

Start crop Point



End crop Point



0-

10

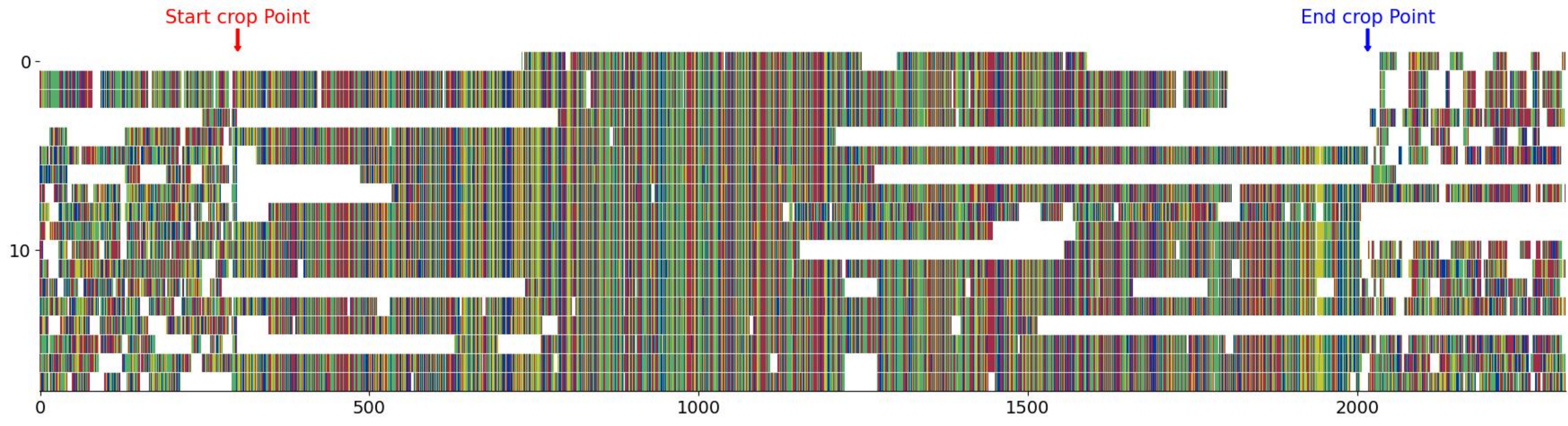
0

