

Start crop Point

End crop Point

MSA length = 5102



Start crop Point

End crop Point

0

10

0

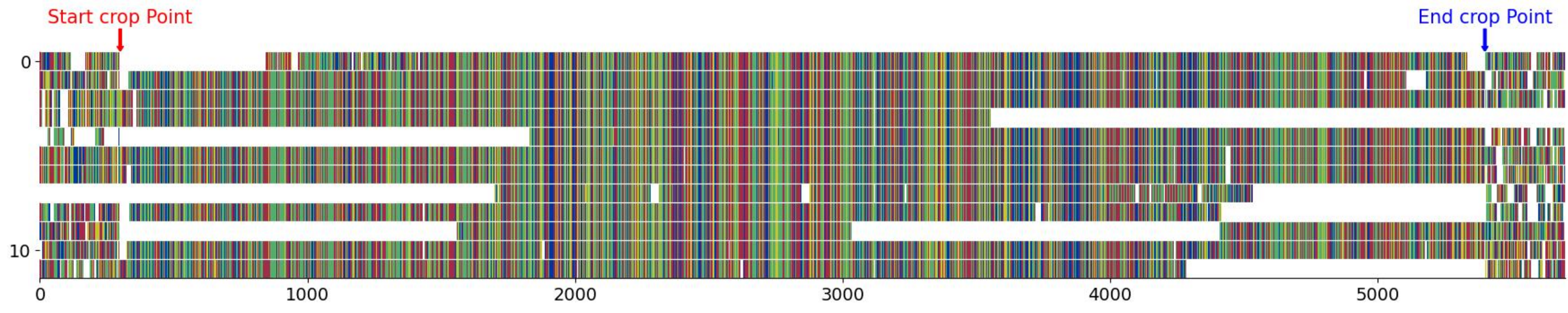
1000

2000

3000

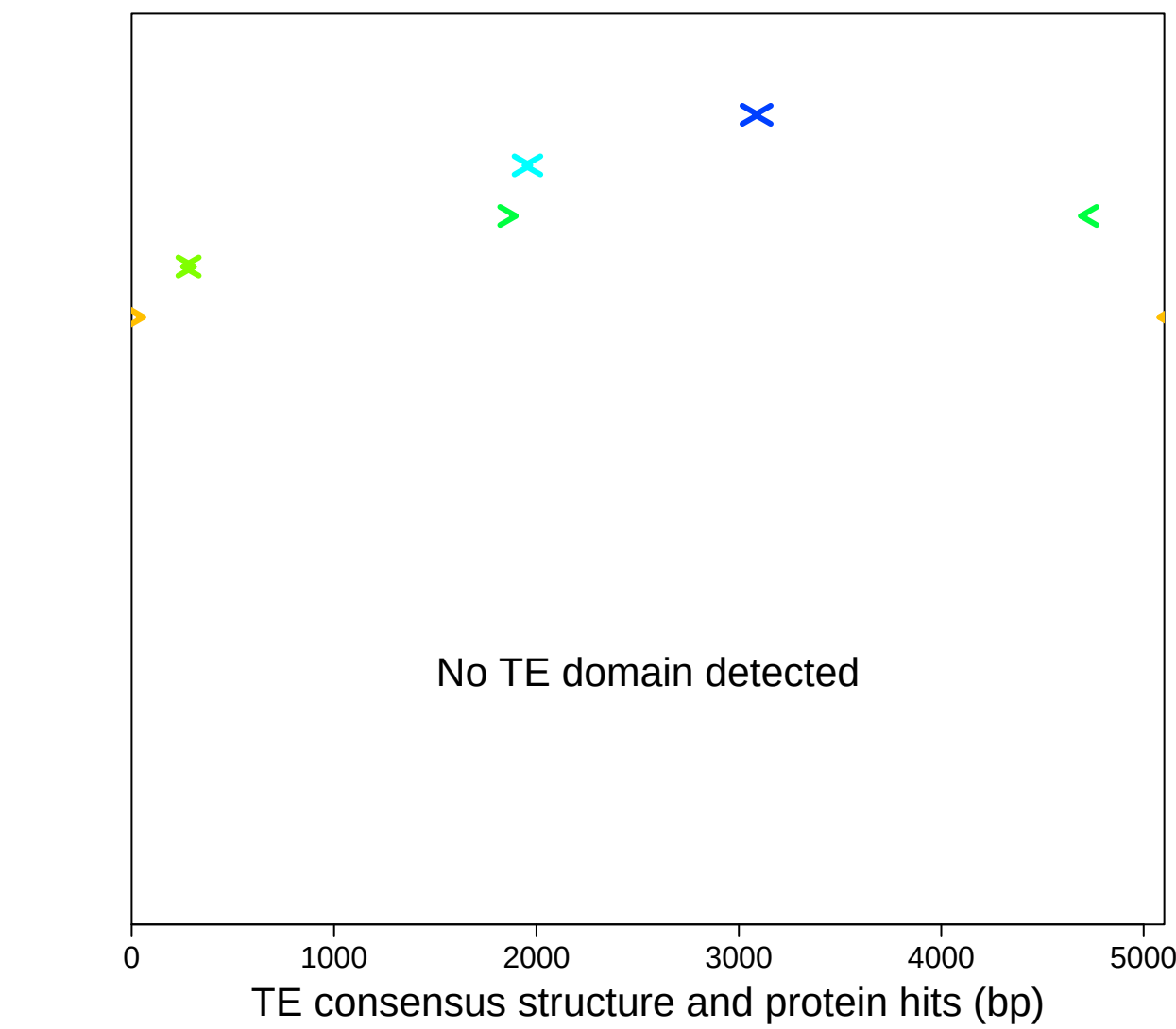
