

Start crop Point



End crop Point



MSA length = 2060



Start crop Point

End crop Point

0 -

10 -

0

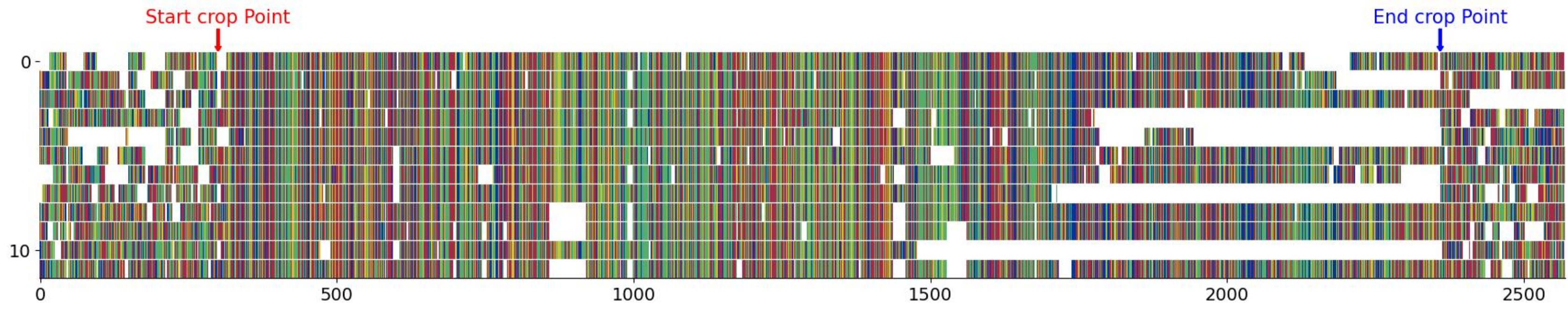
500

1000

1500

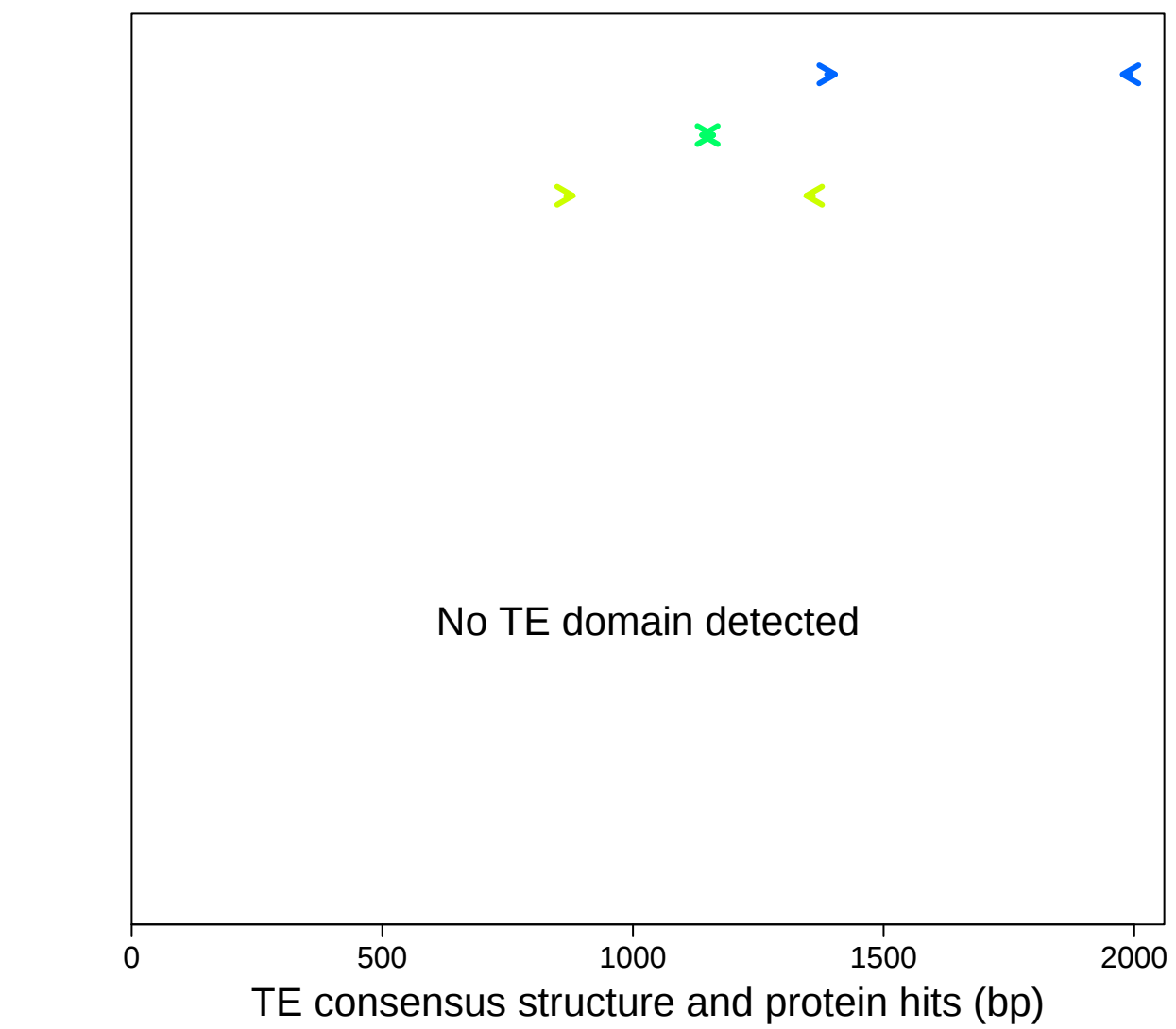
