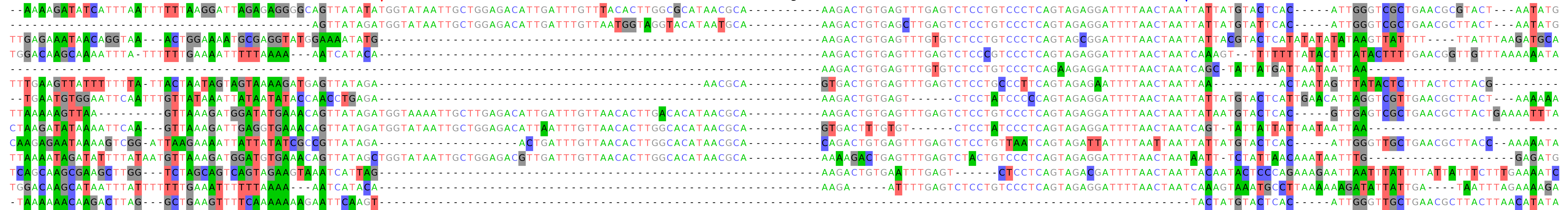


Start crop Point

End crop Point

MSA length = 735



Start crop Point

End crop Point

0

10

0

200

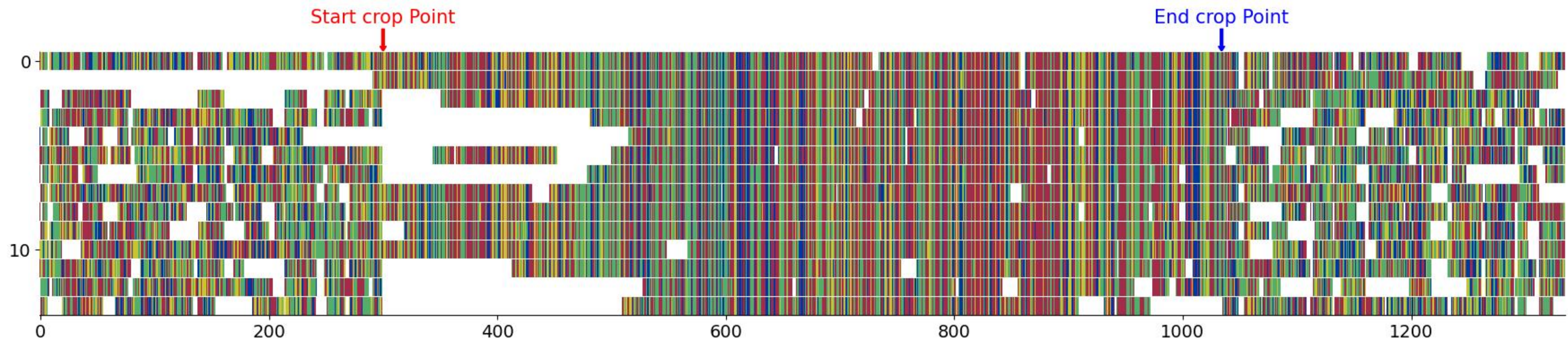
400

600

800

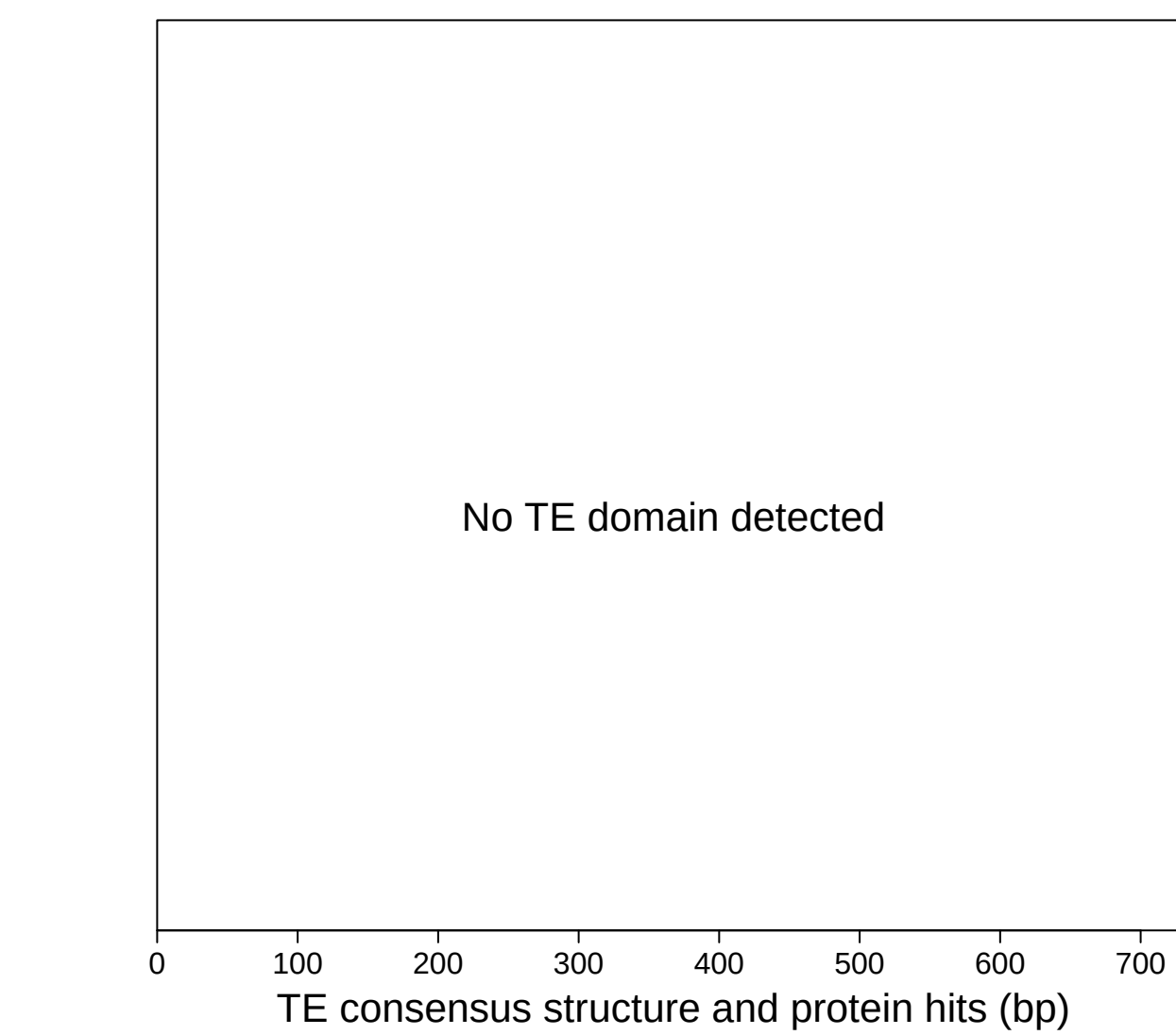