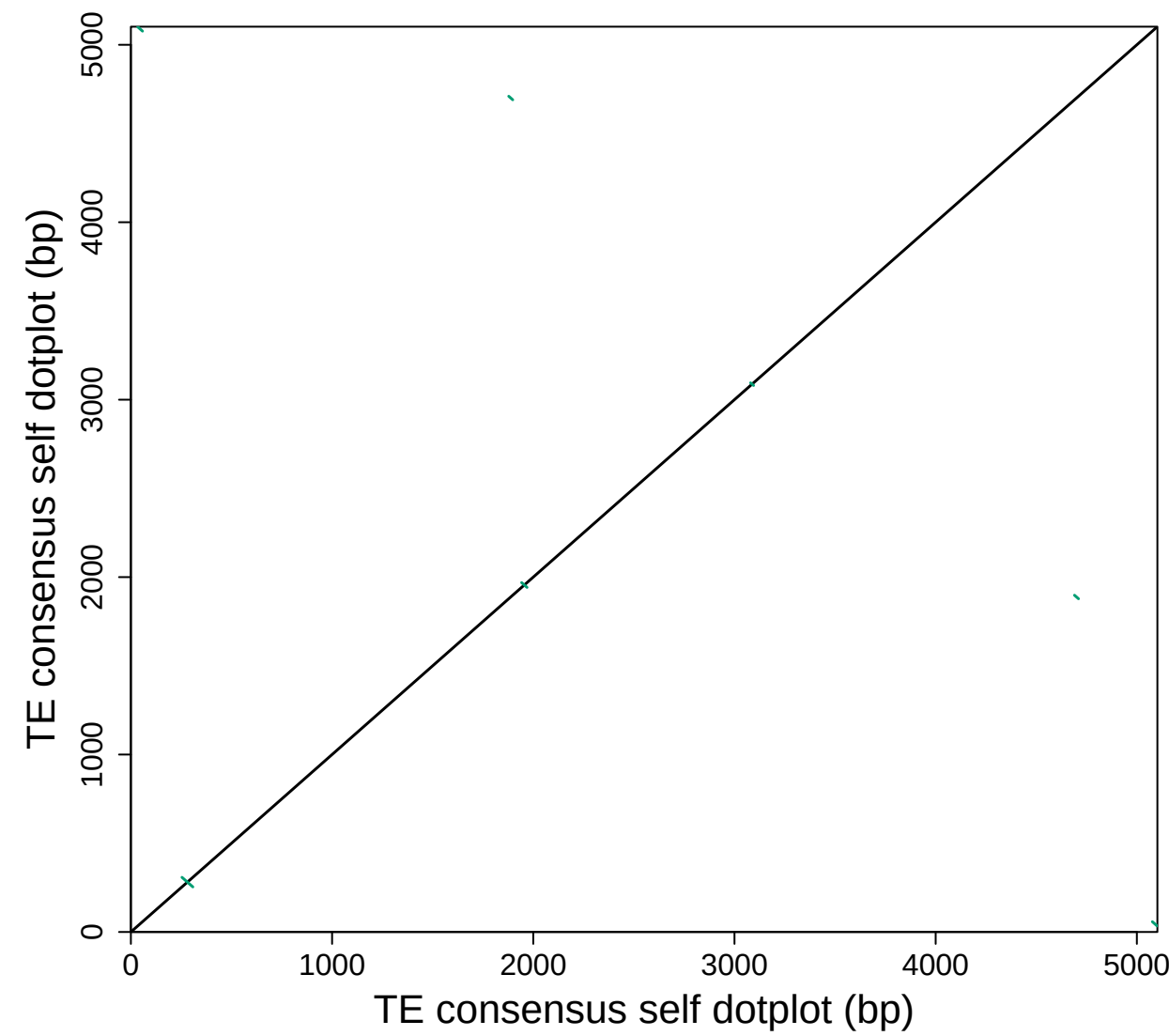
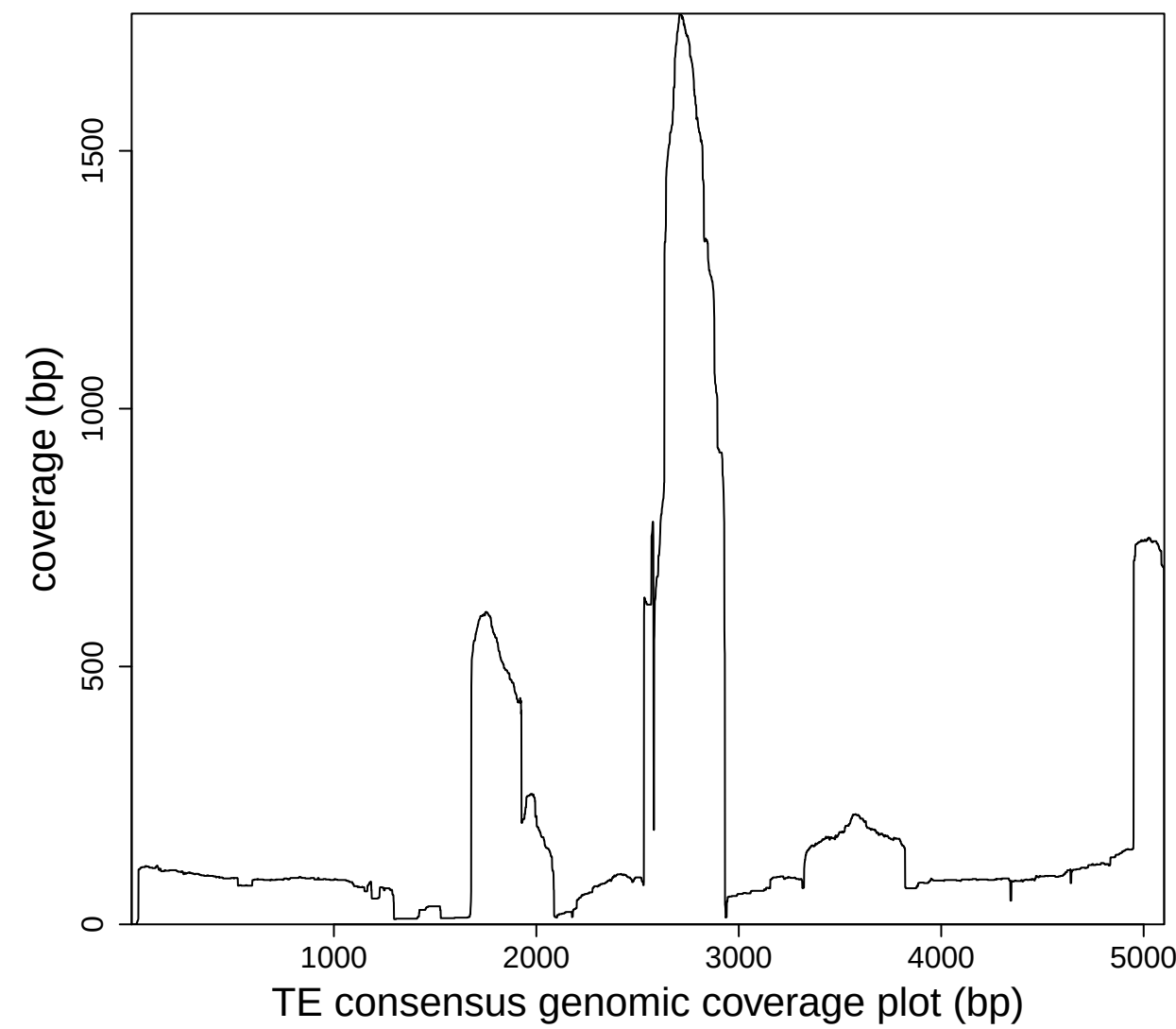
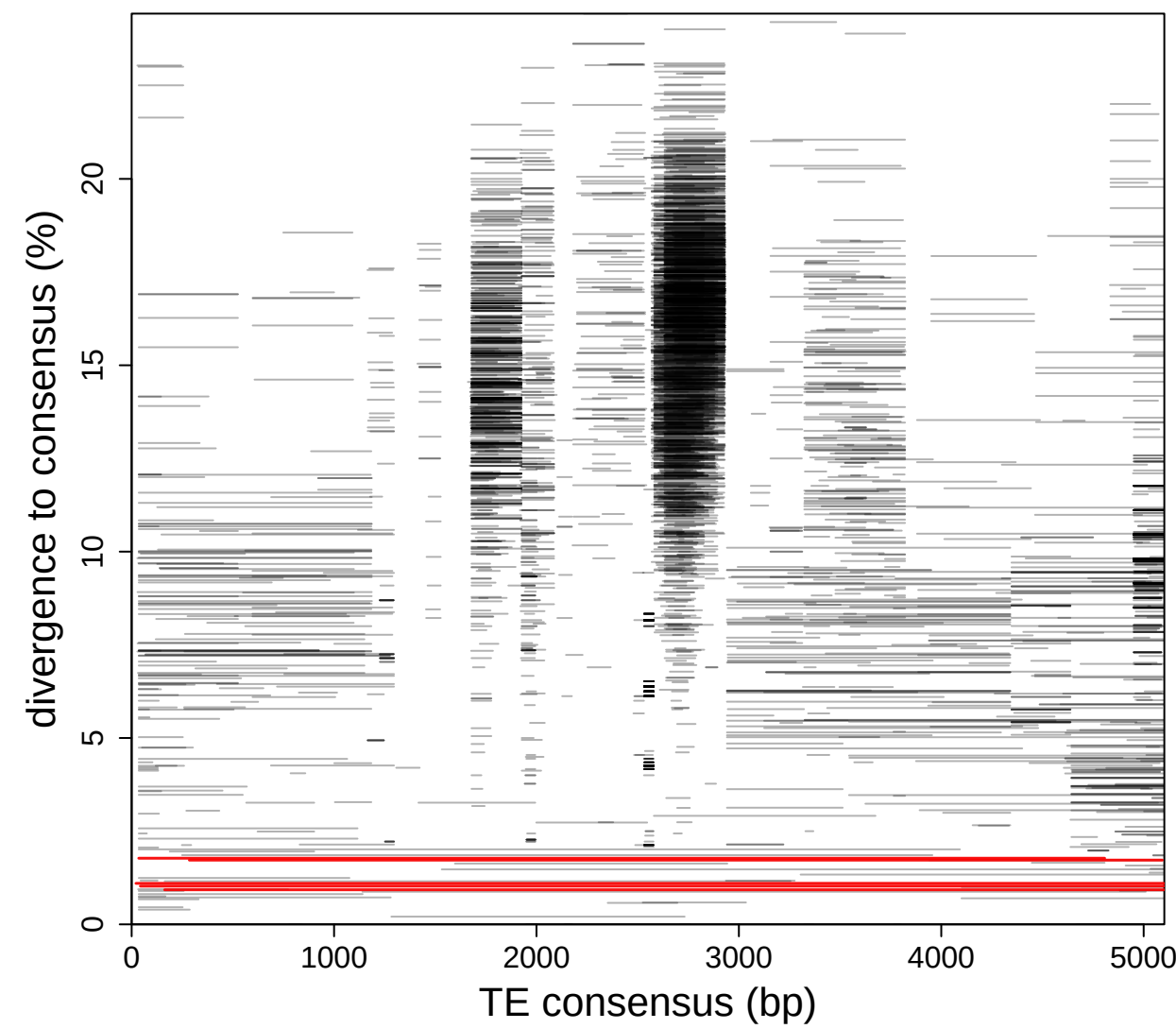
4000