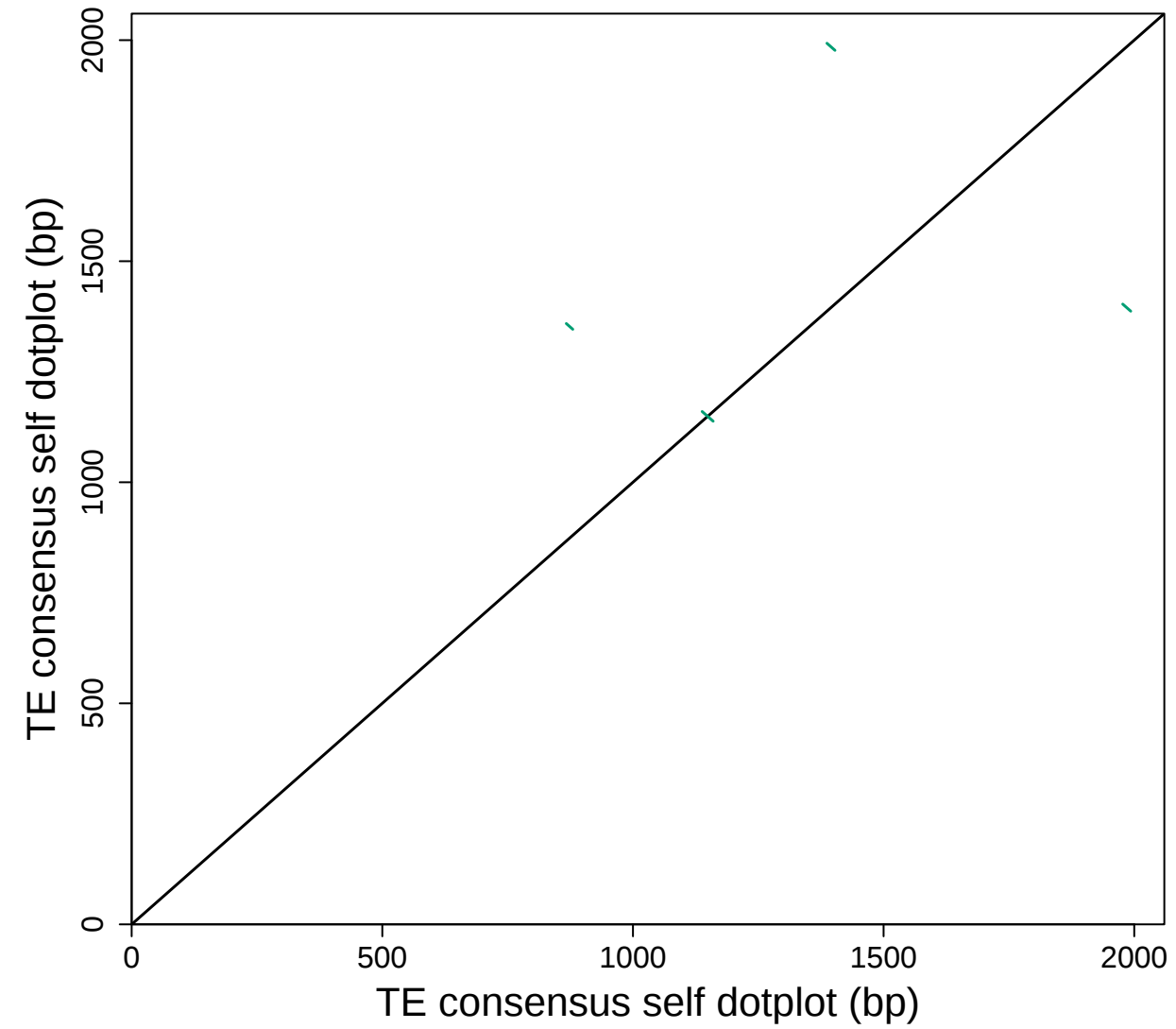
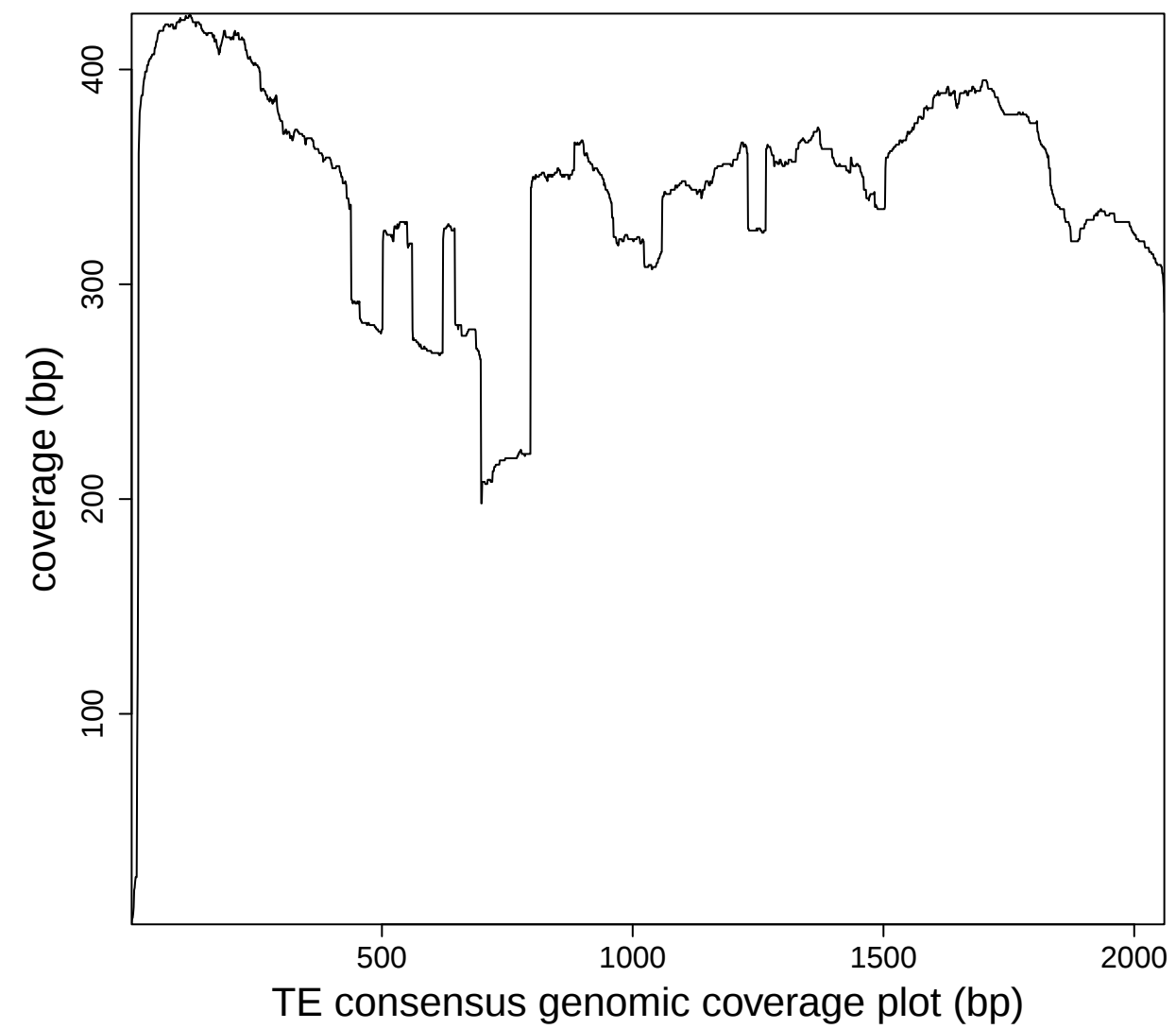
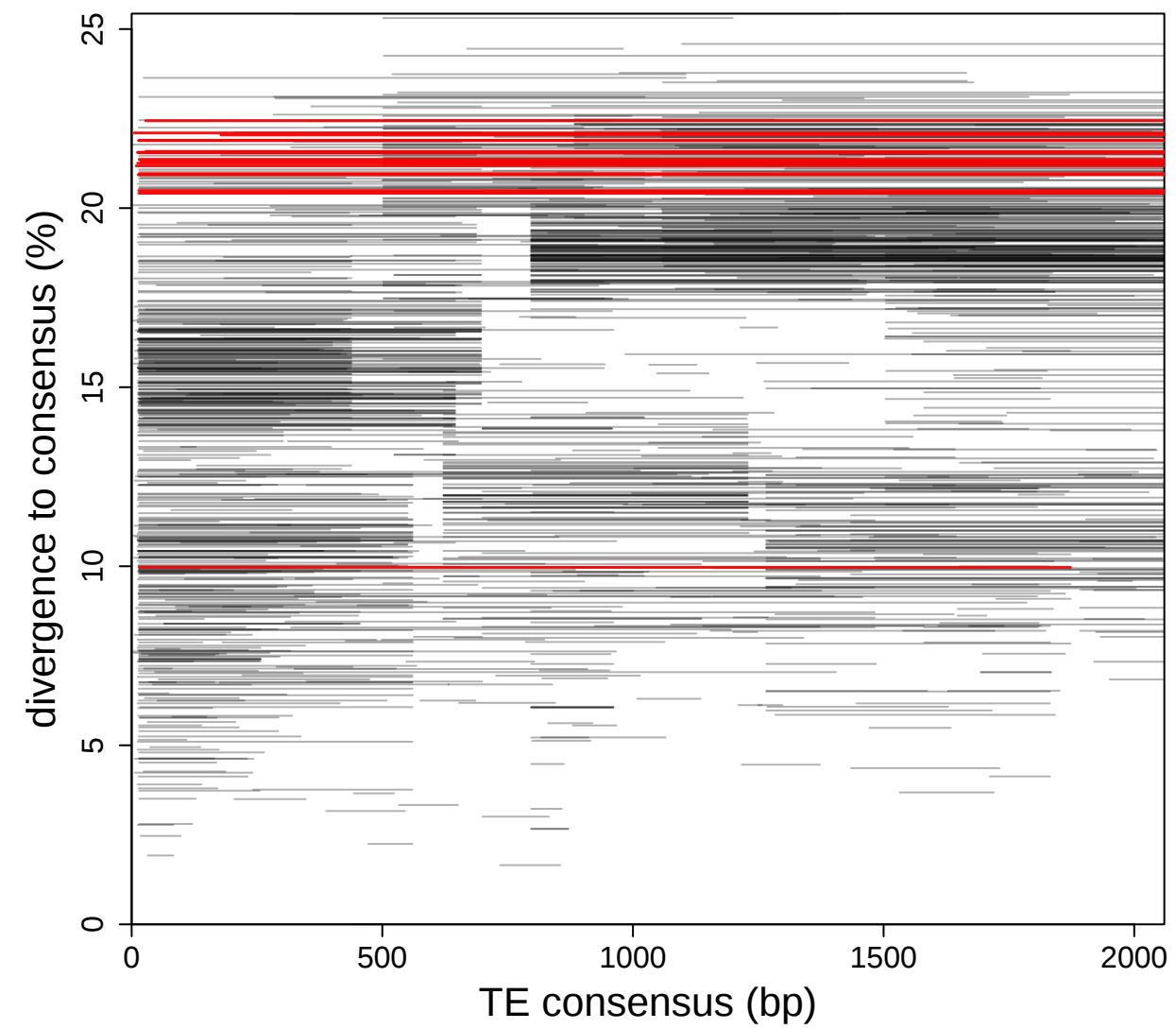
2000