500

1000

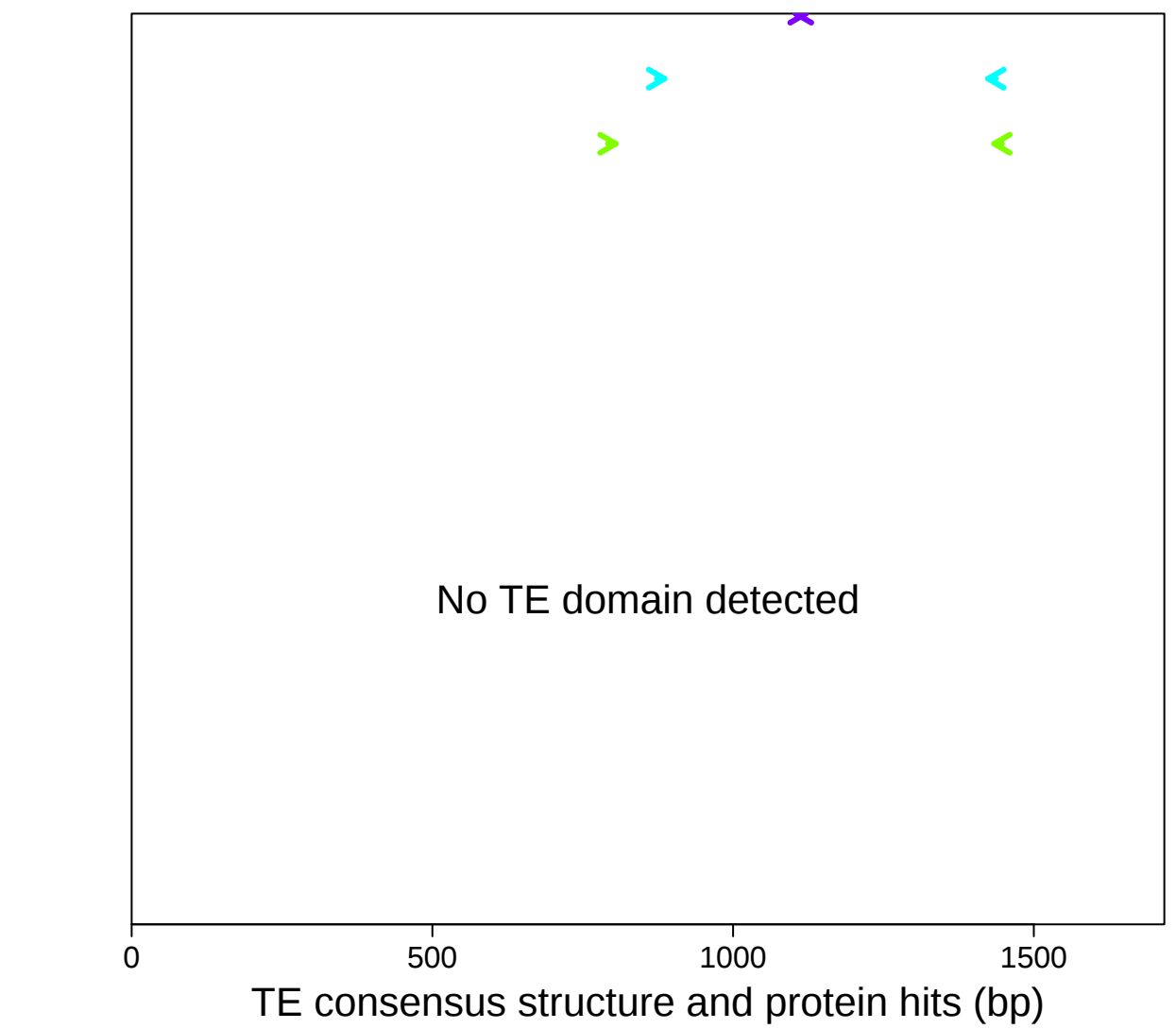
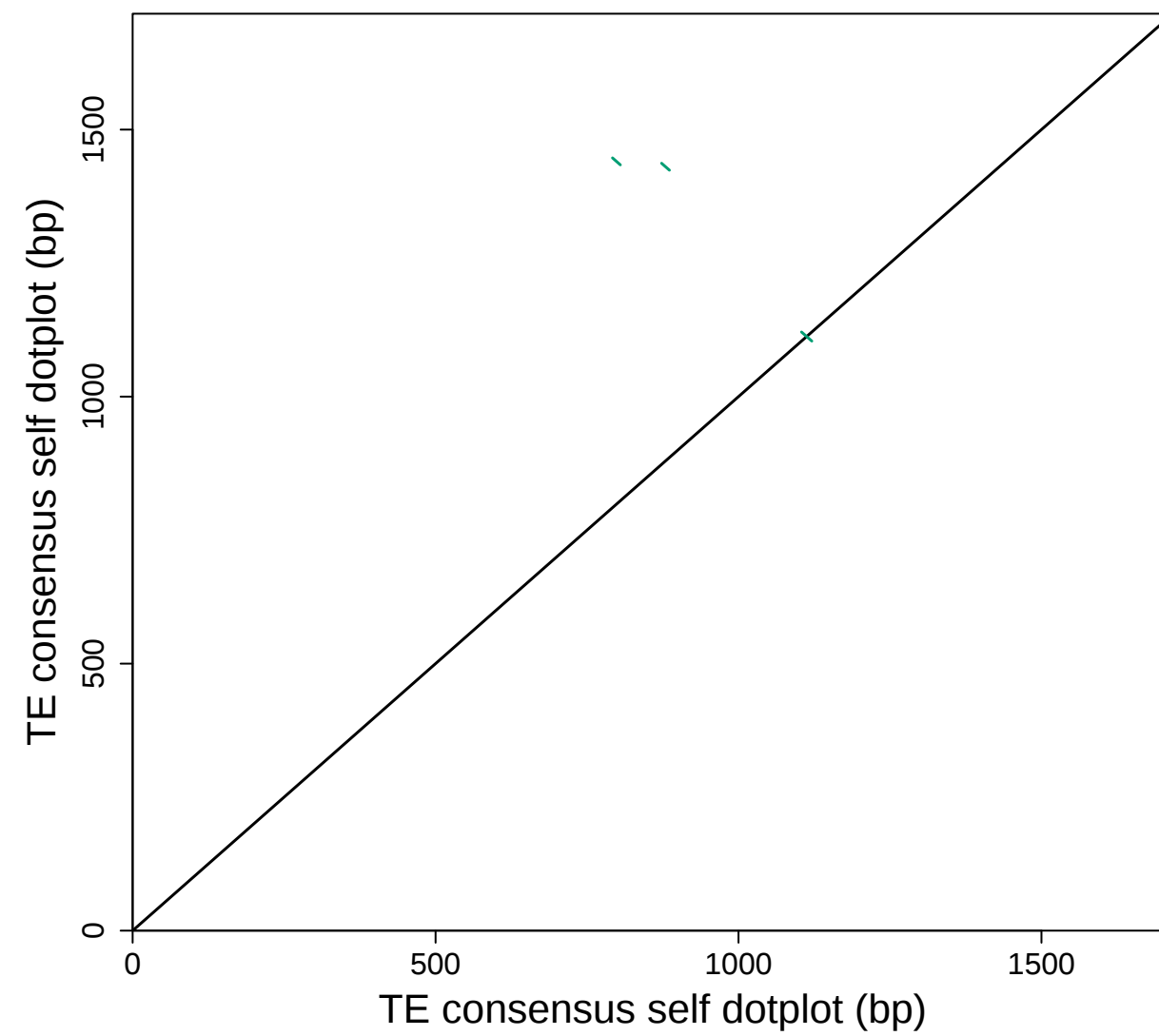