5000



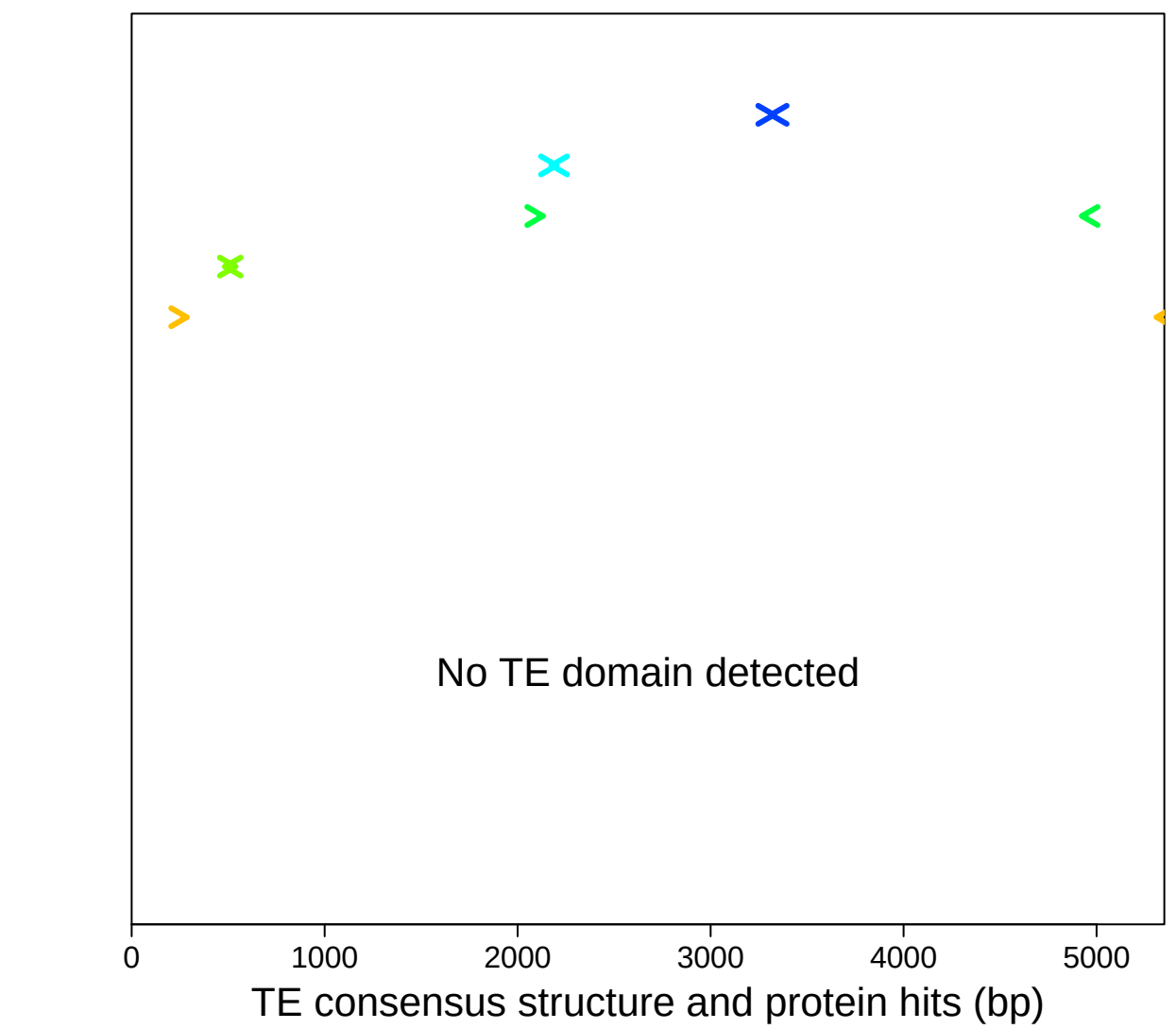
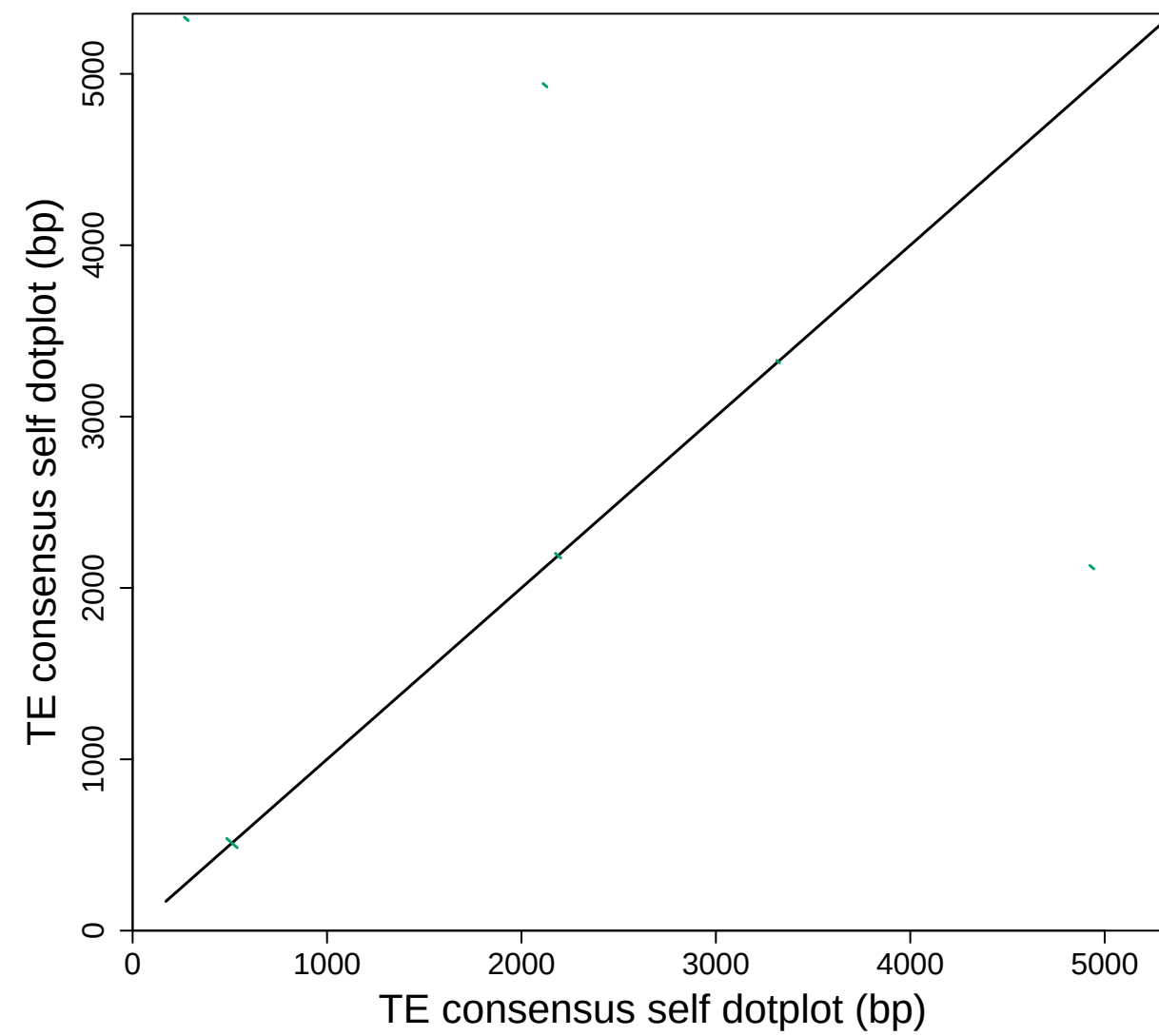
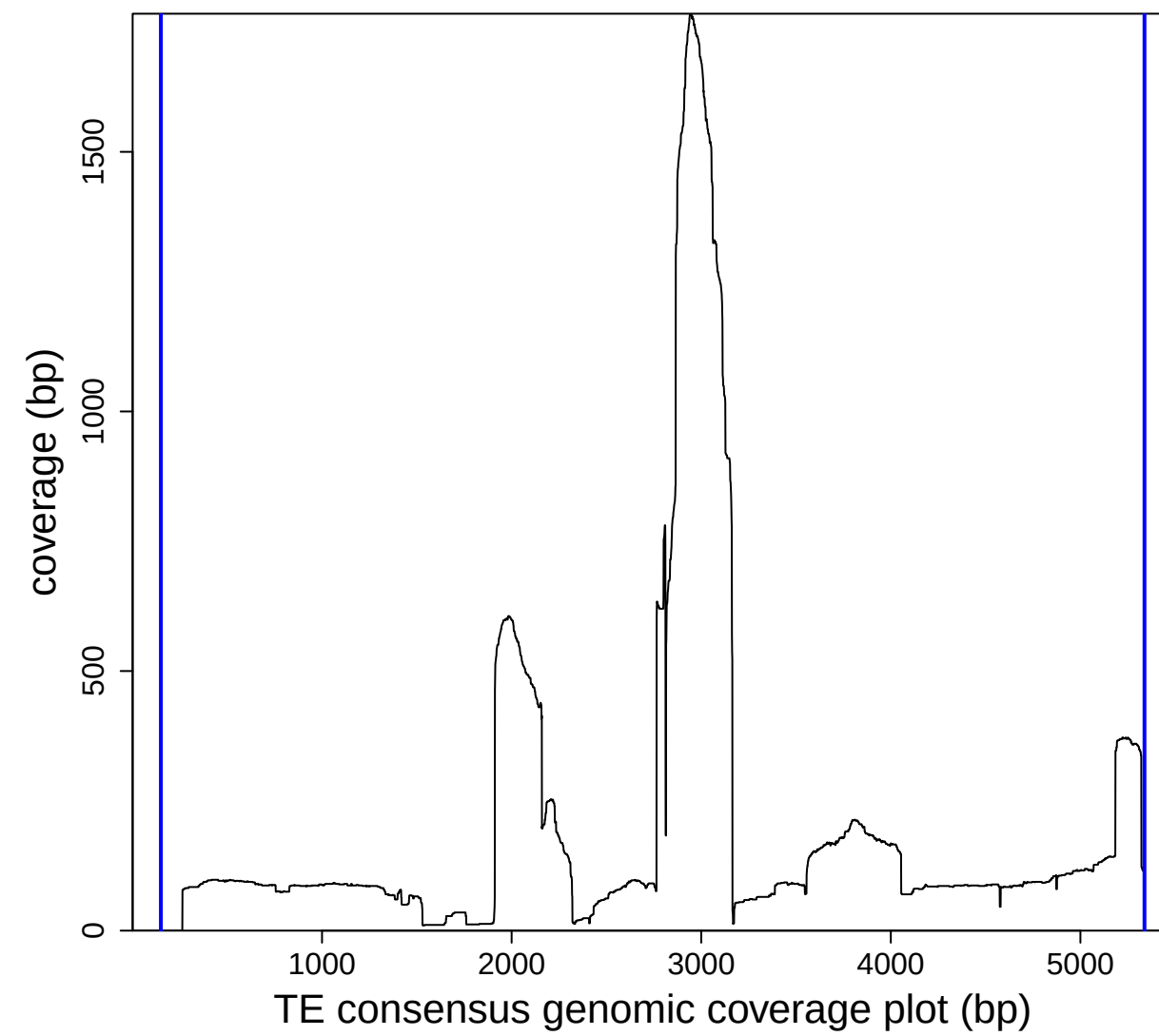