2500



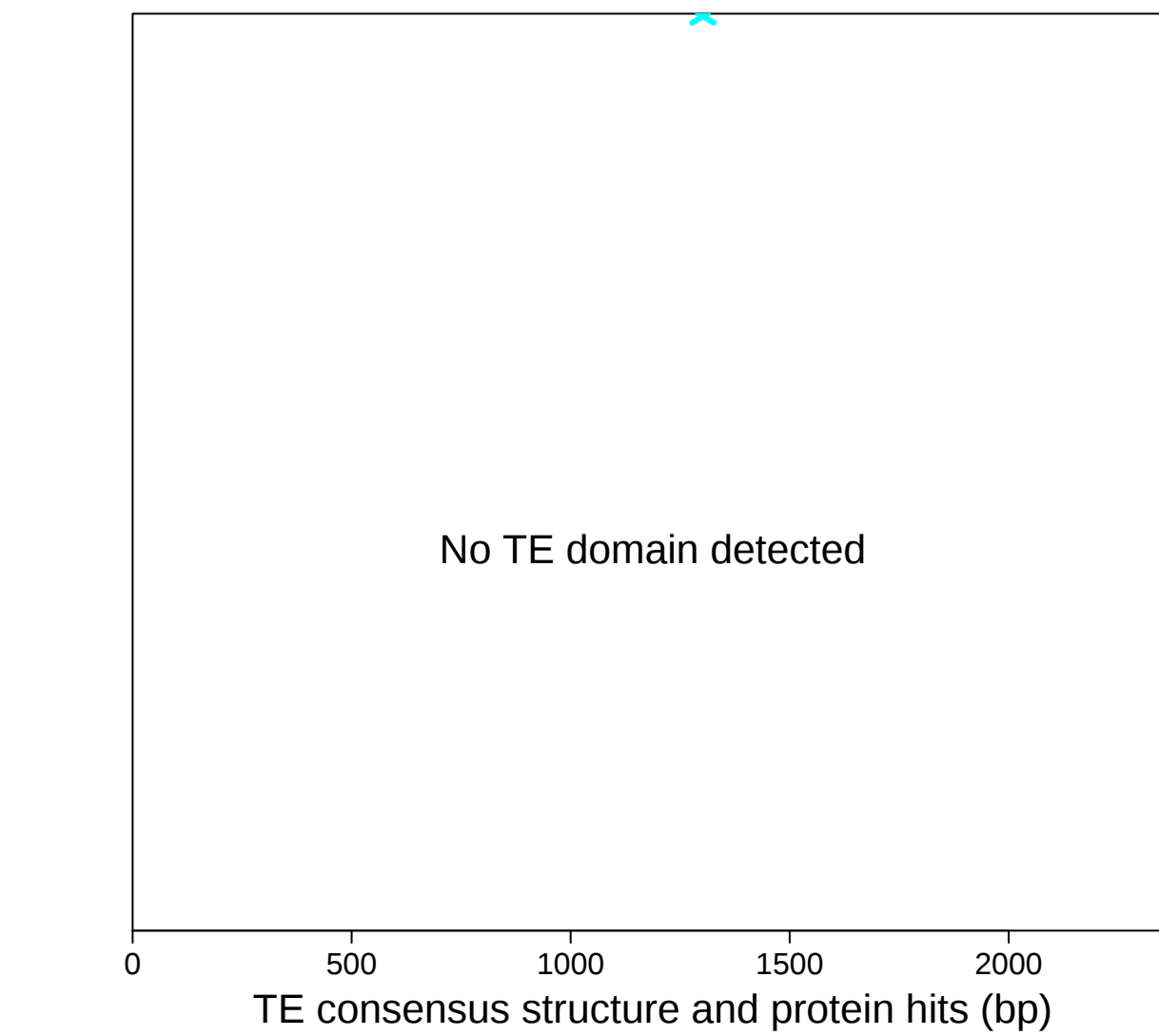
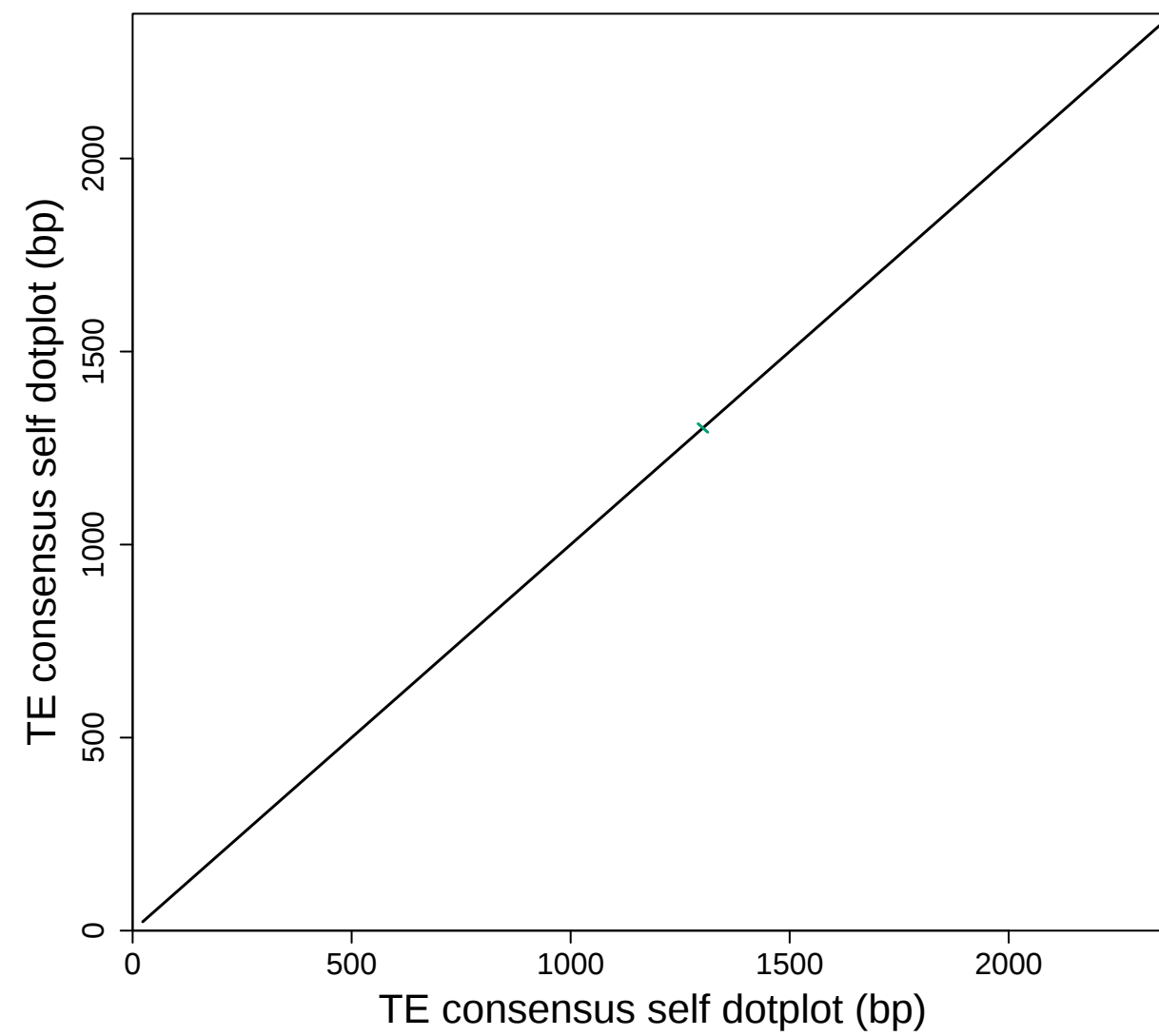