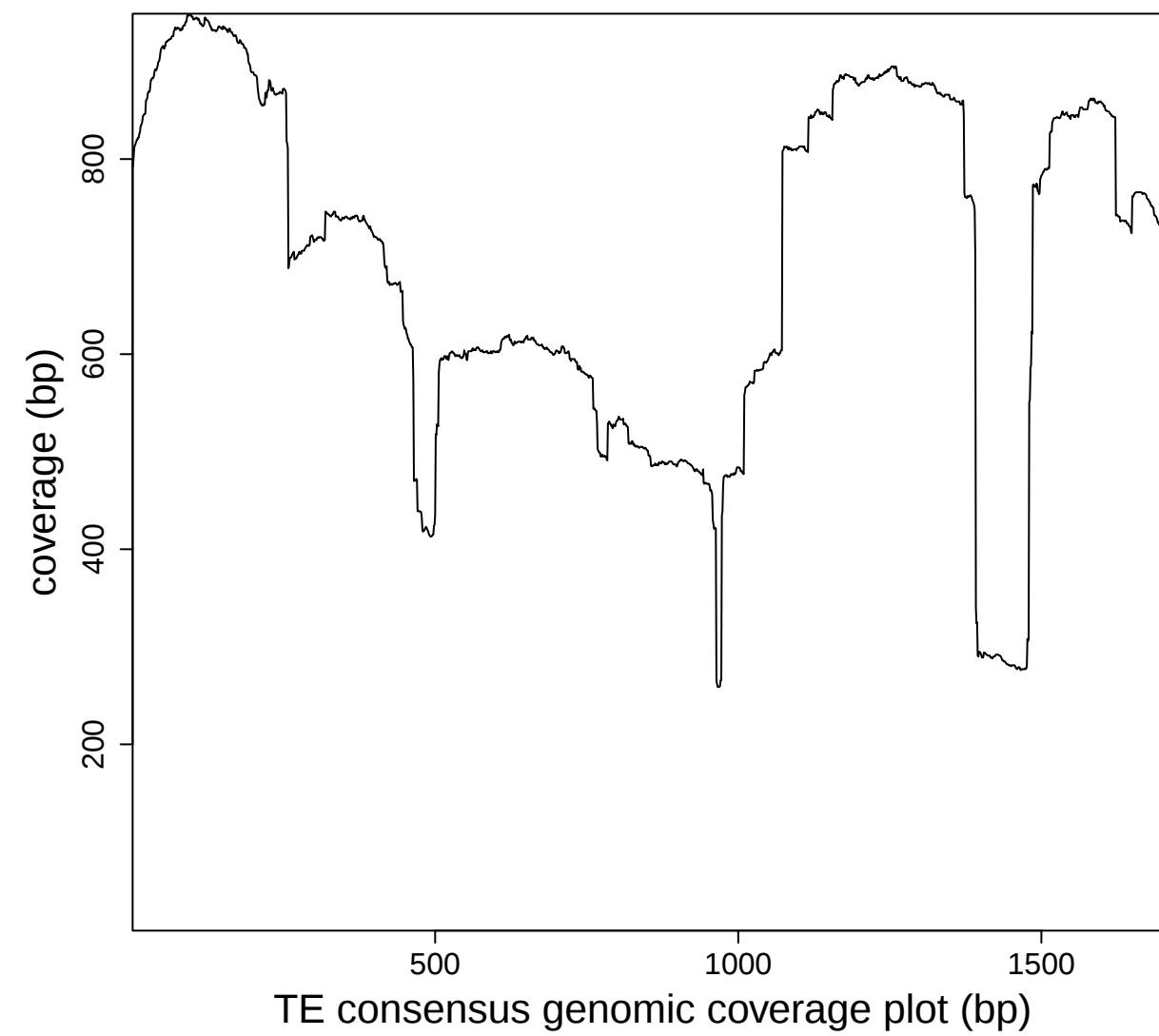
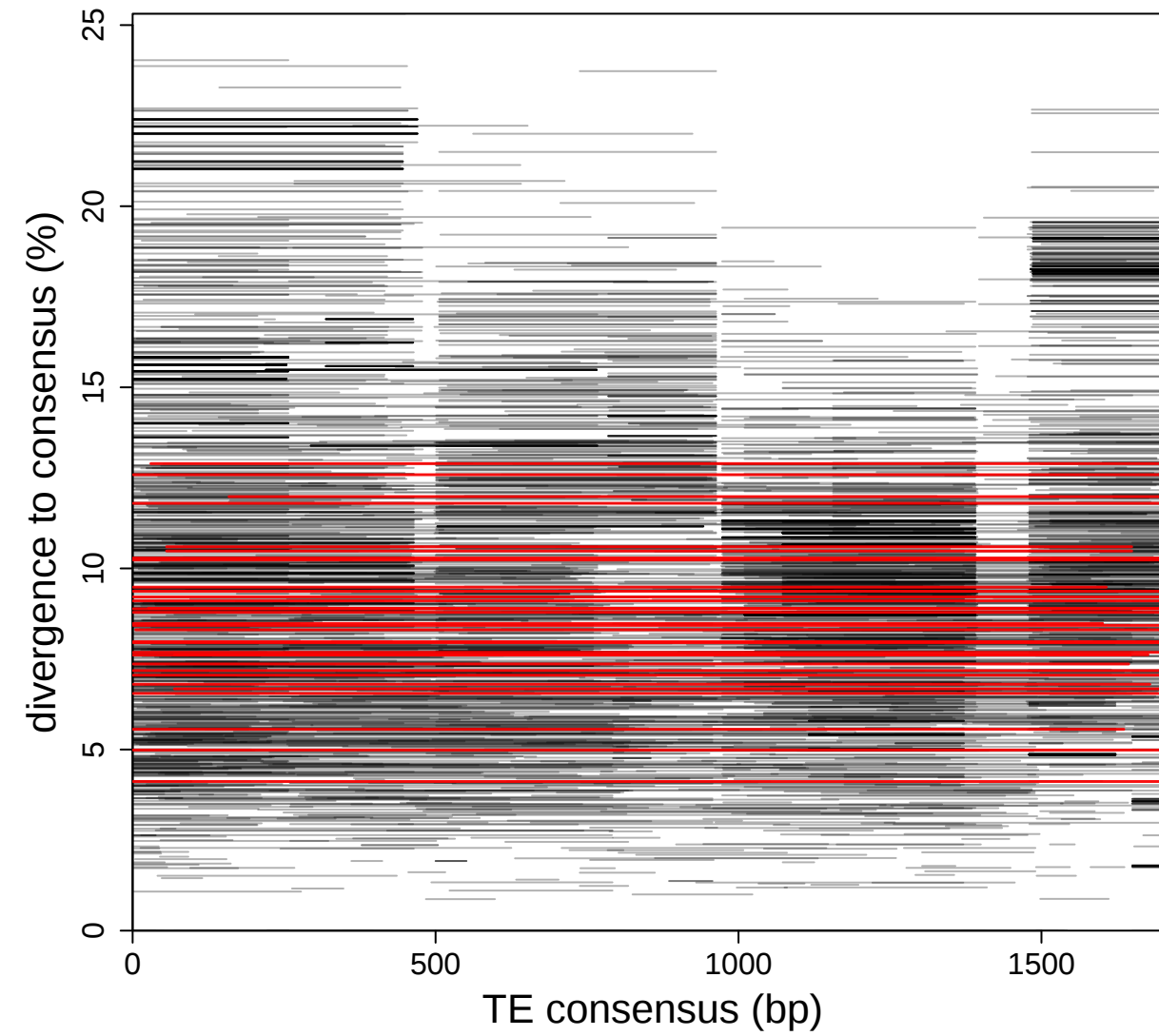
1500

