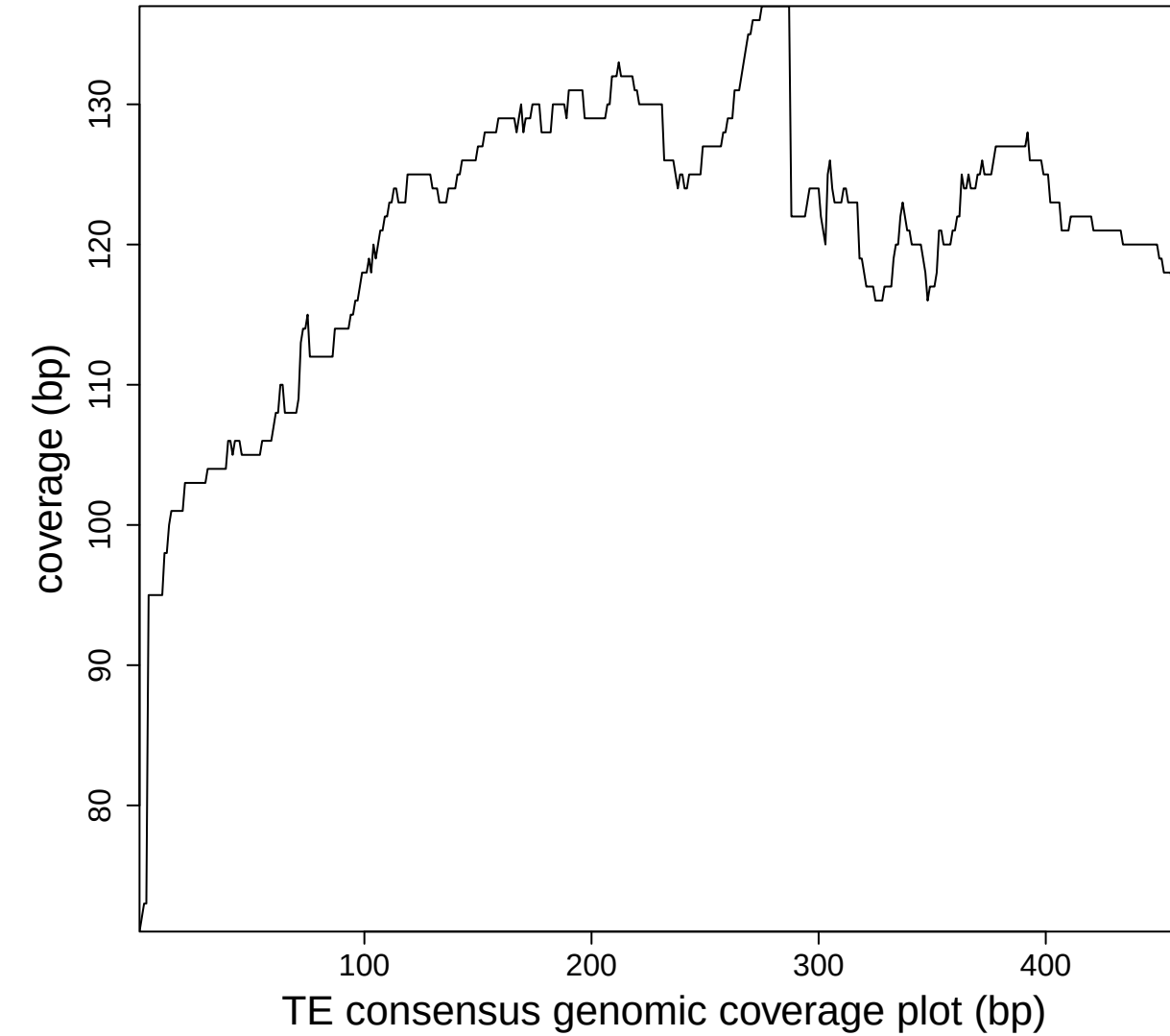