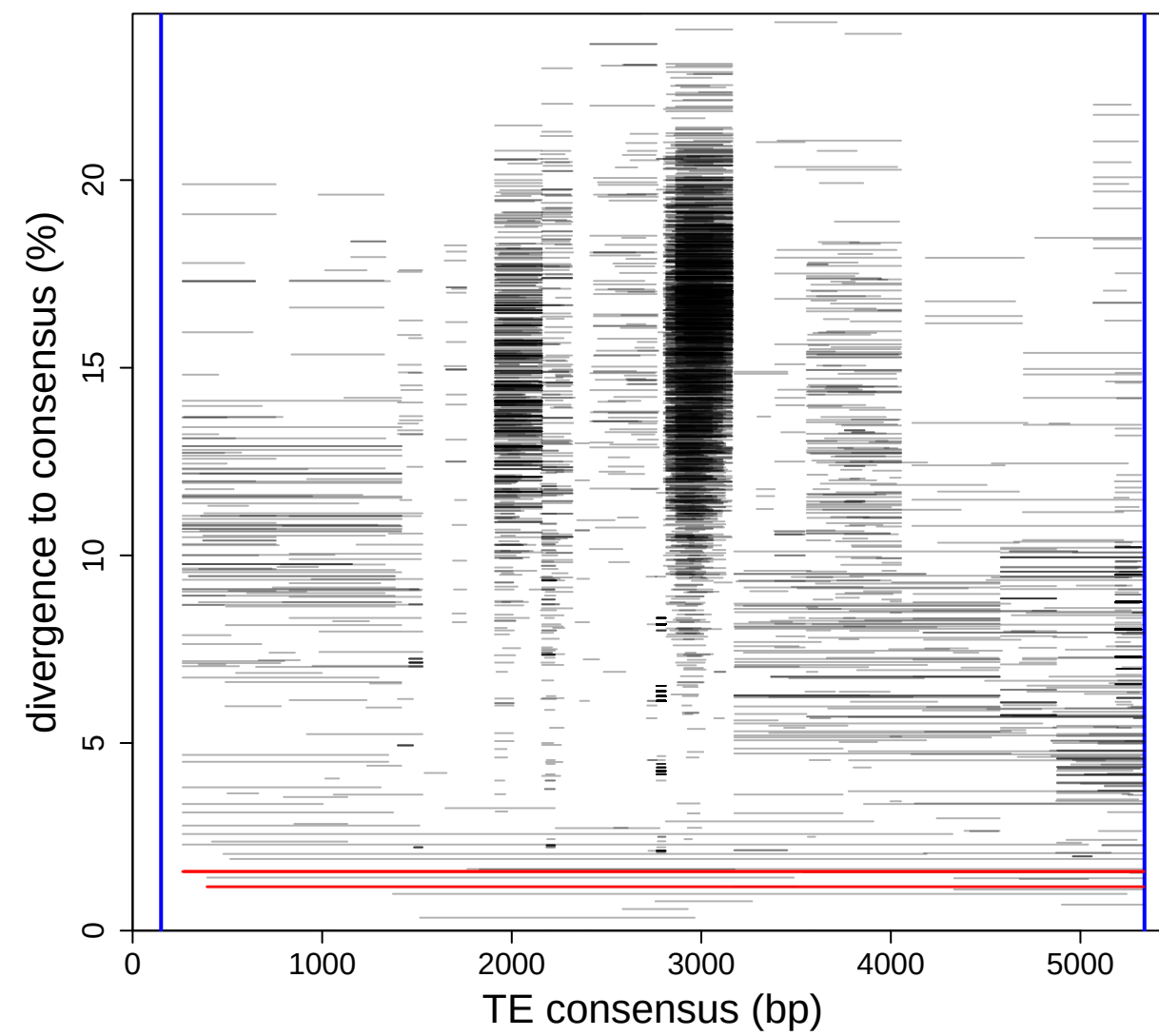
g_2.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcln.fa_ε
size: 5102bp; fragments: 4781; full length: 5 (>=4591.8bp)