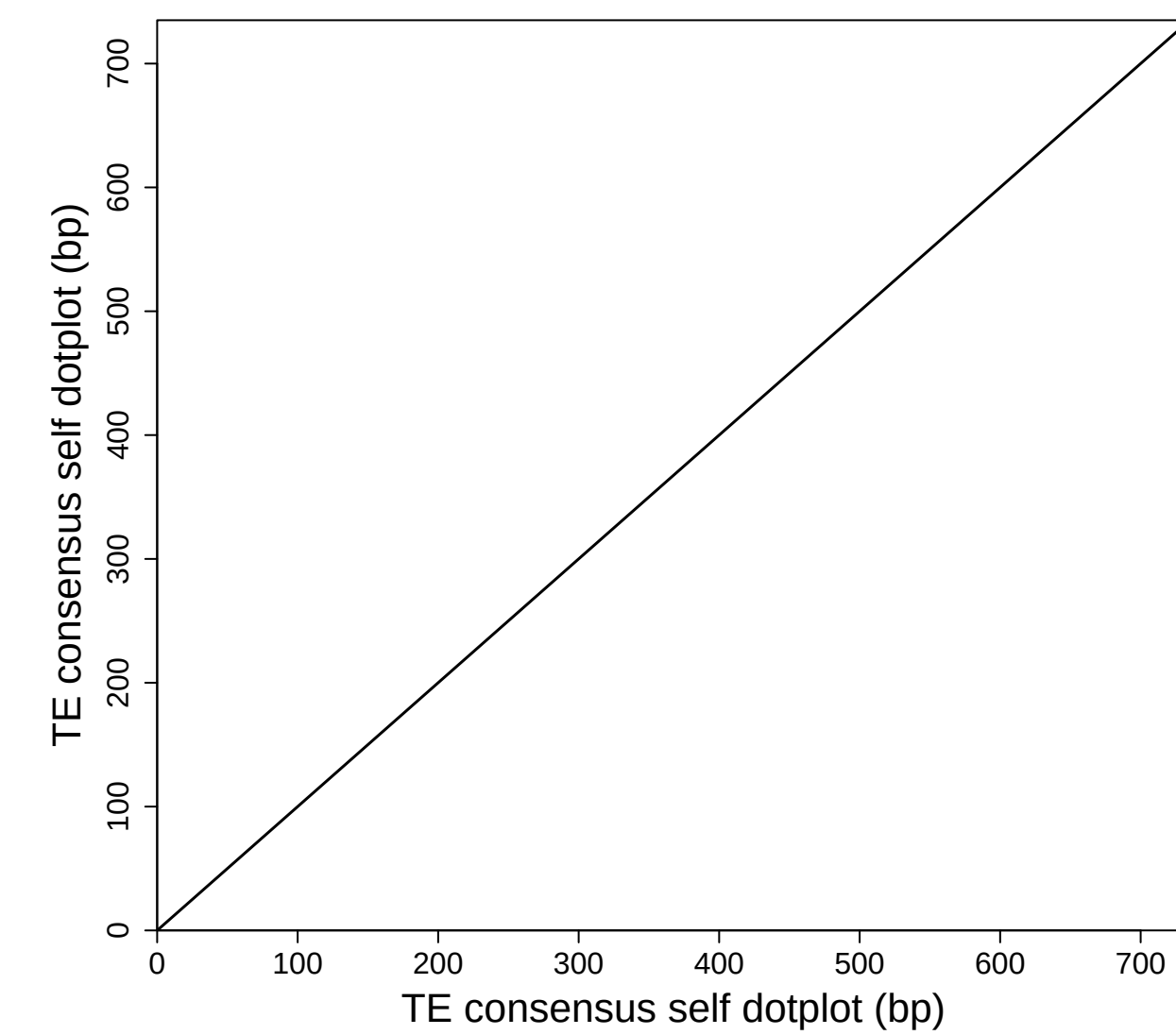
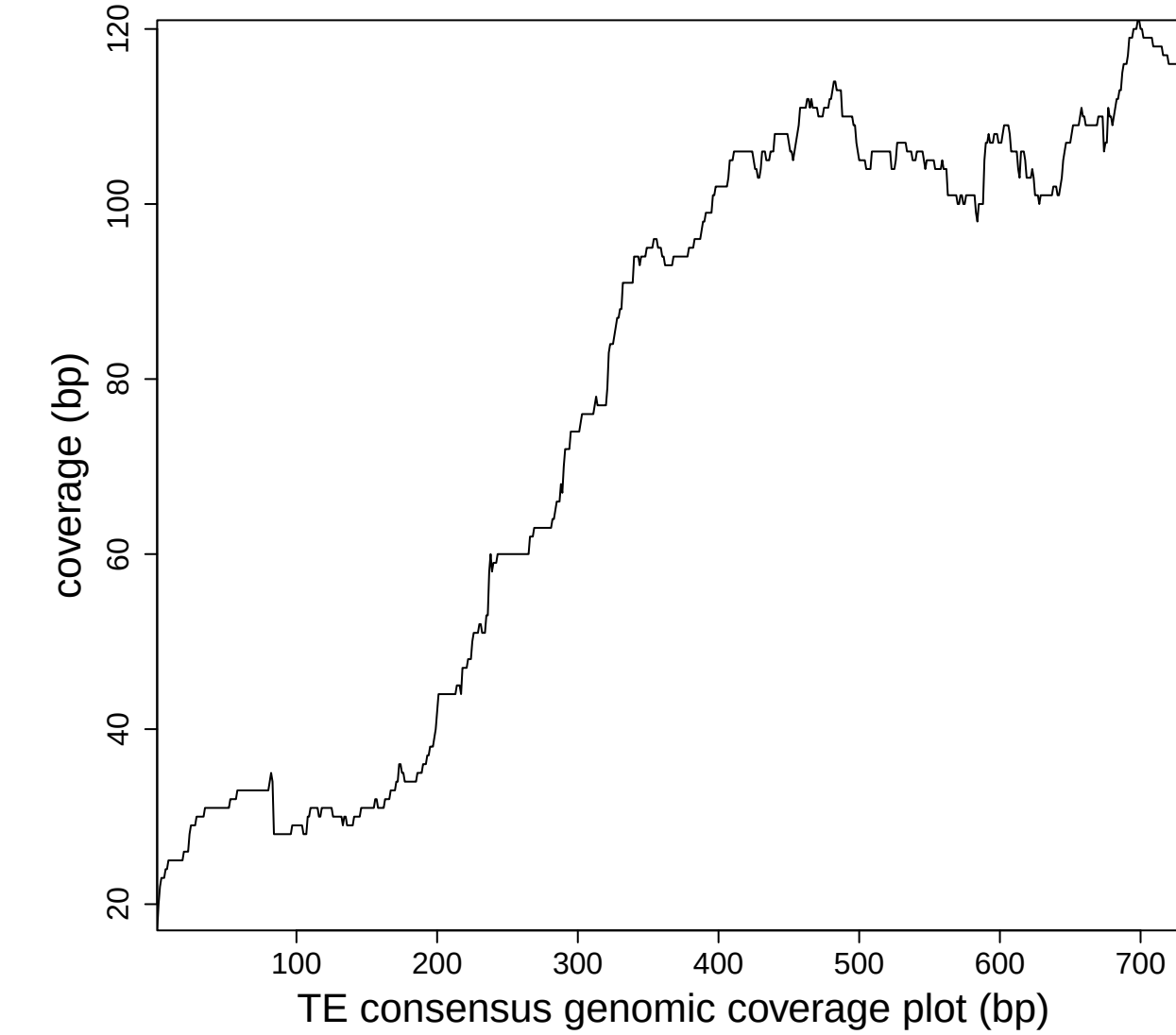
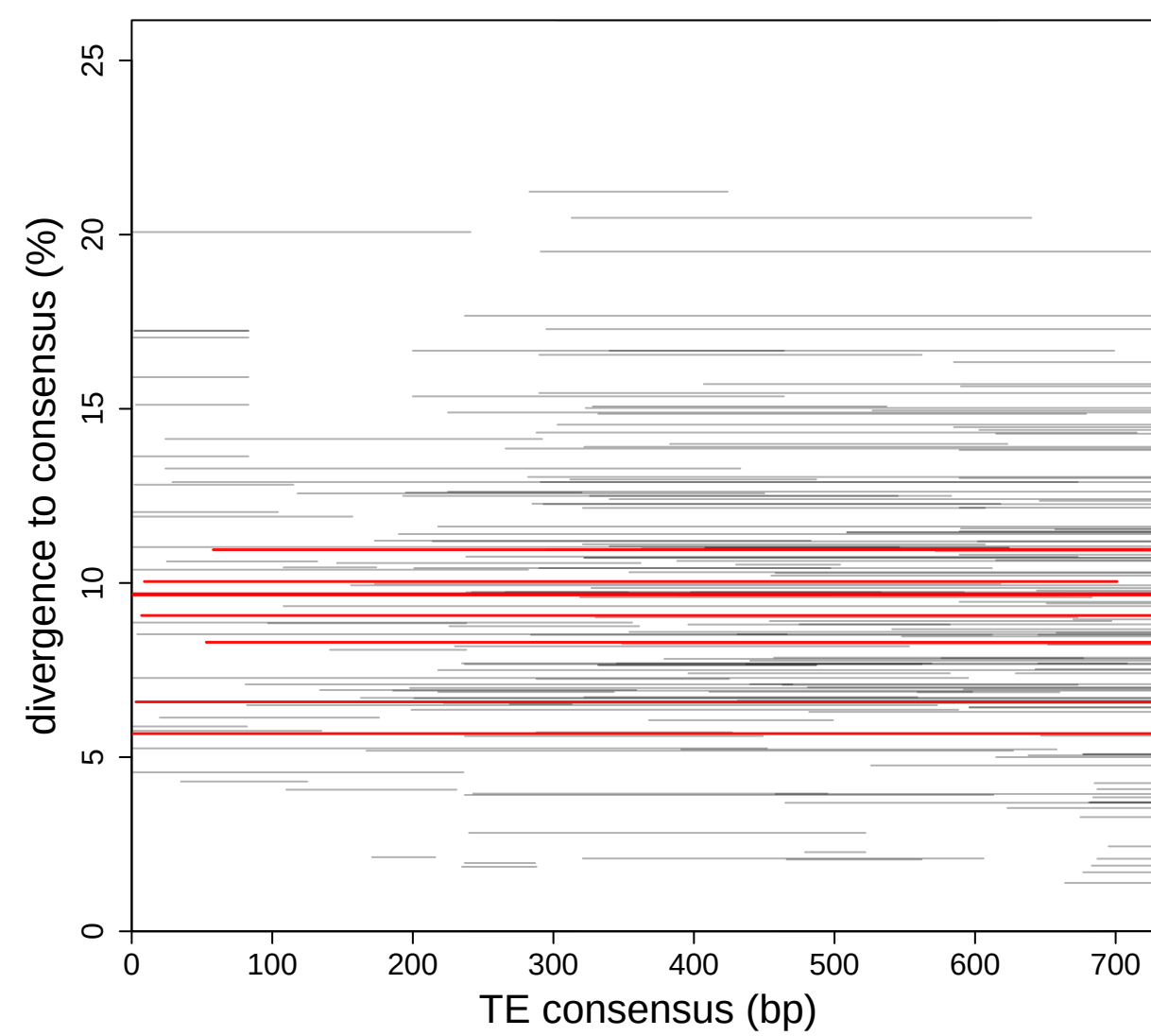
1000