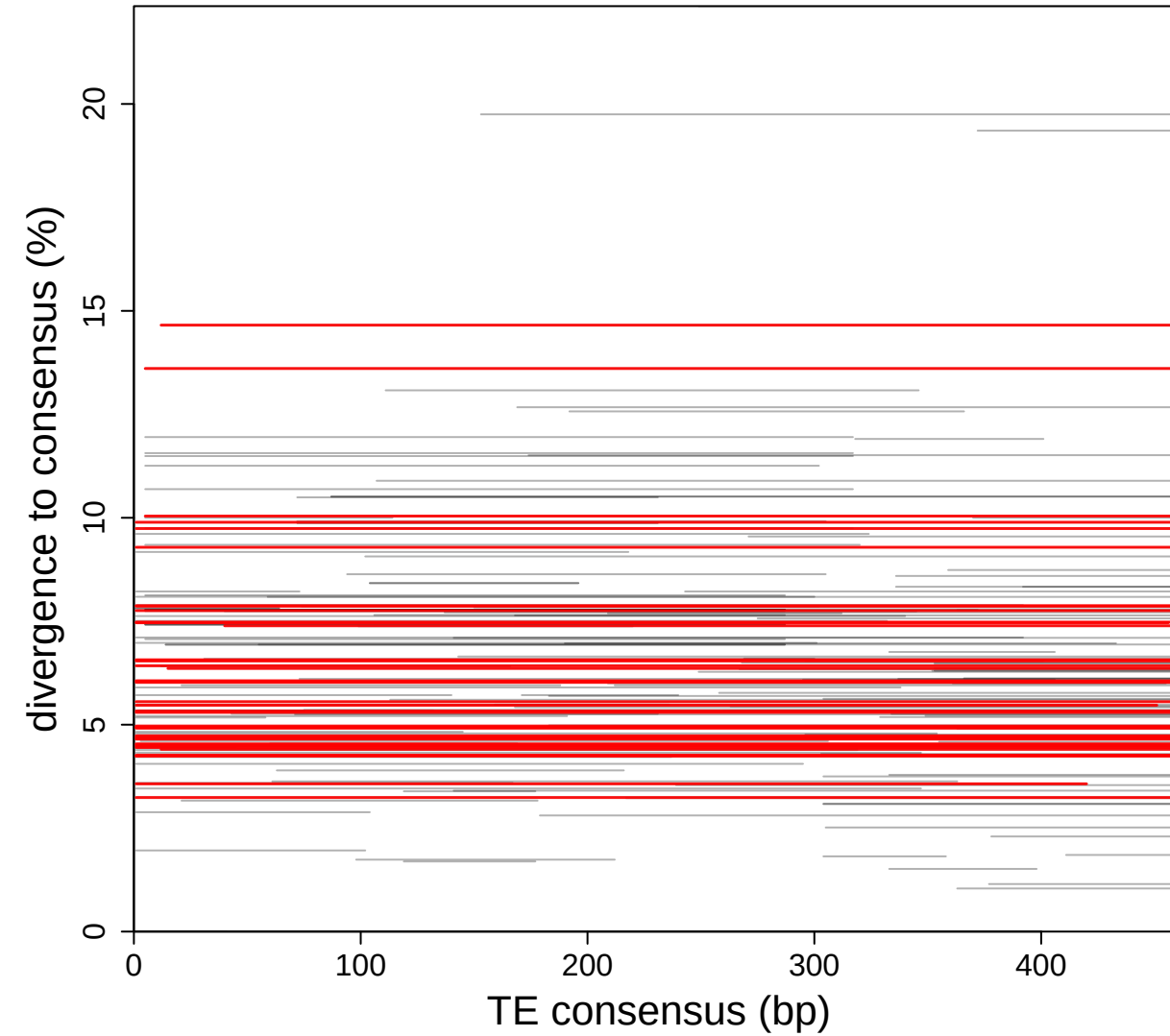