2000



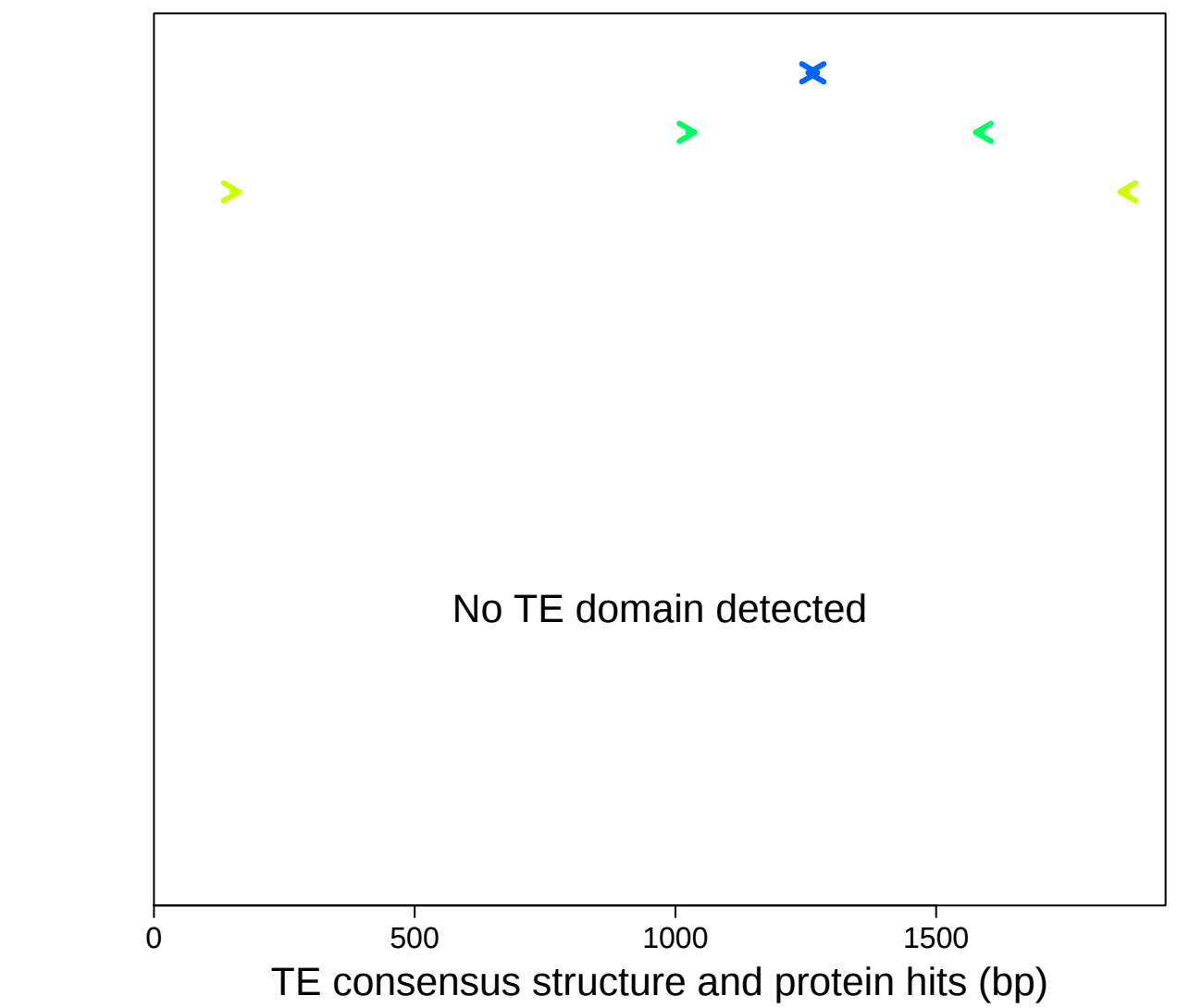
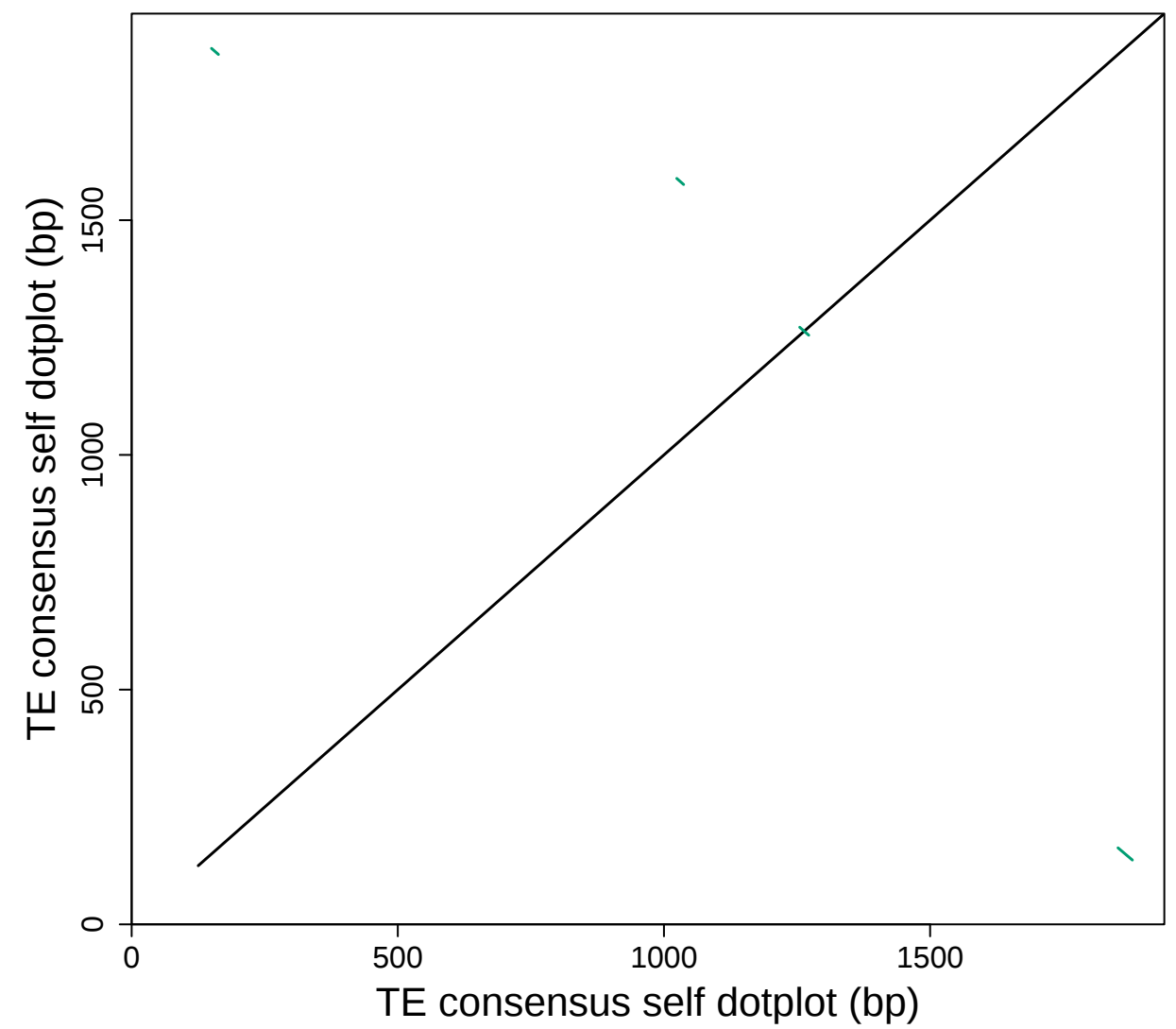
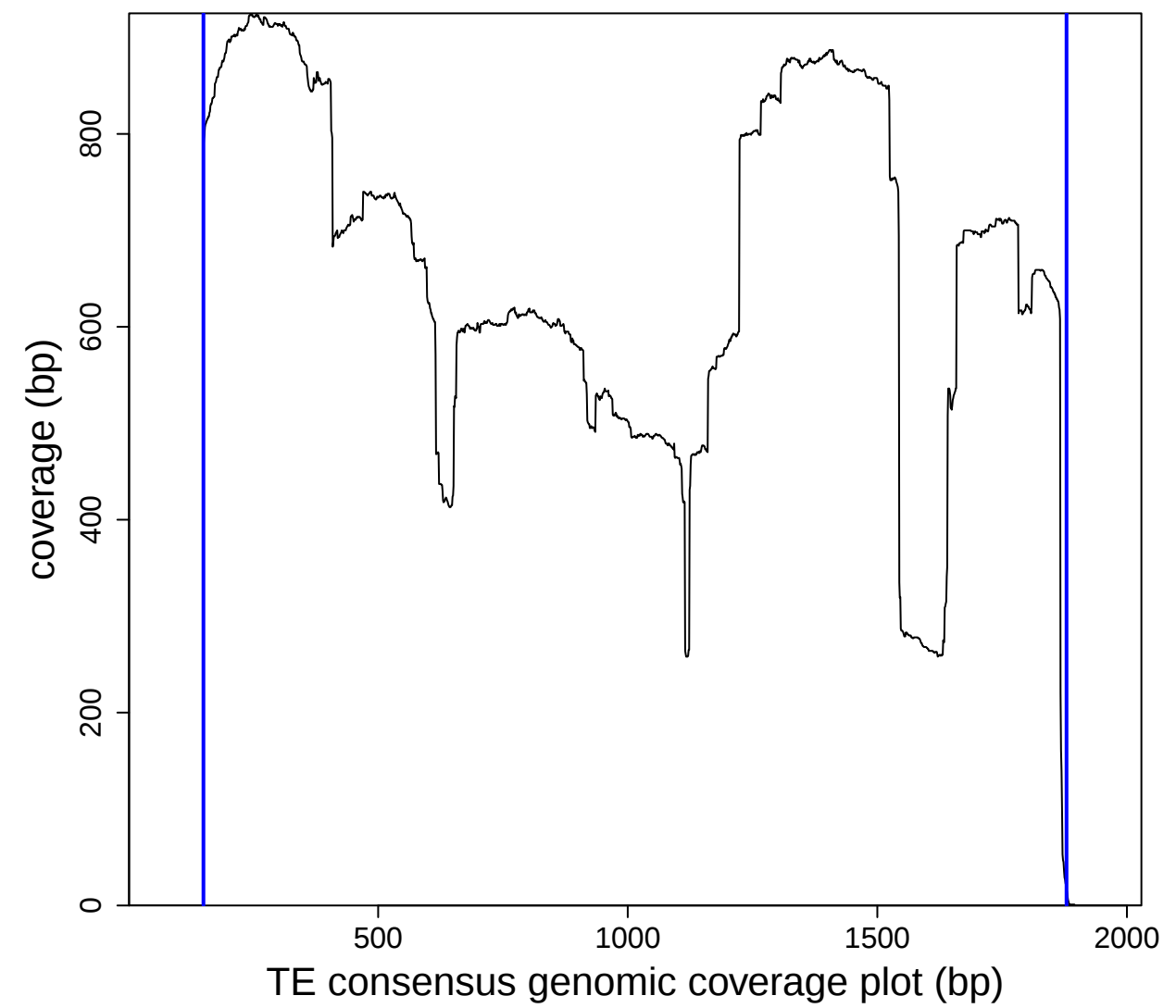