1200



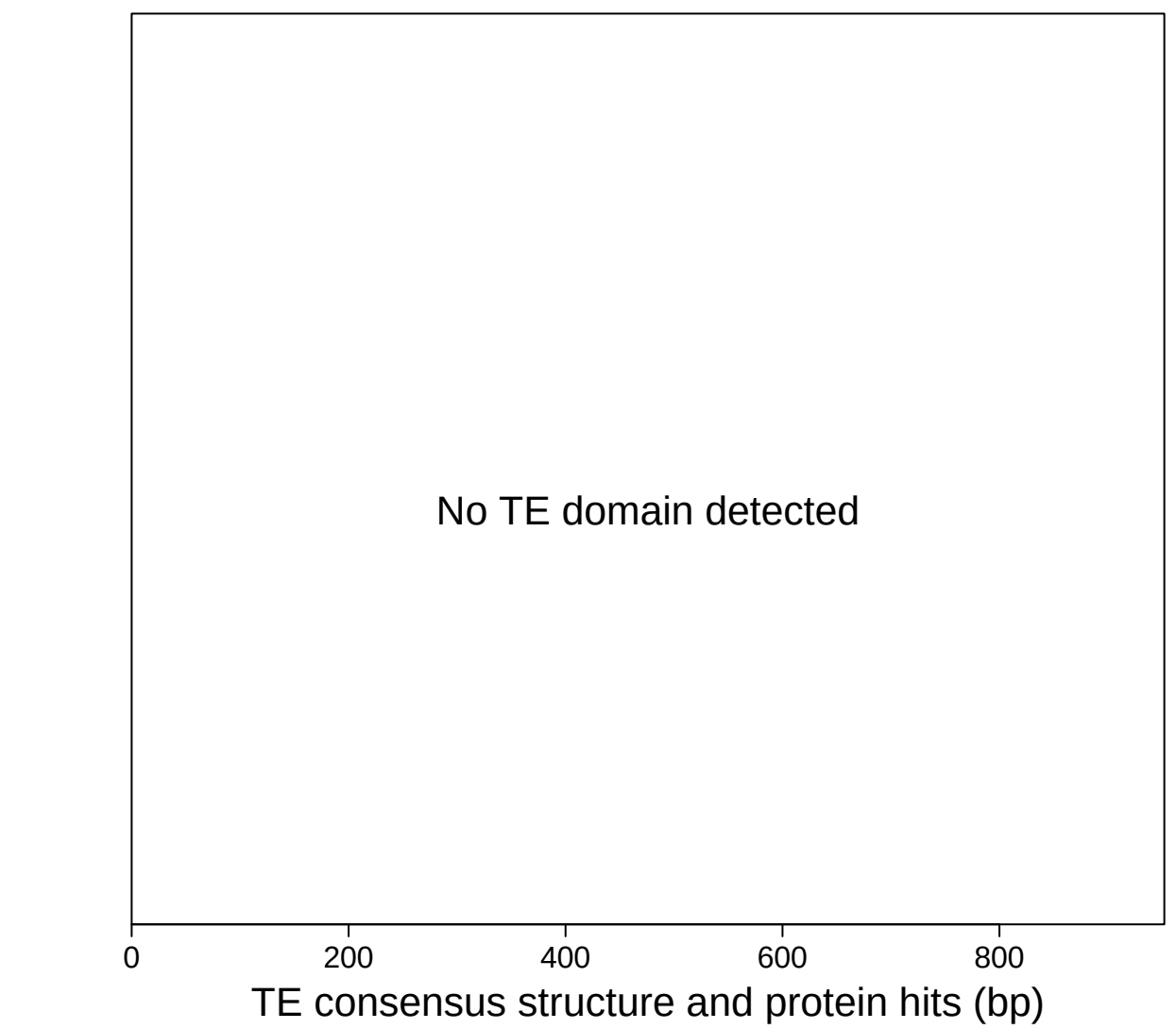
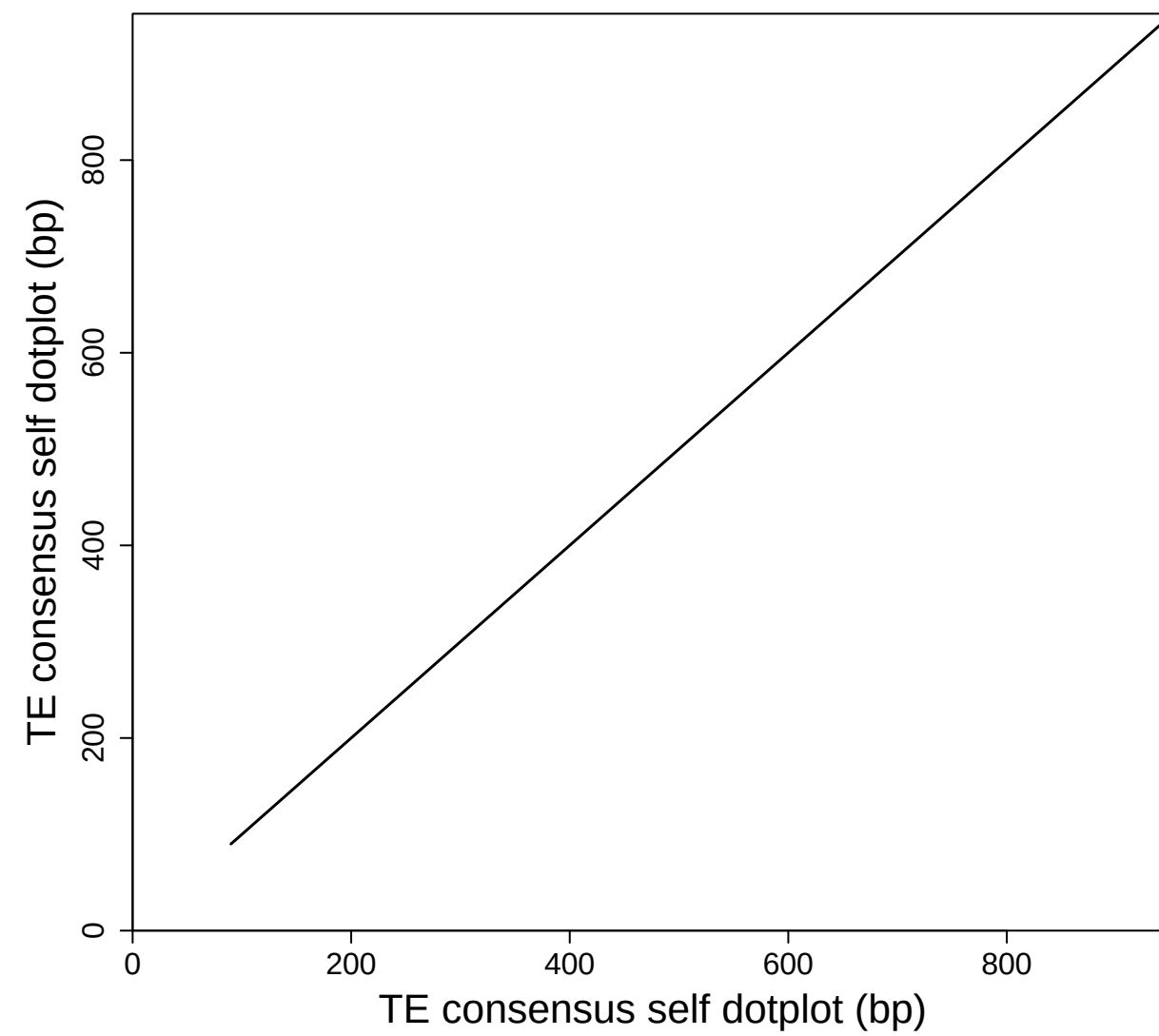
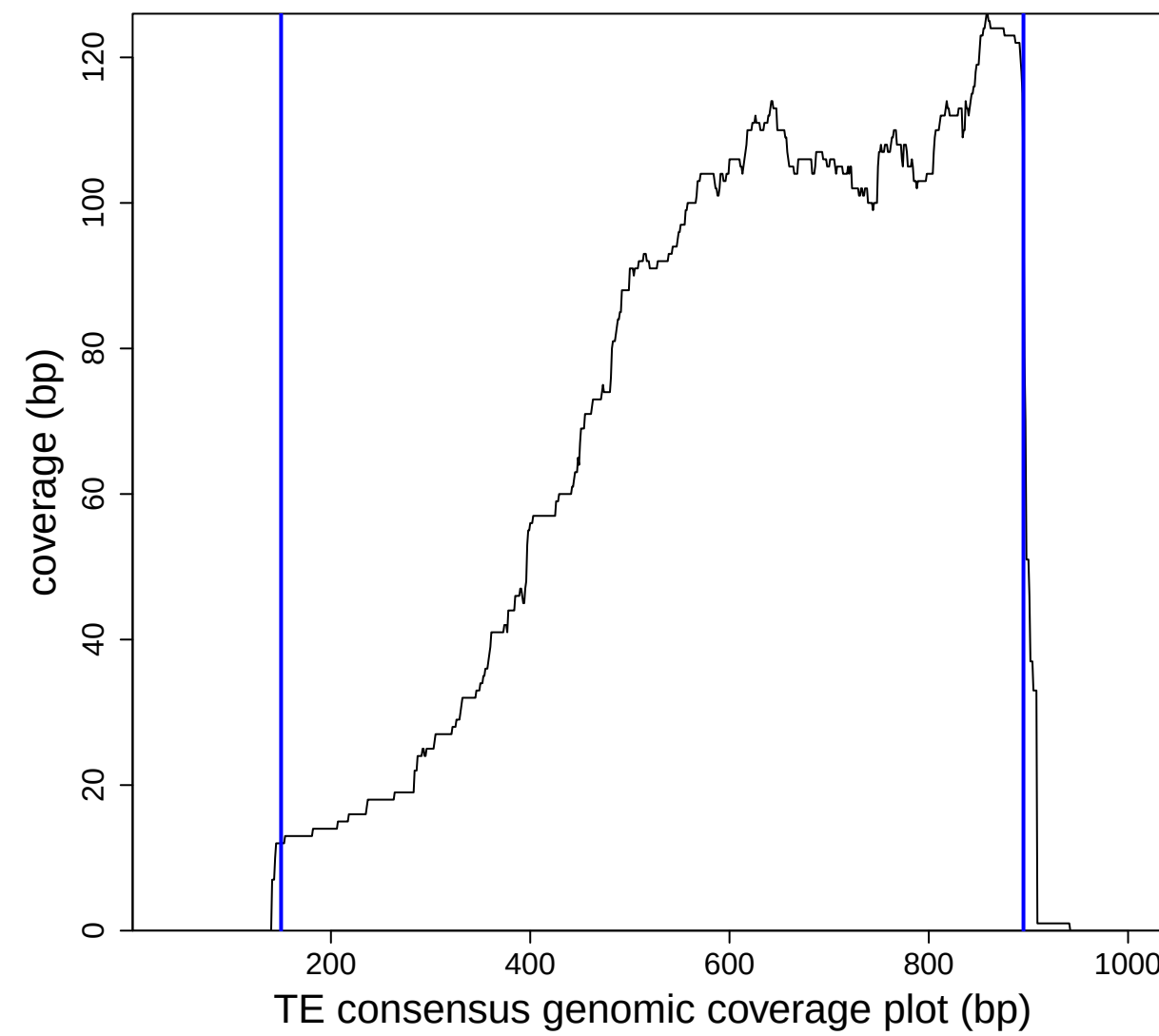