After TEtrimmer 5102 bp

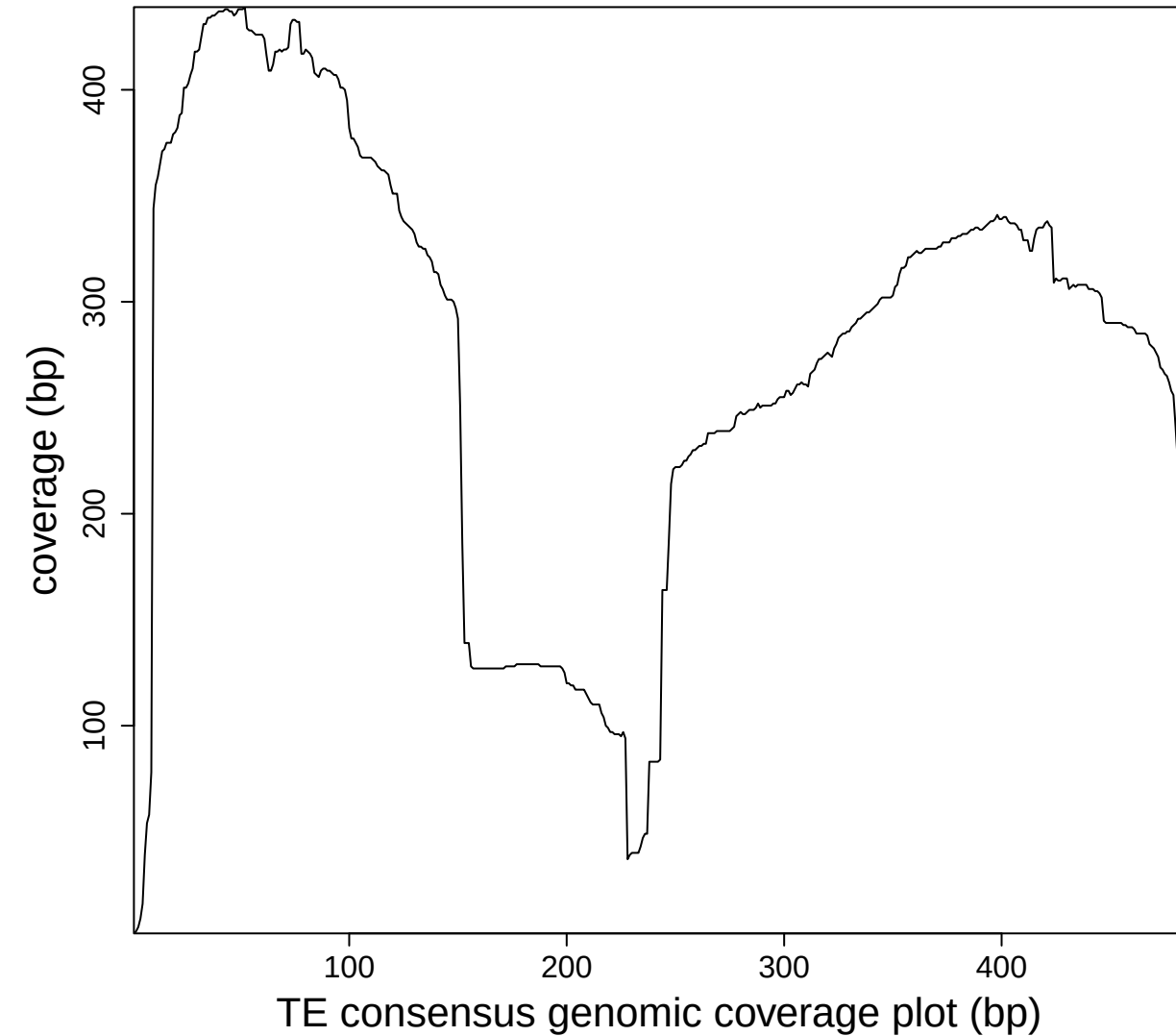
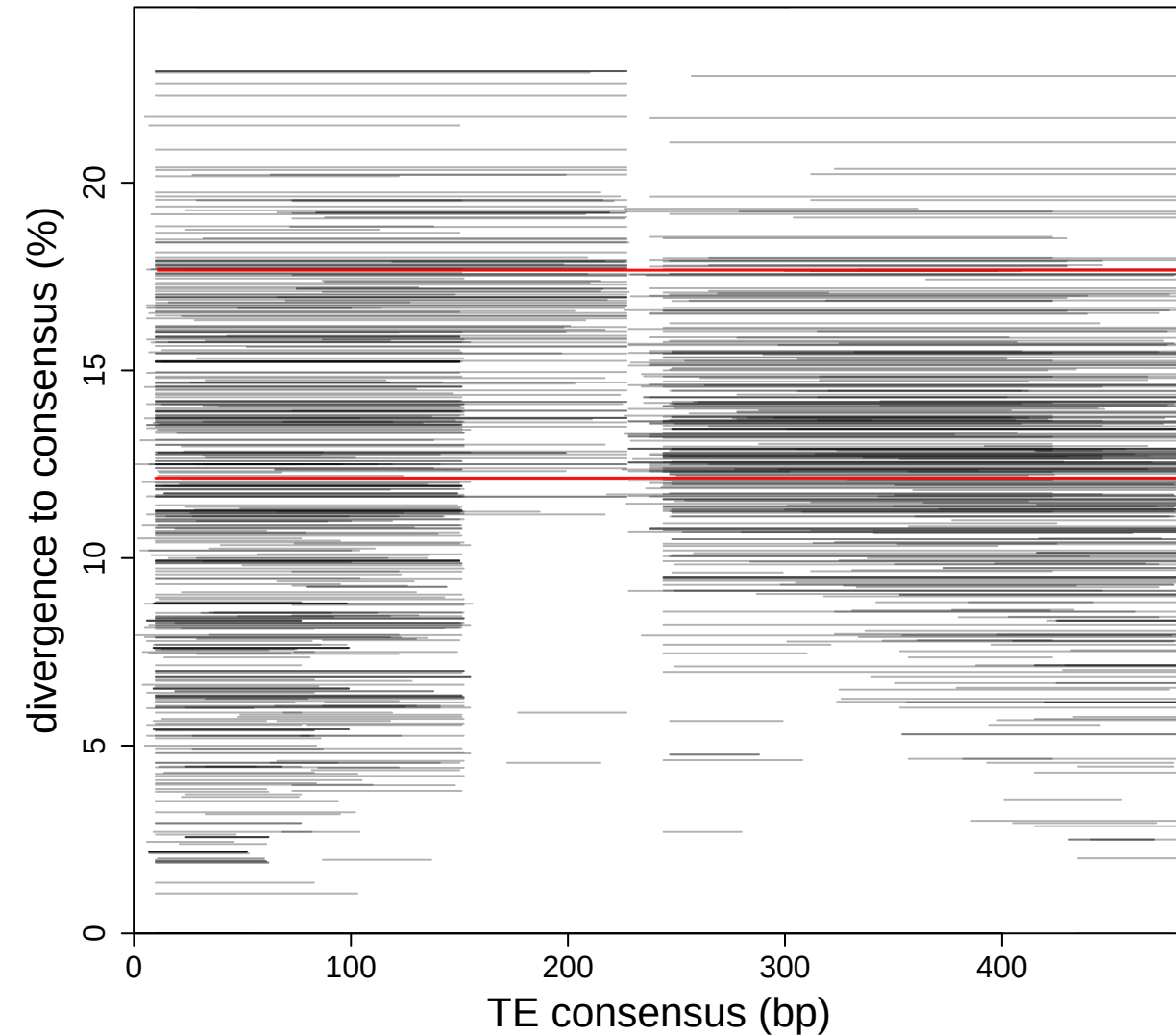


f.bed_g 2.bed_fm 1.bed_0 0 n.bed_g 1.bed_fm 2.bed_0 0 bc
size: 5488bp; fragments: 4365; full length: 3 (>=4939.2bp)