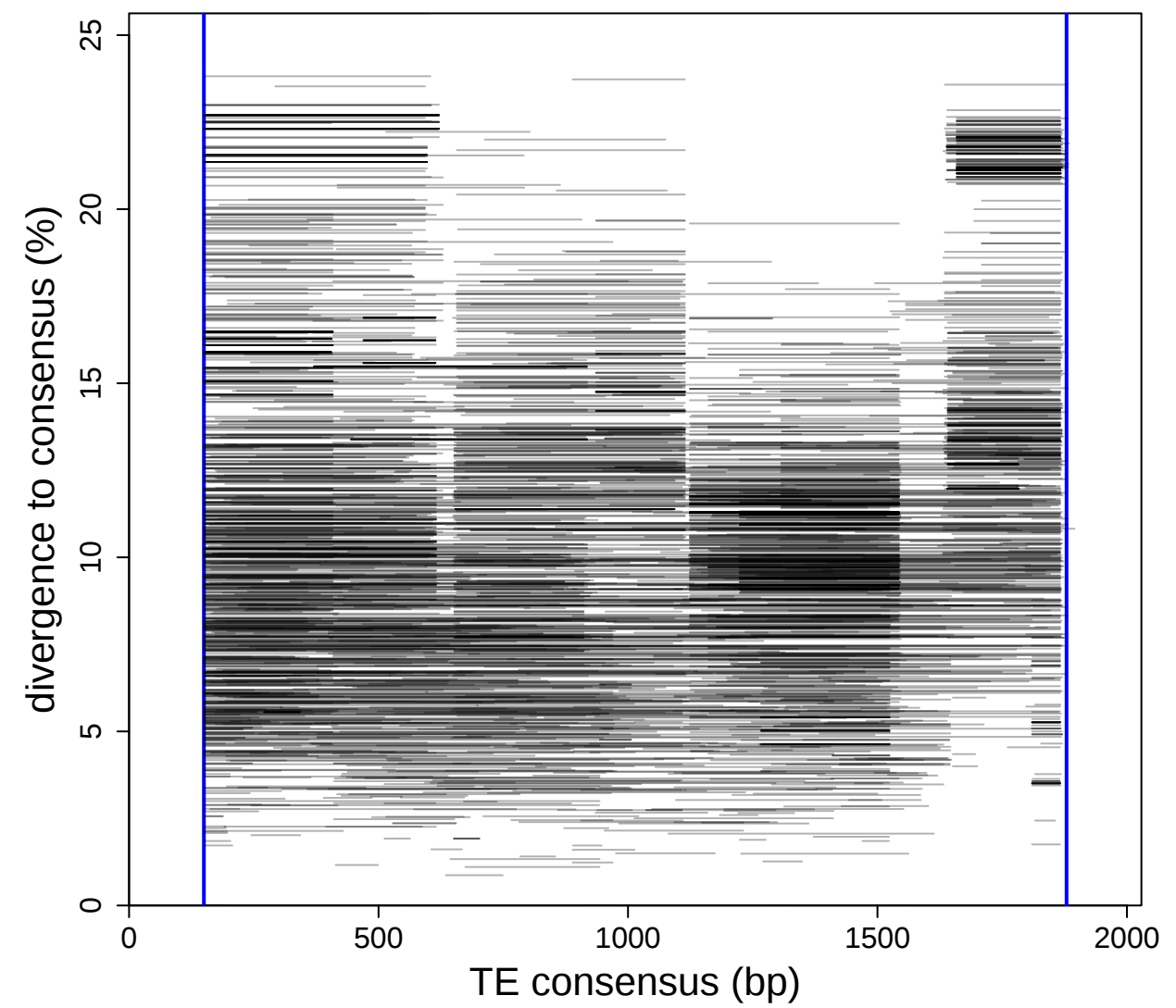
g_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcln.fa_a
size: 1717bp; fragments: 3409; full length: 32 (>=1545.3bp)