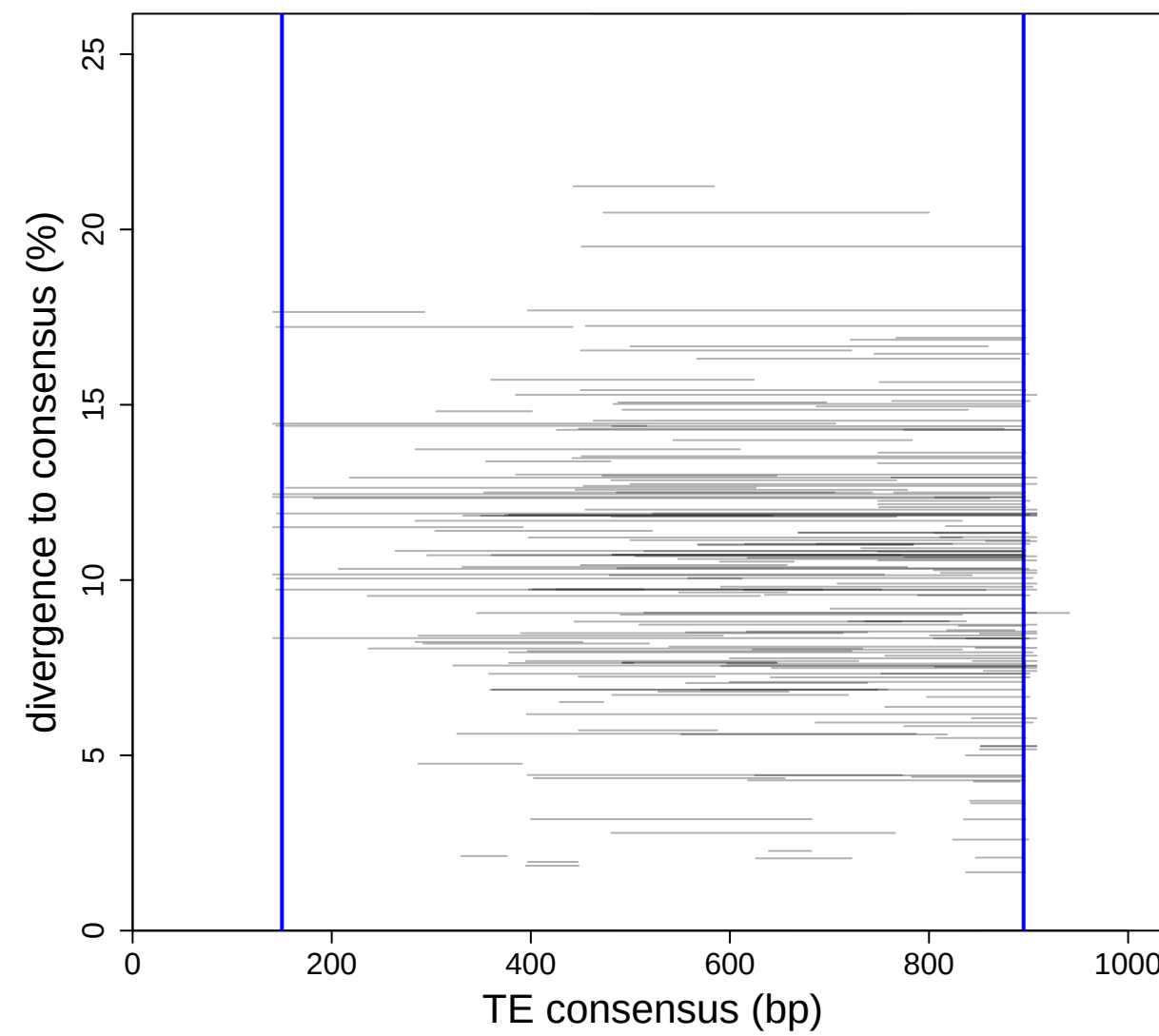
g_2.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcln.fa_ε
size: 735bp; fragments: 221; full length: 8 (>=661.5bp)