uf.bed g 2.bed fm 1.bed 0 0 n.bed g 1.bed fm 2.bed 0 0 b
size: 4404bp; fragments: 165; full length: 0 (>=3963.6bp)

After TEtrimmer Extended plot Blue lines are boundaries

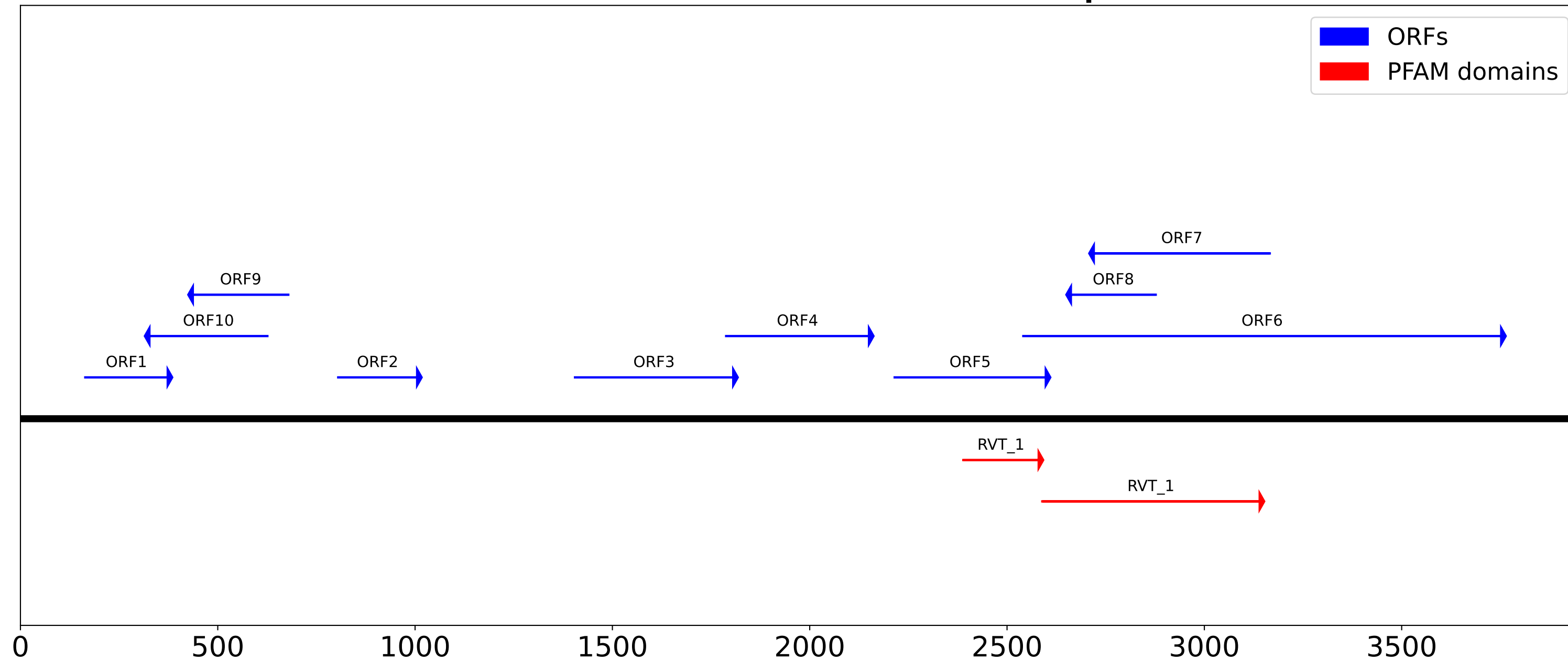


TE: rnd_5_family_10926
size: 463bp; fragments: 214; full length: 44 (≥ 416.7 bp)