After TEtrimmer 1717 bp

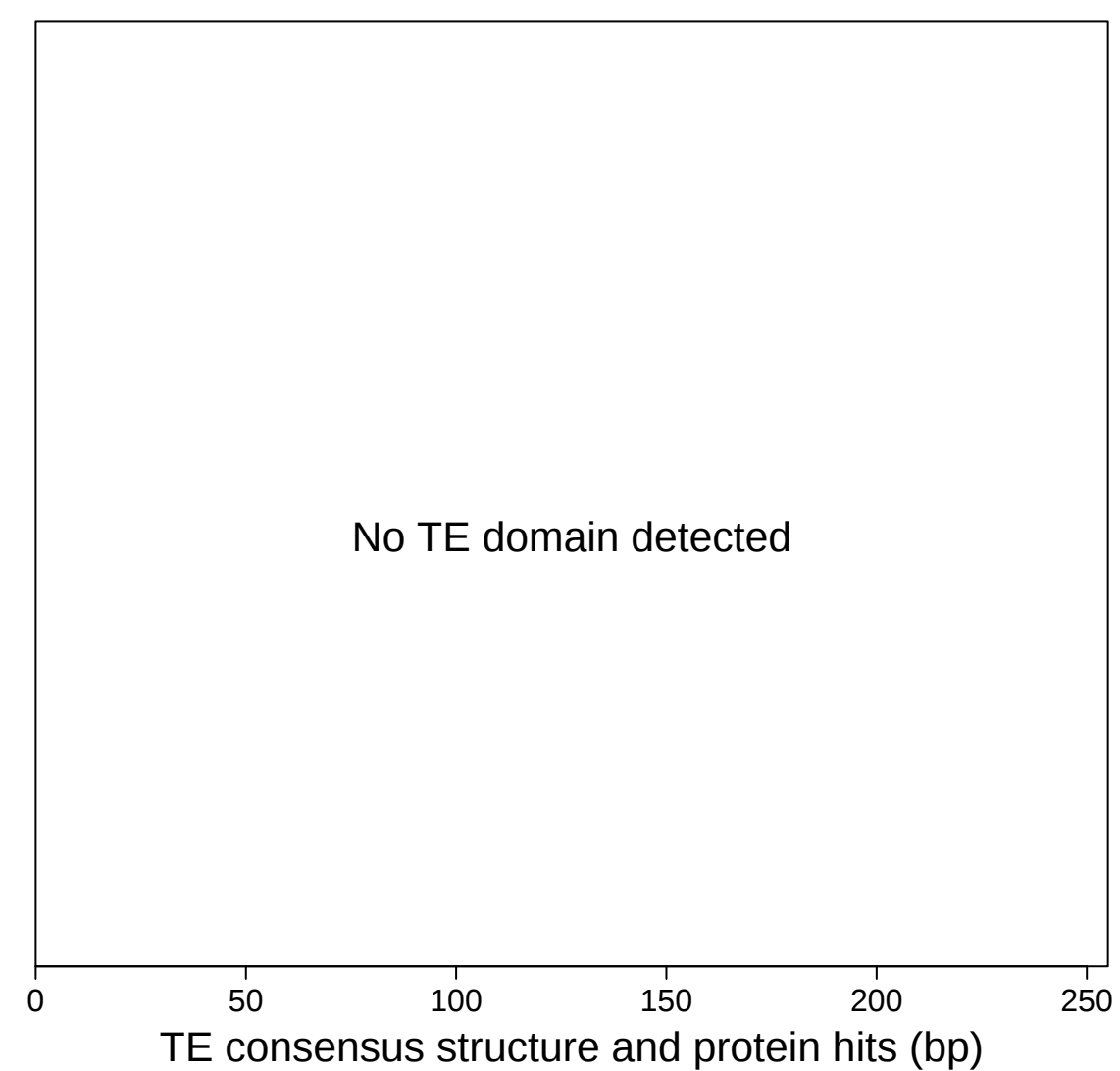
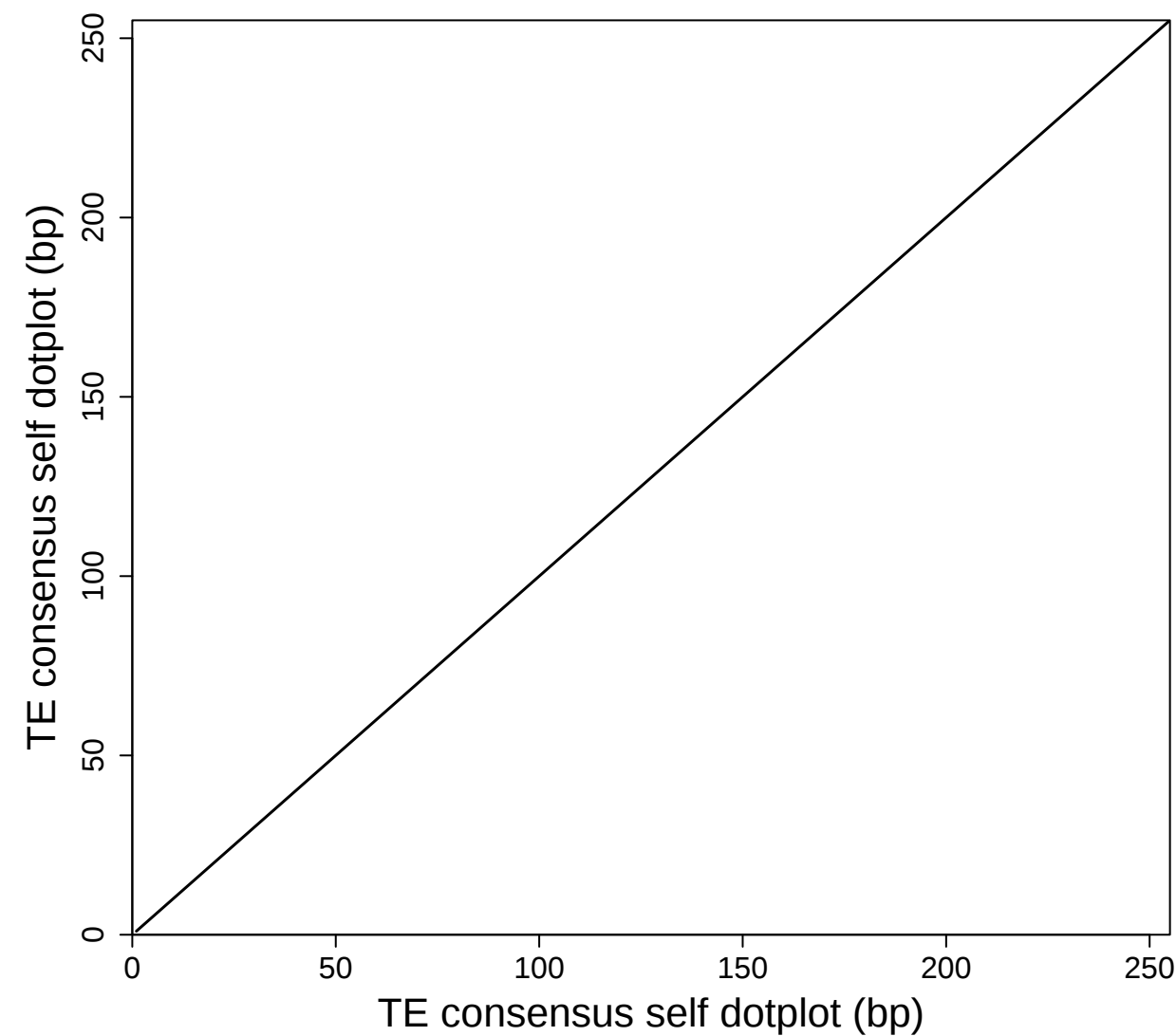