After TEtrimmer 735 bp

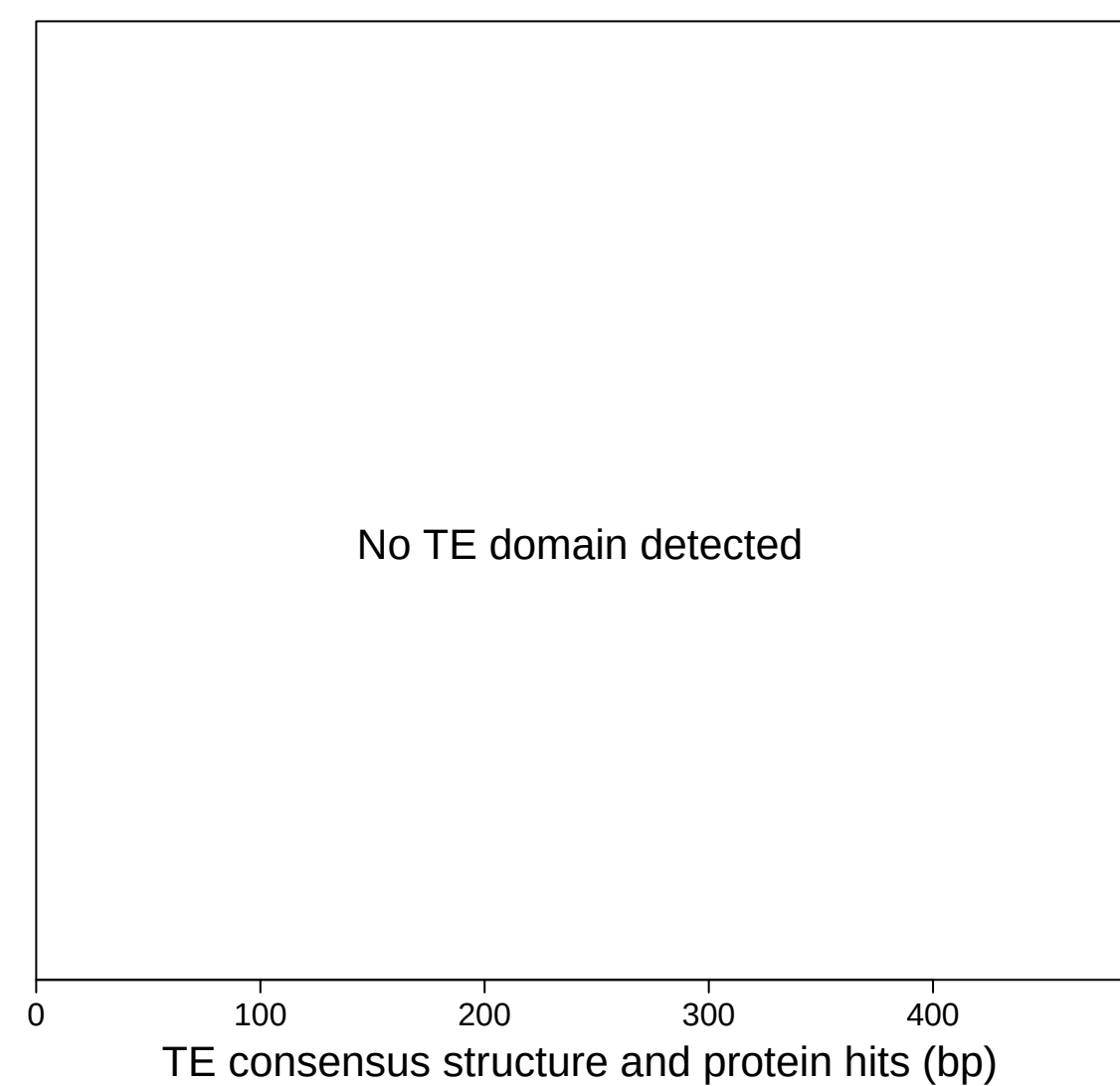
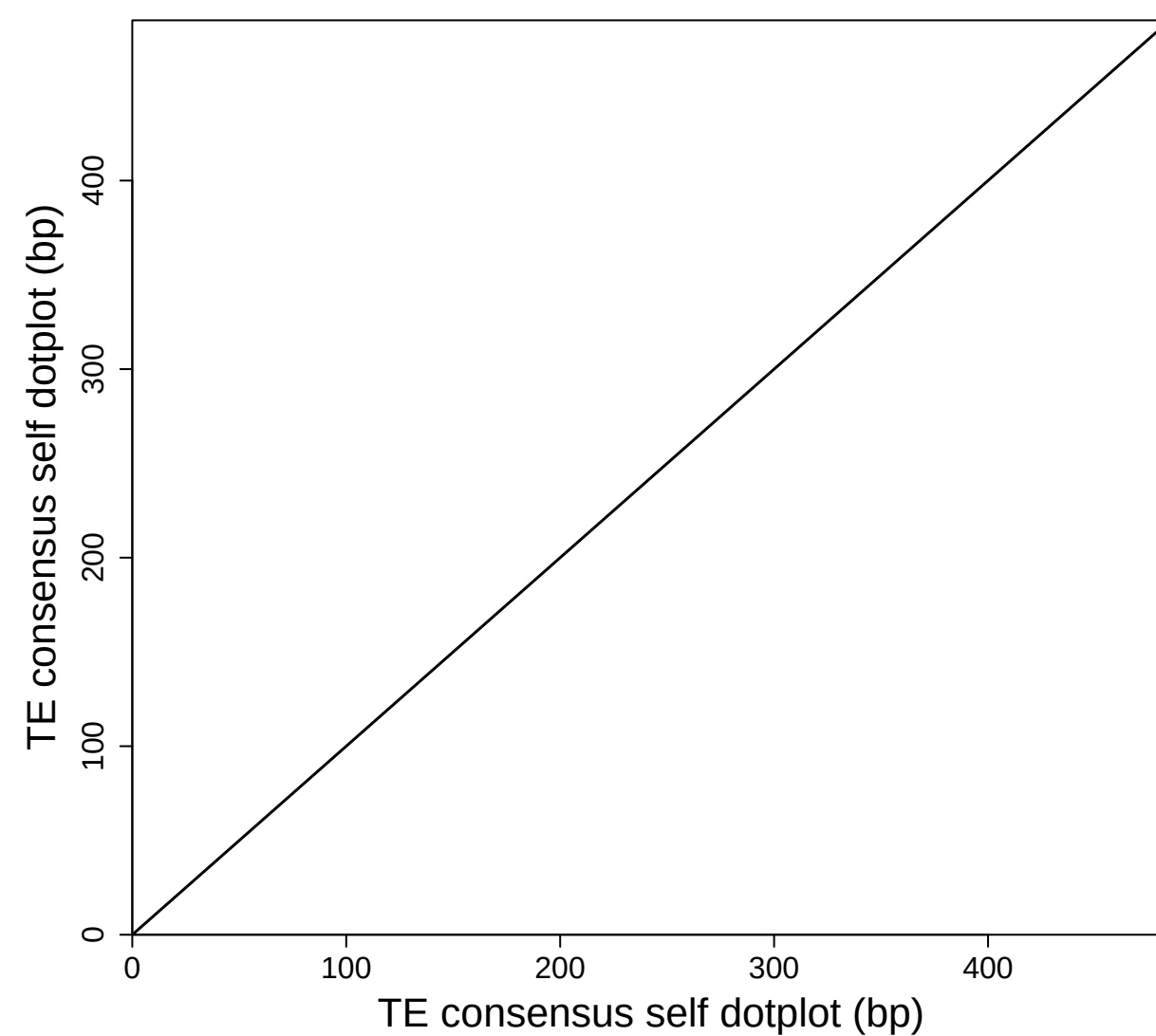