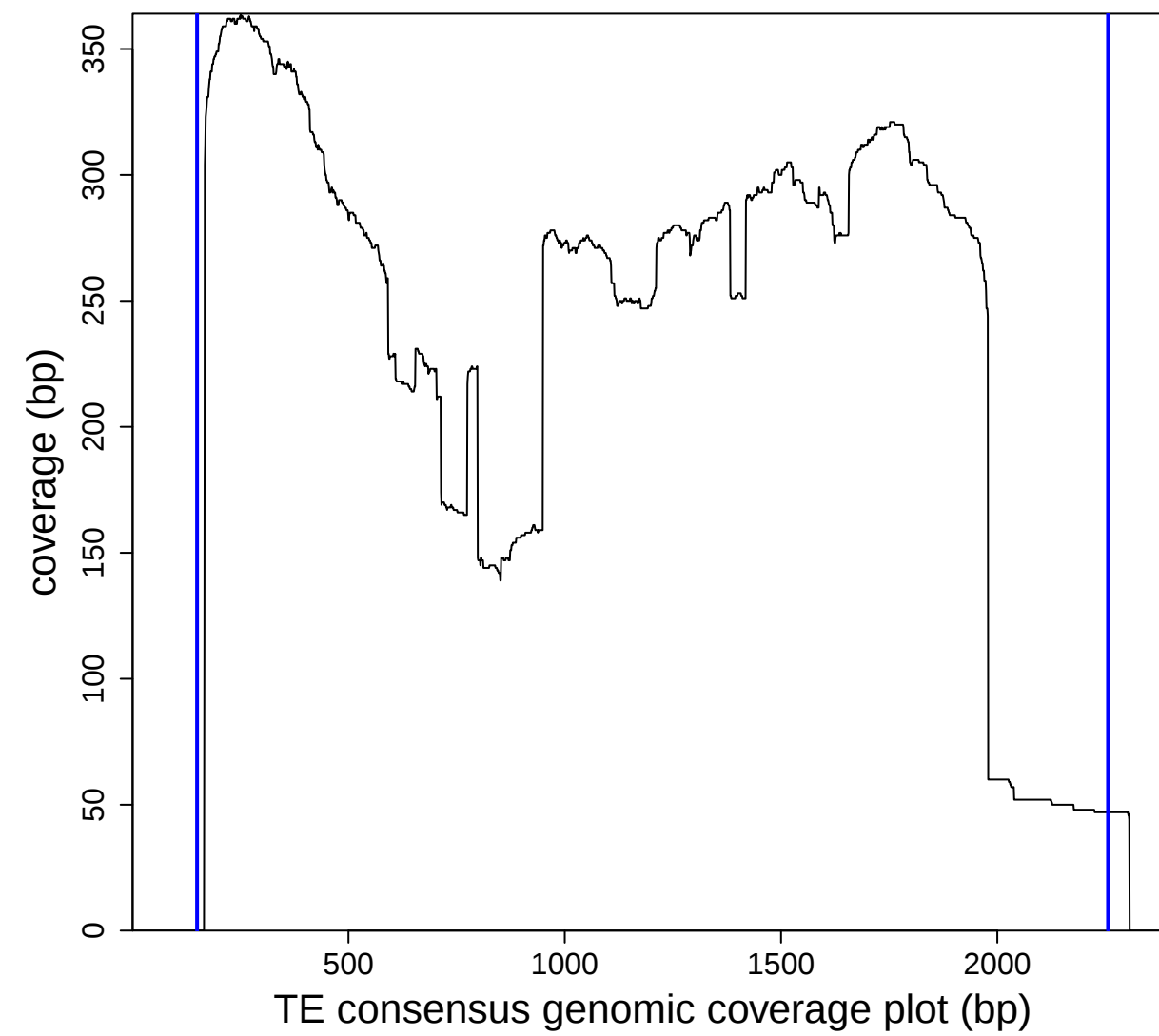
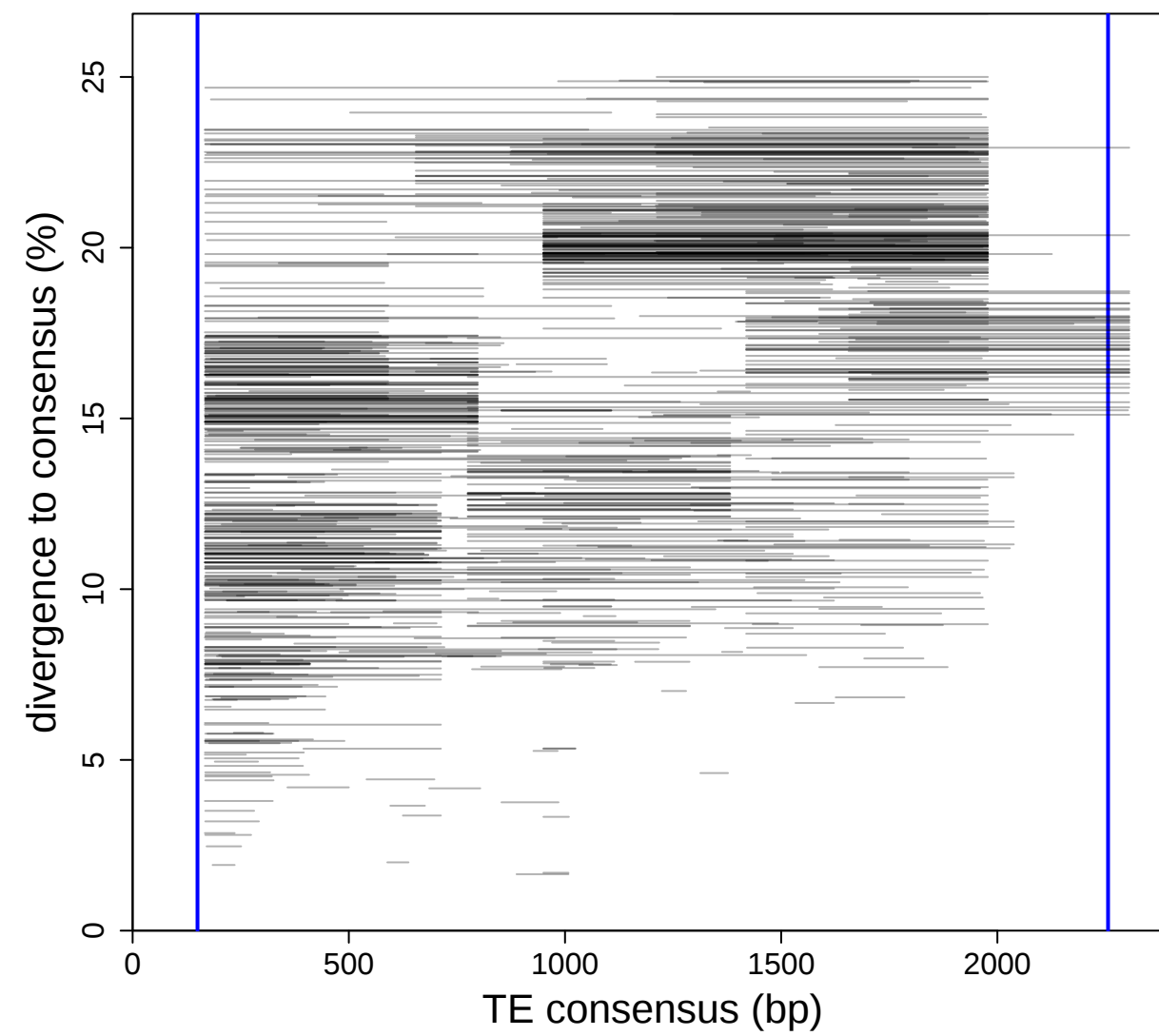
a.b.bed_uf.bed_g_4.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.f
size: 2060bp; fragments: 1138; full length: 19 (≥ 1854 bp)