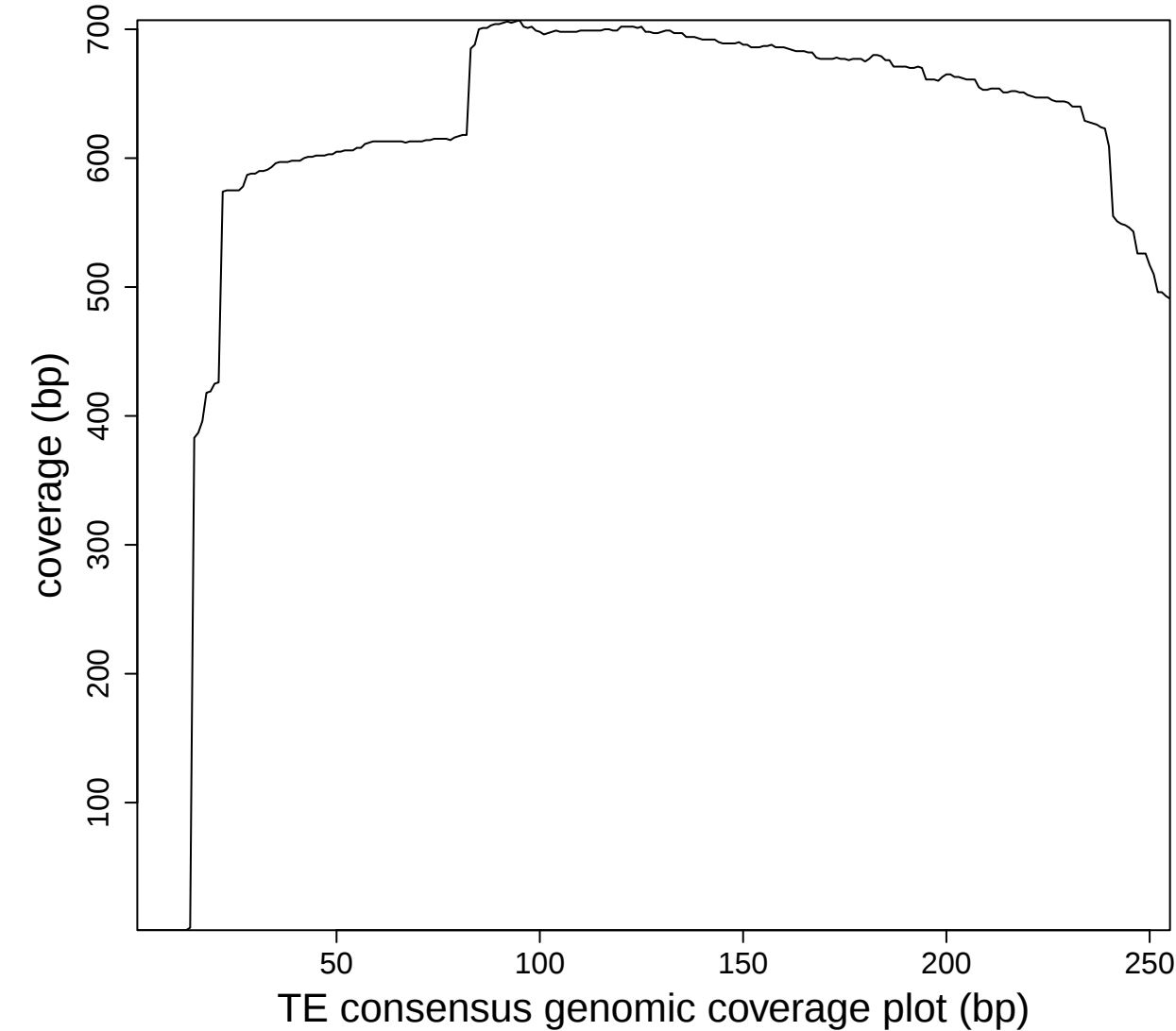
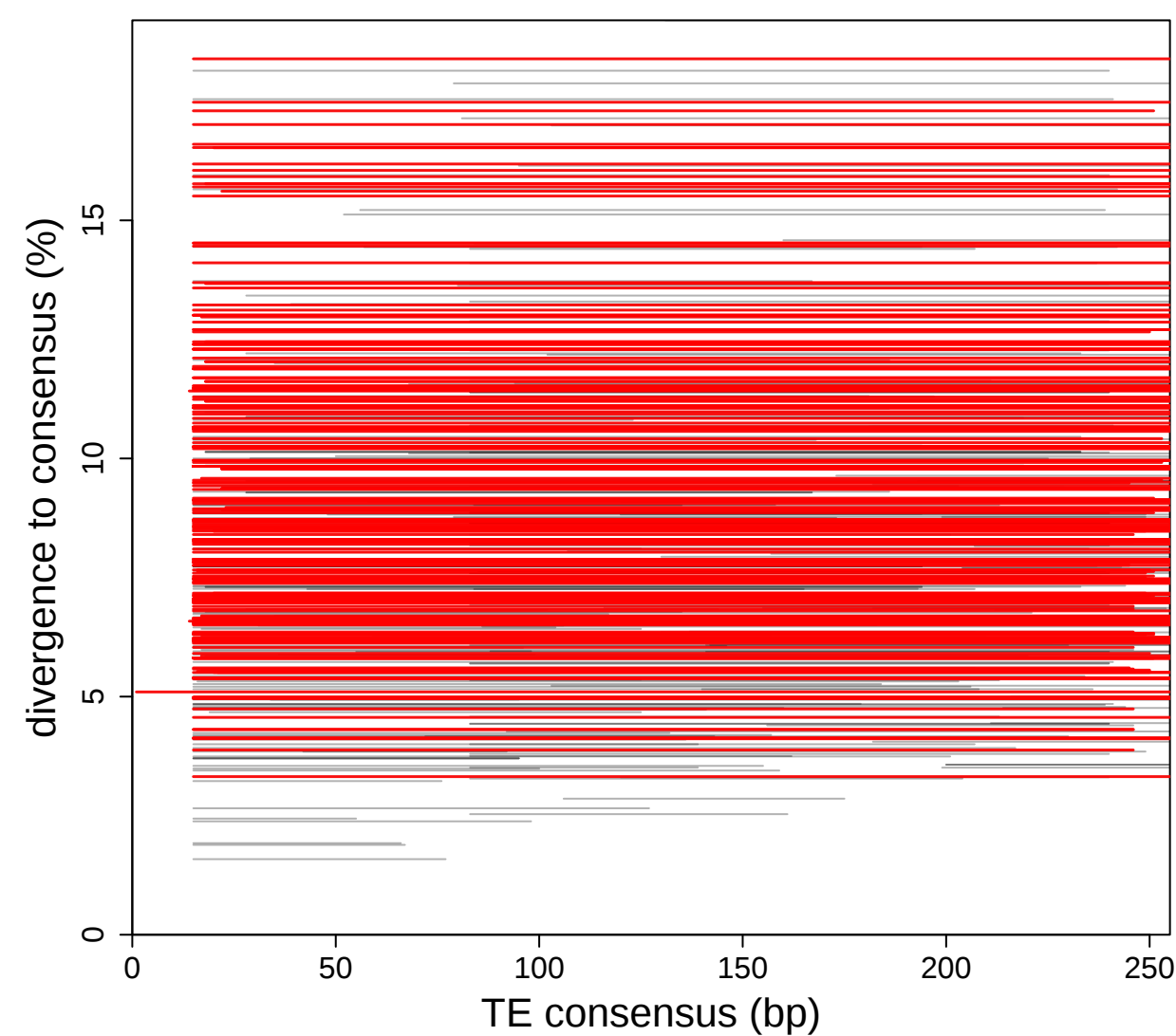