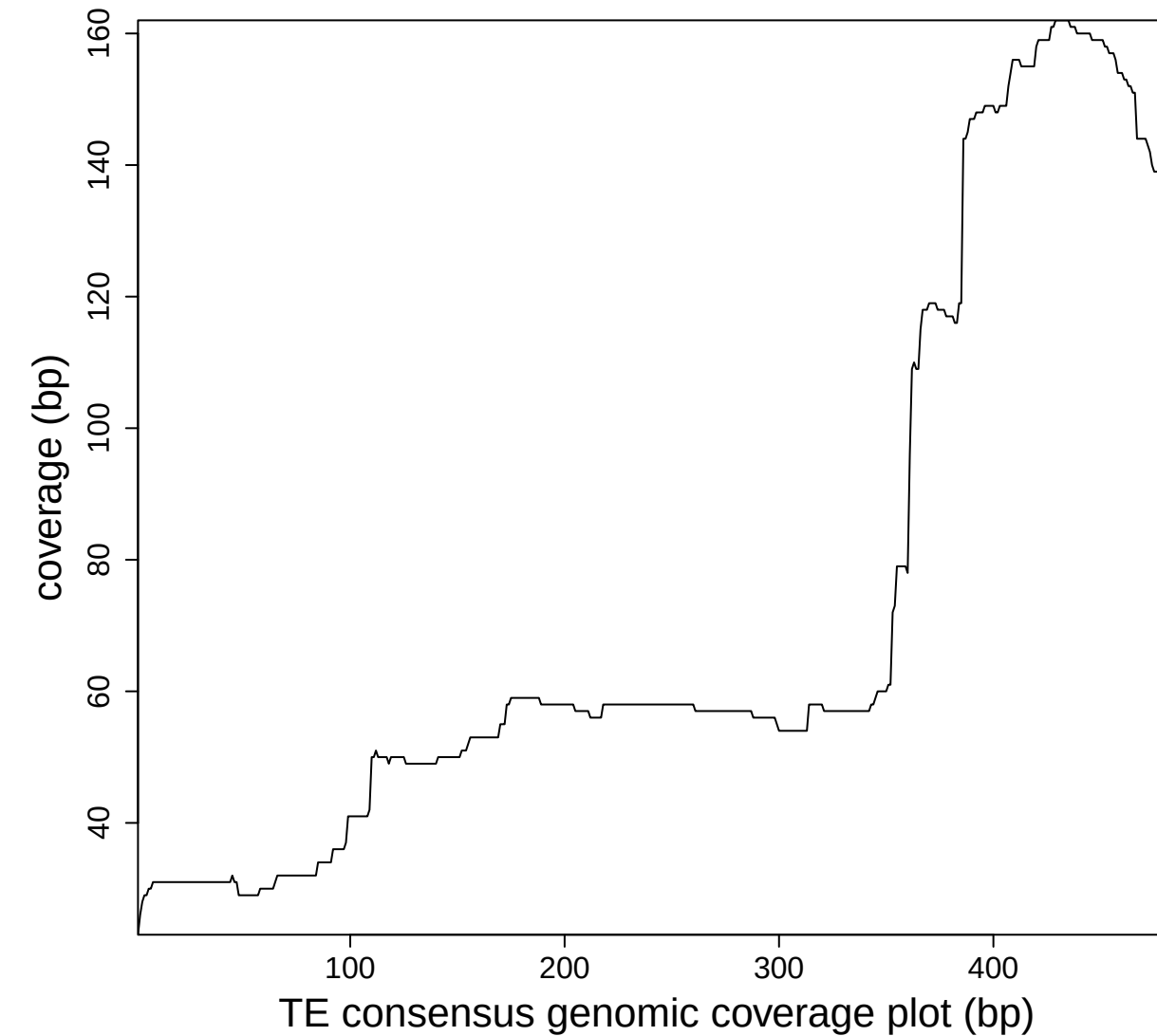
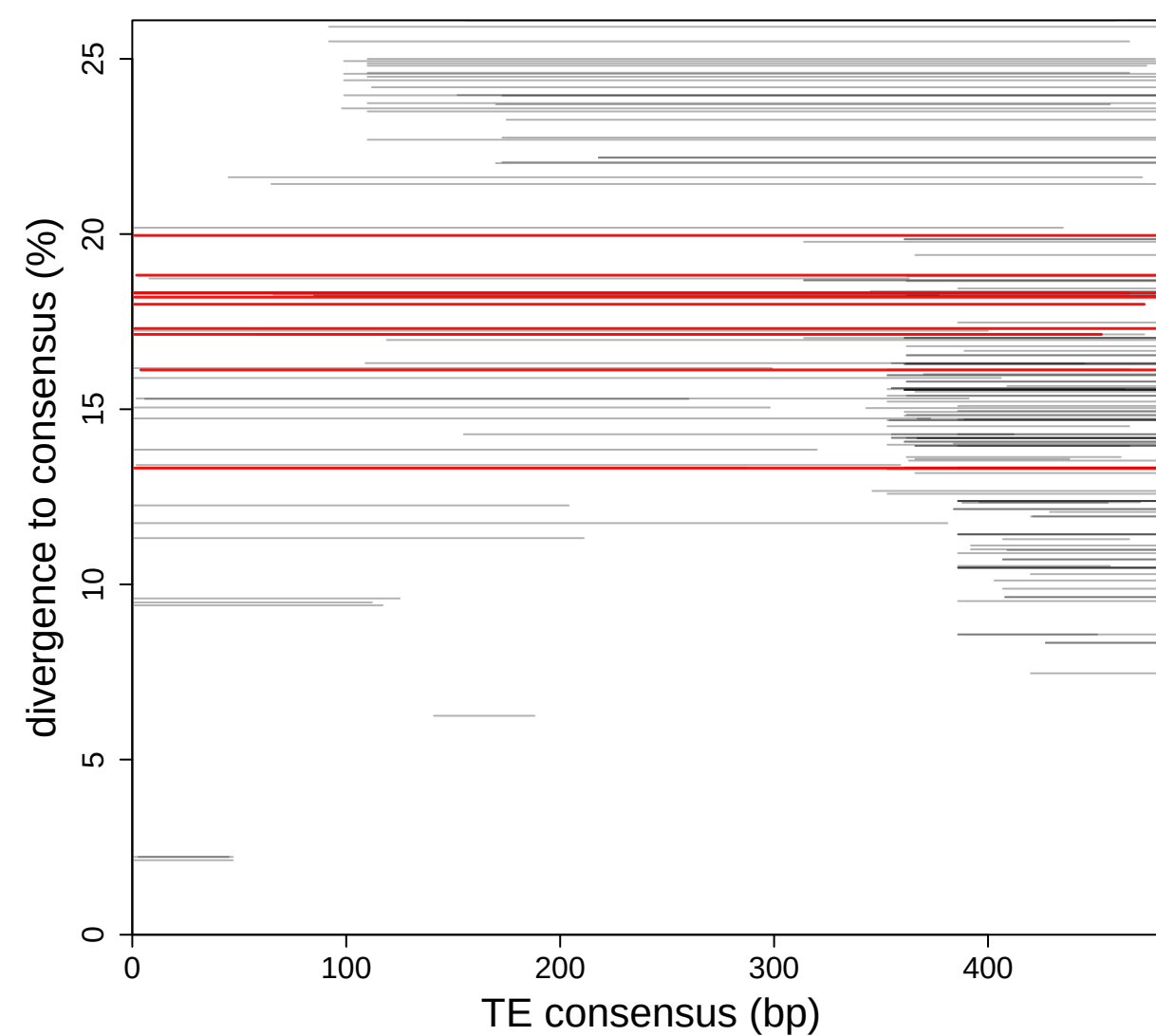