After TEtrimmer 2060 bp

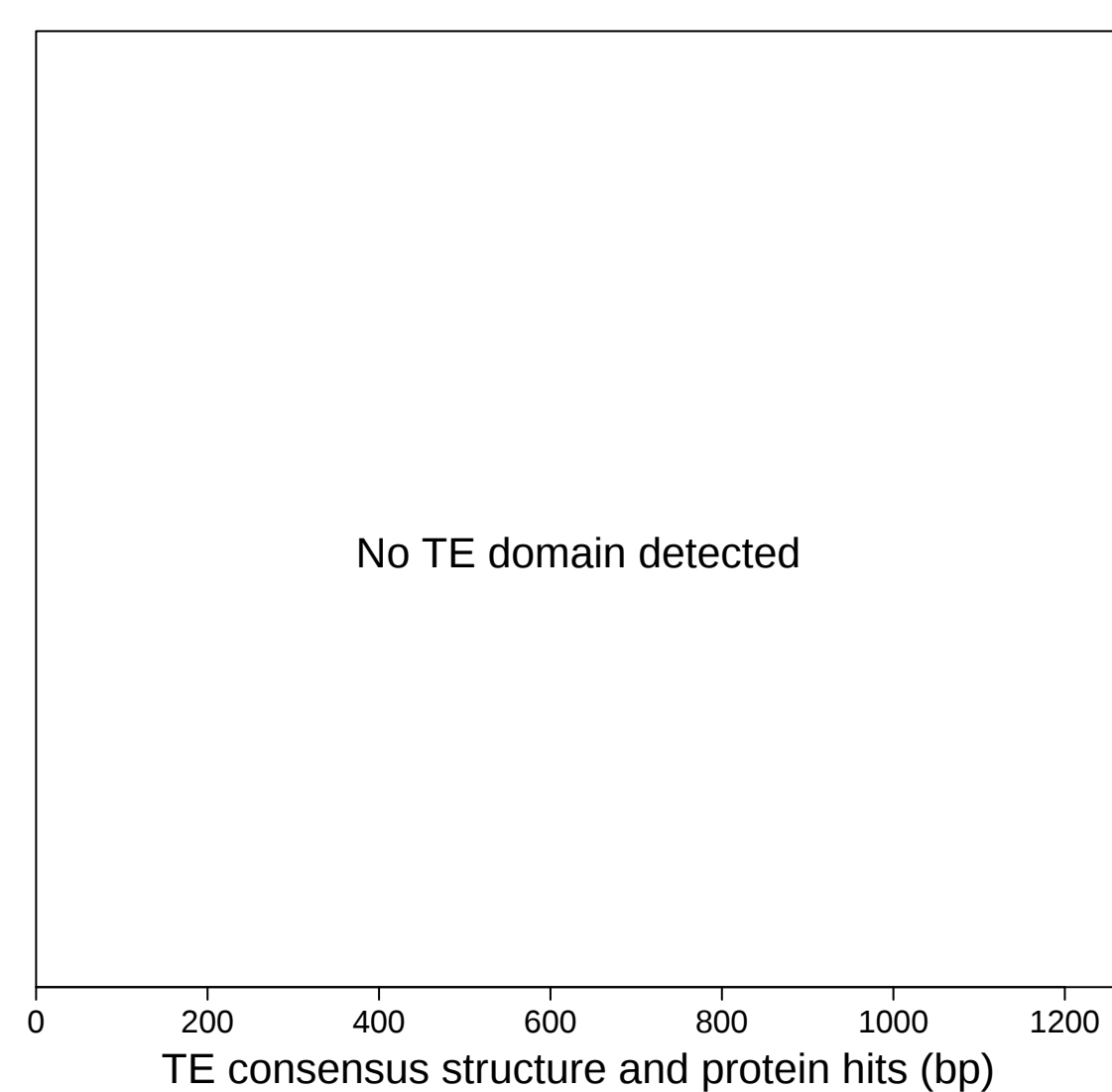