f.bed_g_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcl
size: 2029bp; fragments: 3178; full length: 0 (>=1826.1bp)

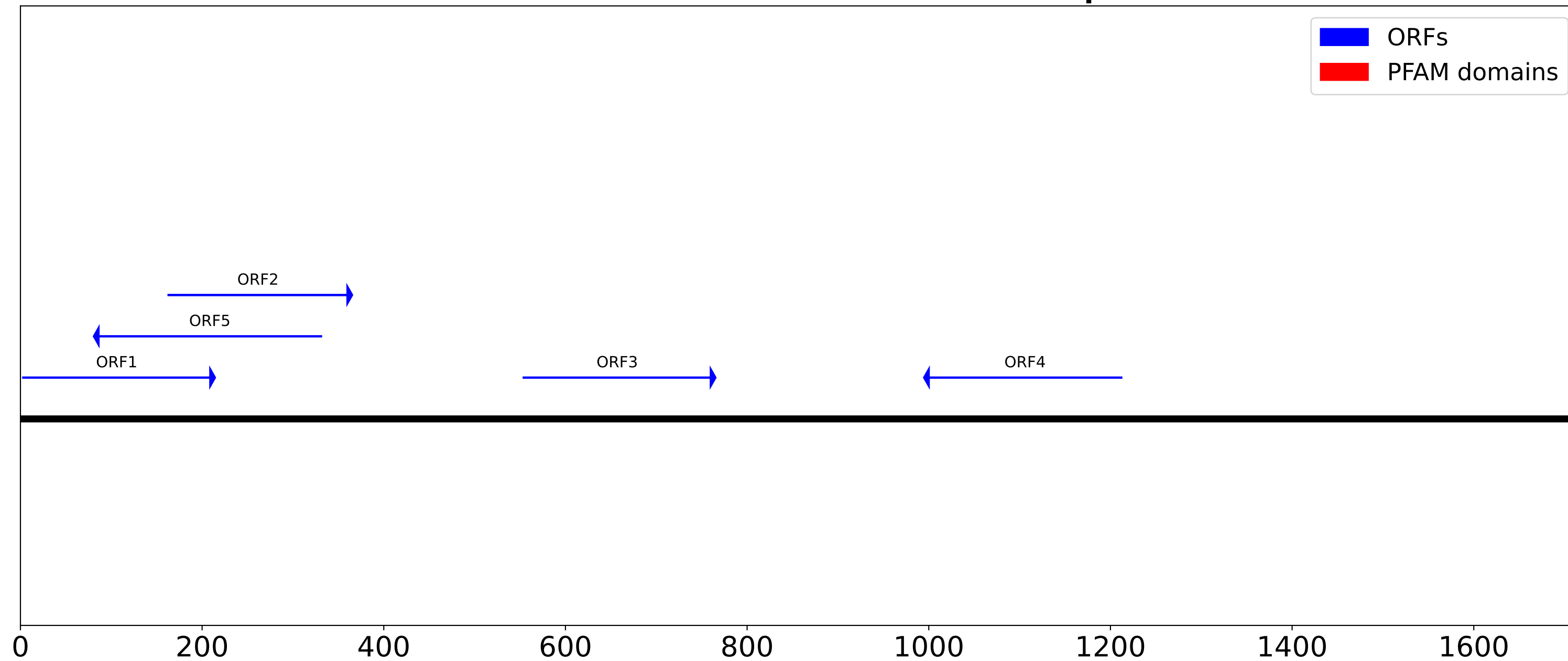
After TEtrimmer Extended plot Blue lines are boundaries



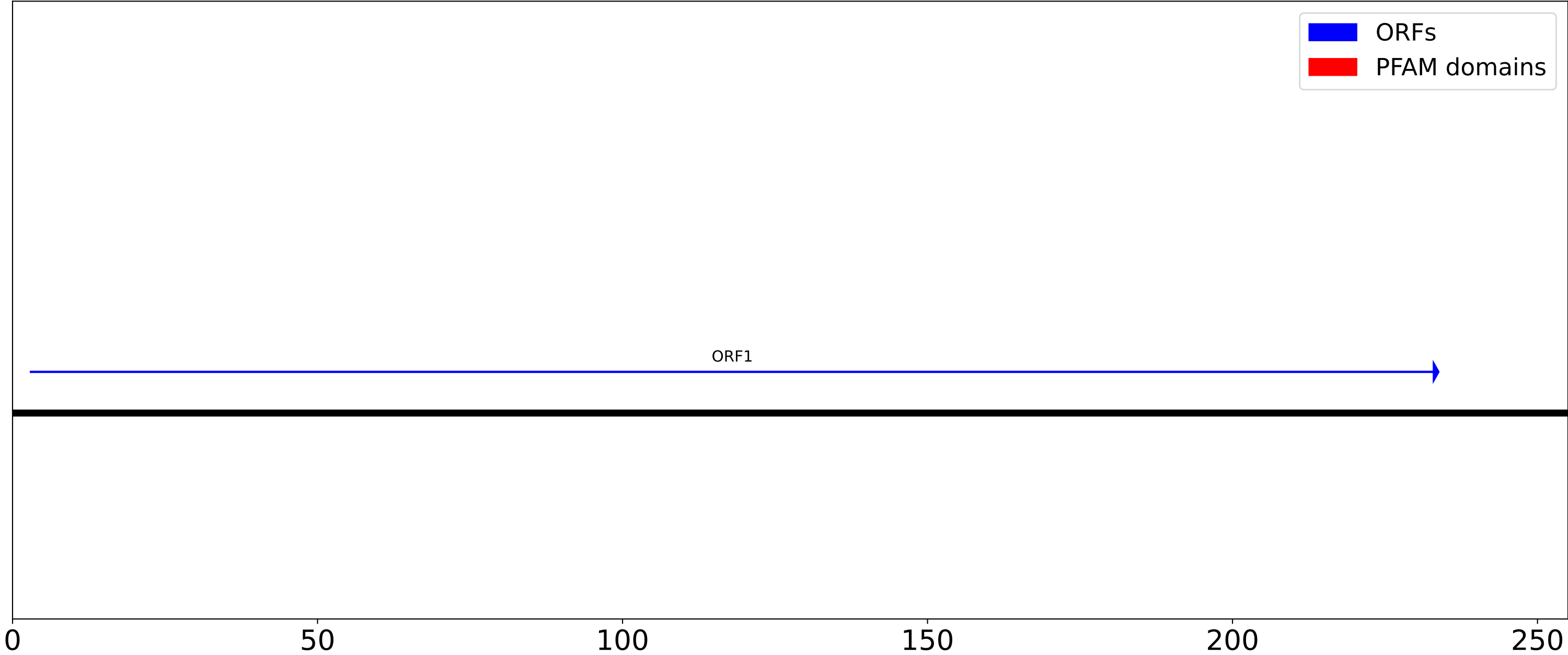
TE: rnd_1_family_797
size: 255bp; fragments: 764; full length: 452 (≥ 229.5 bp)



After TEtrimmer ORF and PFAM domain plot



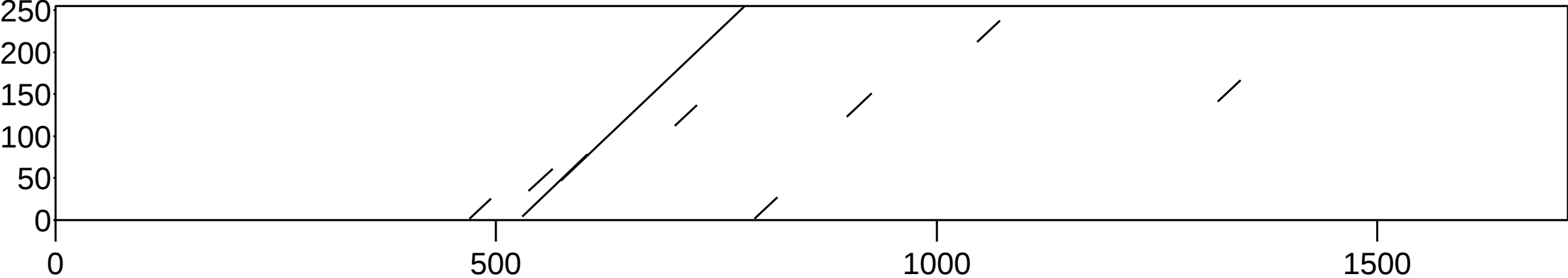
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

Isensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)