f.bed_g_2.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bc
size: 1045bp; fragments: 207; full length: 0 (>=940.5bp)

After TEtrimmer Extended plot Blue lines are boundaries

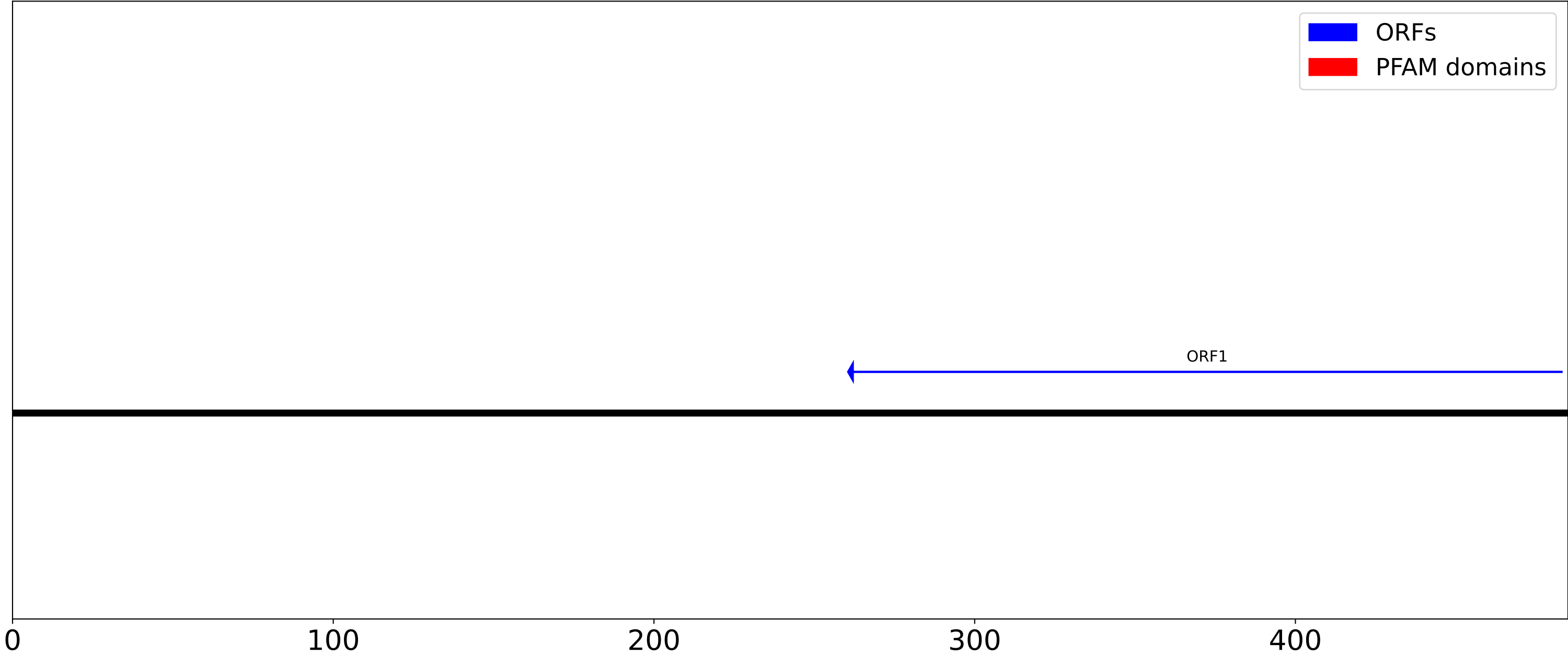


TE: rnd_5_family_4123
size: 485bp; fragments: 185; full length: 10 (≥ 436.5 bp)



Before TEtrimmer 485 bp

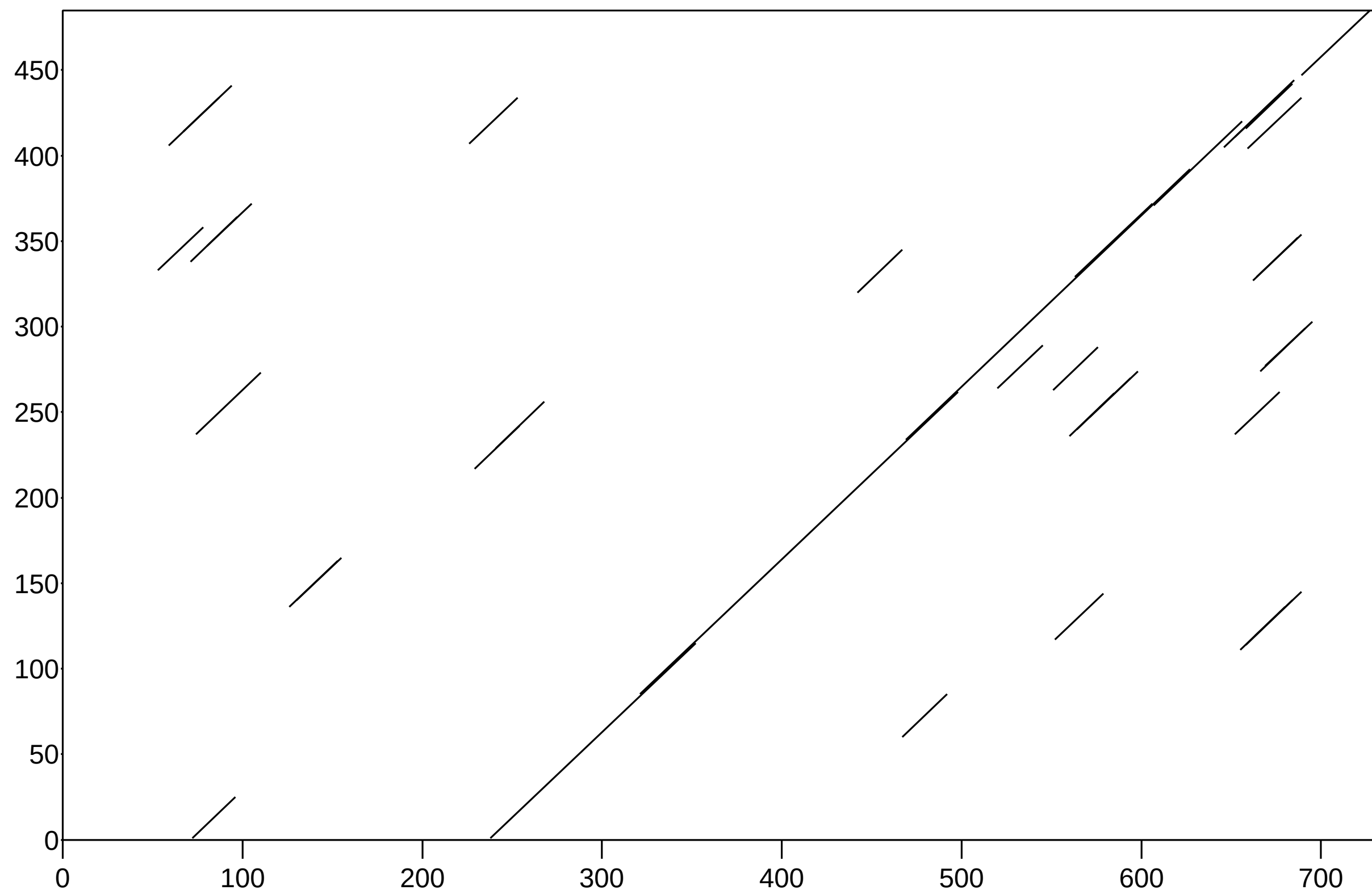
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)