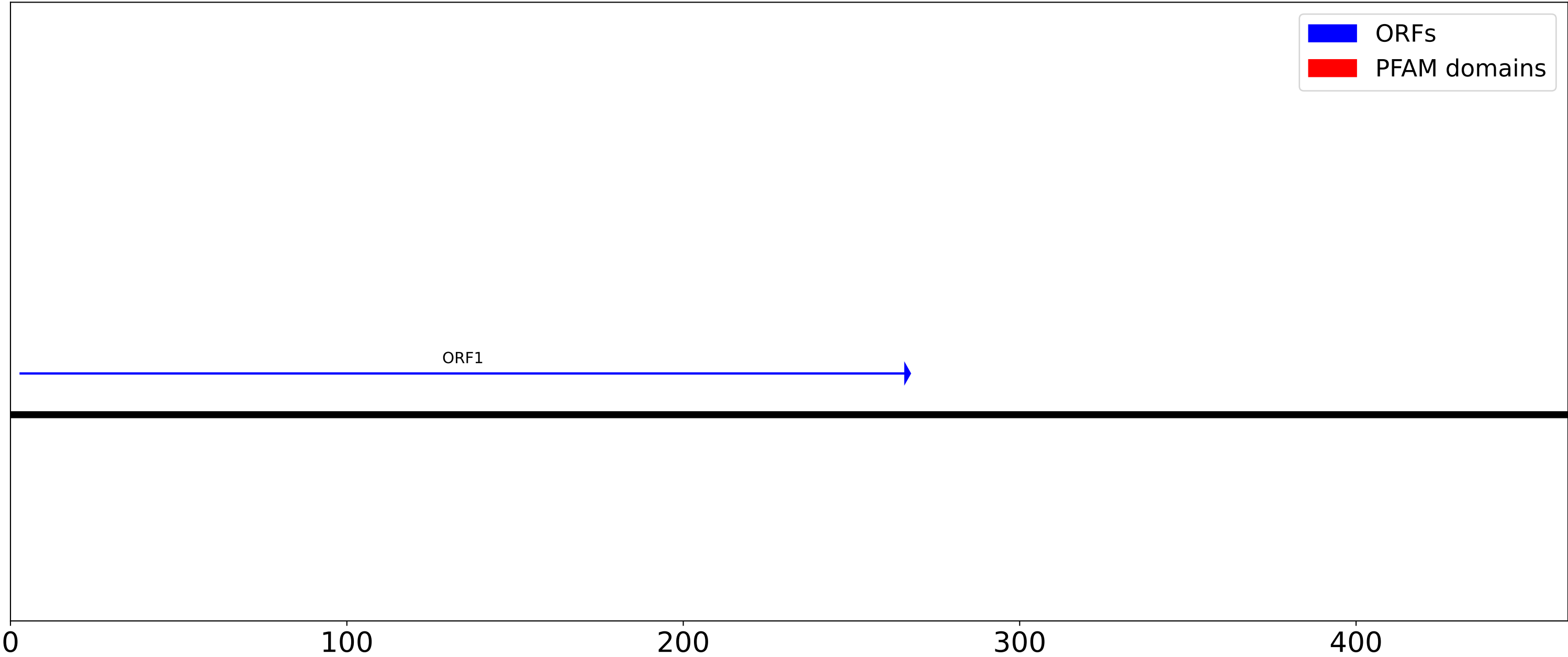


Before TEtrimmer 463 bp

After TEtrimmer ORF and PFAM domain plot



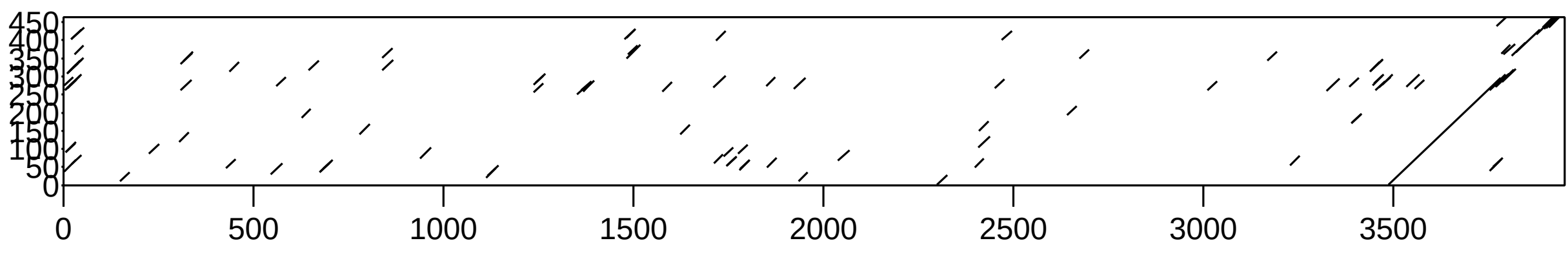
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

sensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)