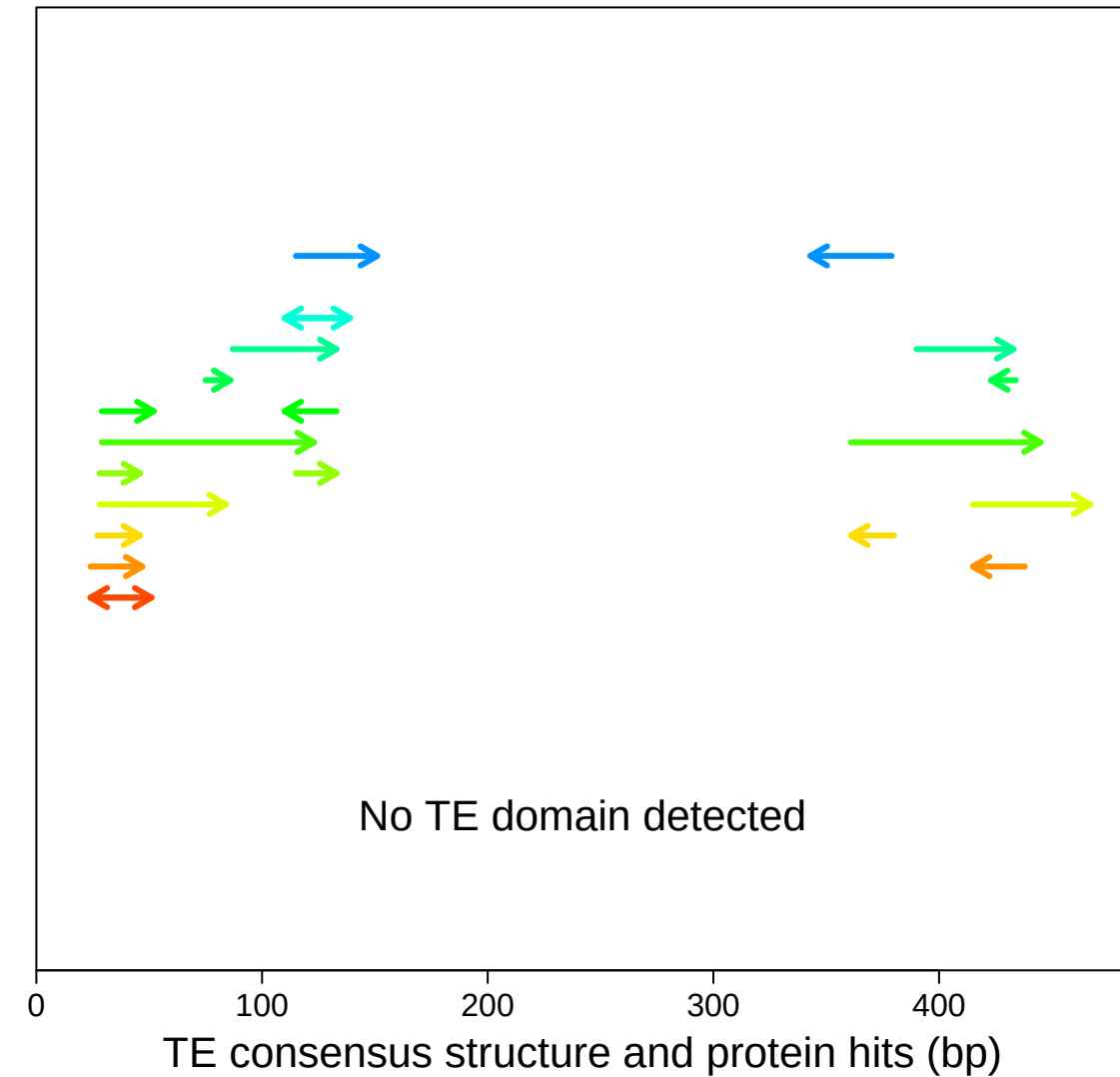
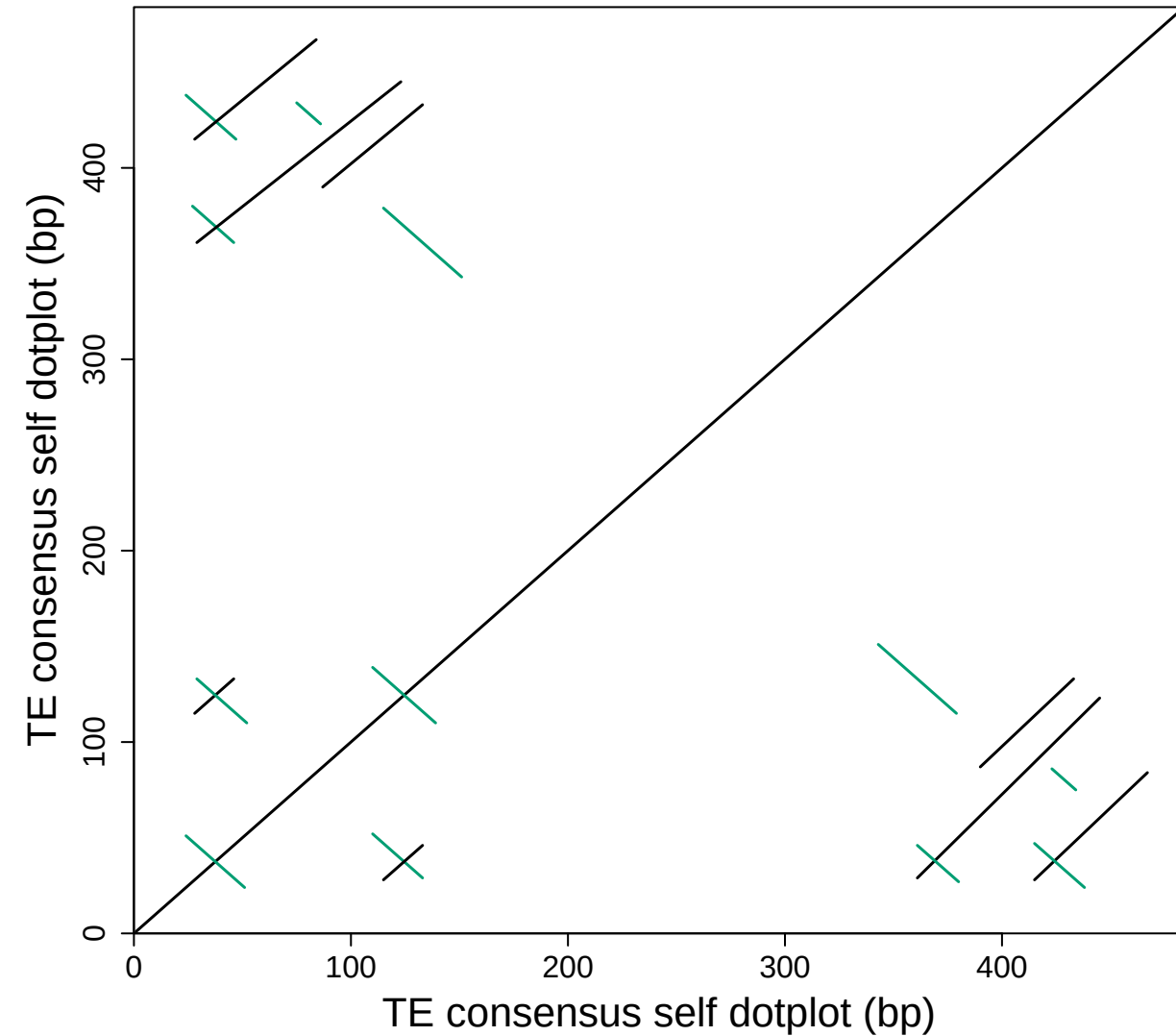
After TETrimmer Extended plot Blue lines are boundaries



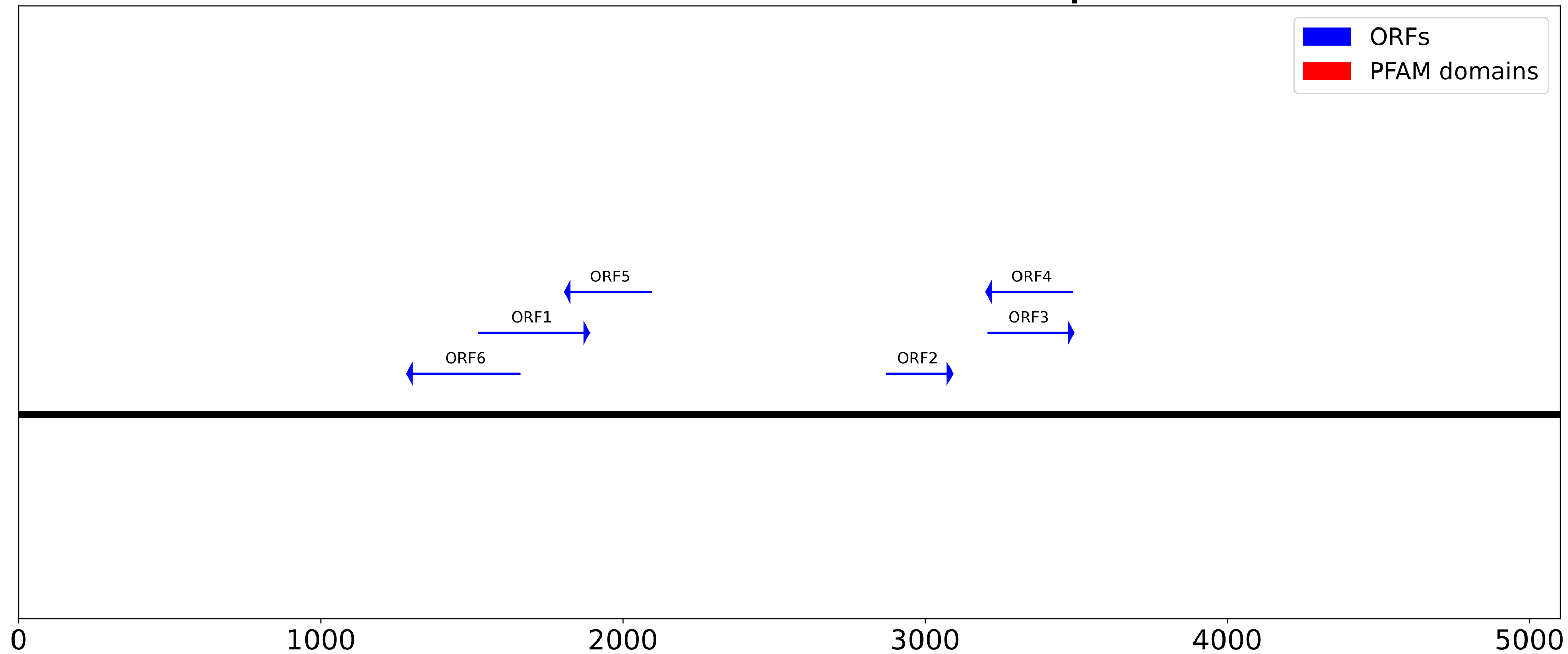
TE: rnd_4_family_1396
size: 484bp; fragments: 884; full length: 2 (≥ 435.6 bp)



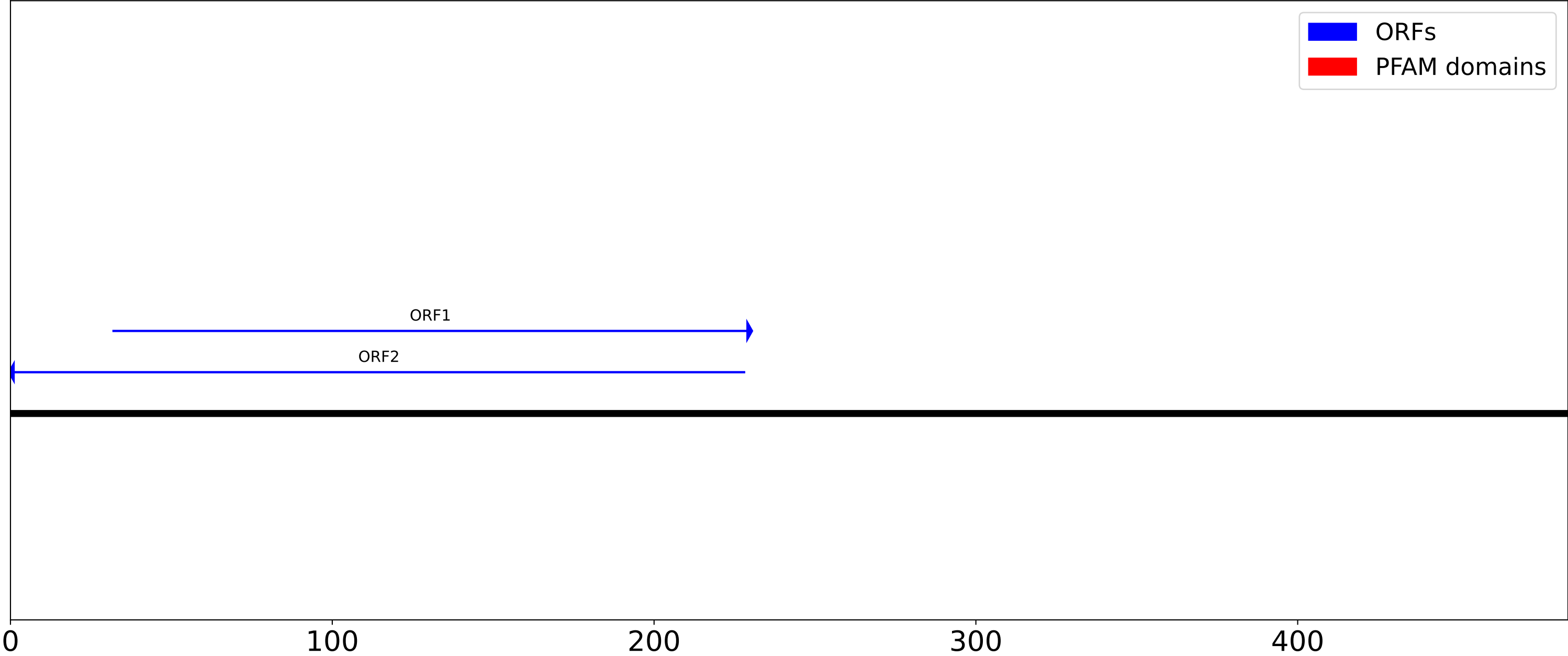
Before TEtrimmer 484 bp



After TEtrimmer ORF and PFAM domain plot



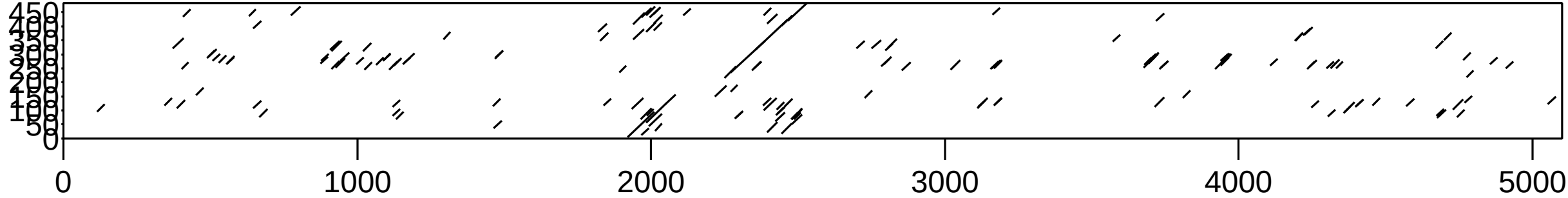
Before TEtrimmer ORF and PFAM domain plot



Dotplot

(windowsize = 25, threshold = 50.00 09/10/25)

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)