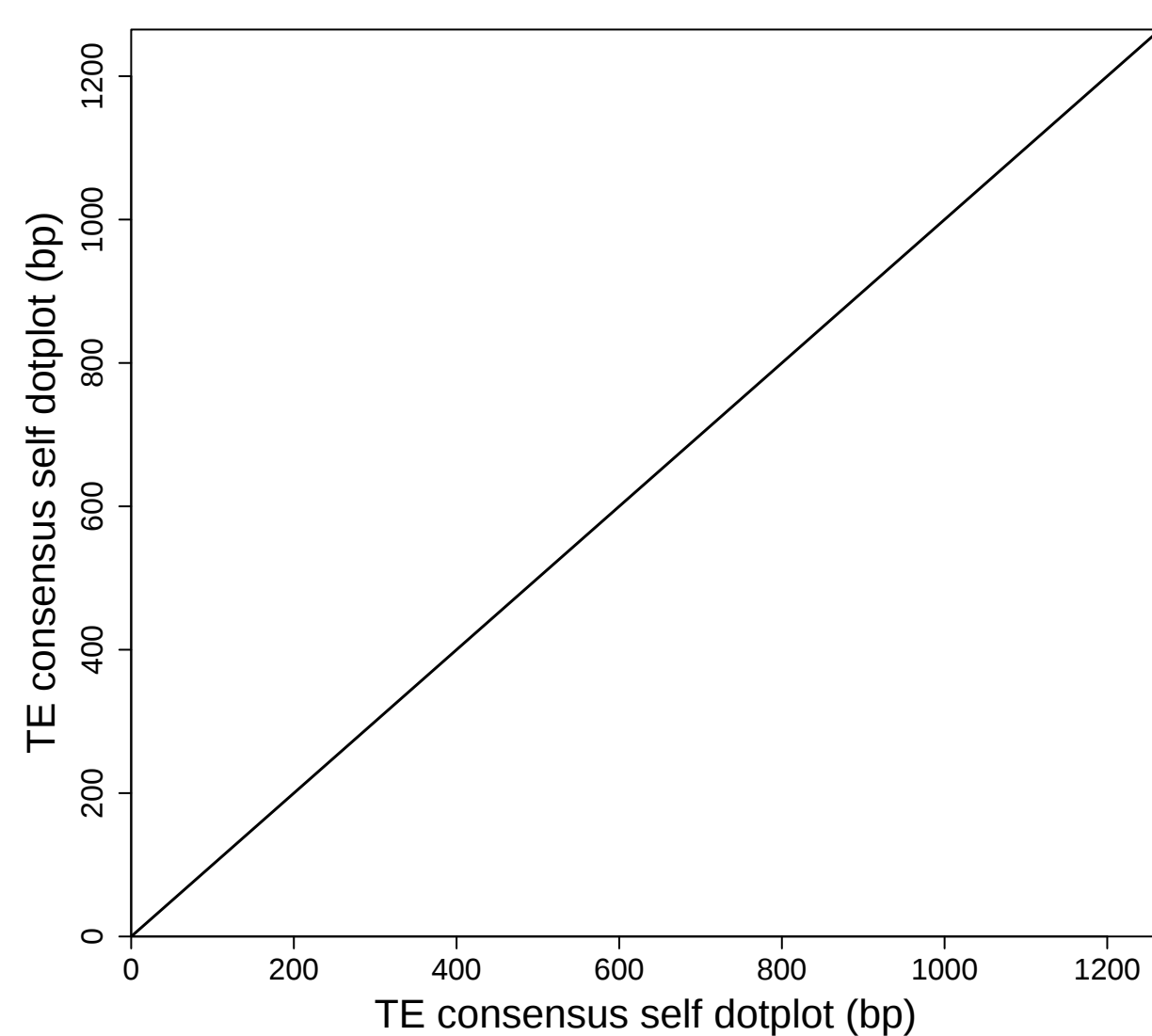
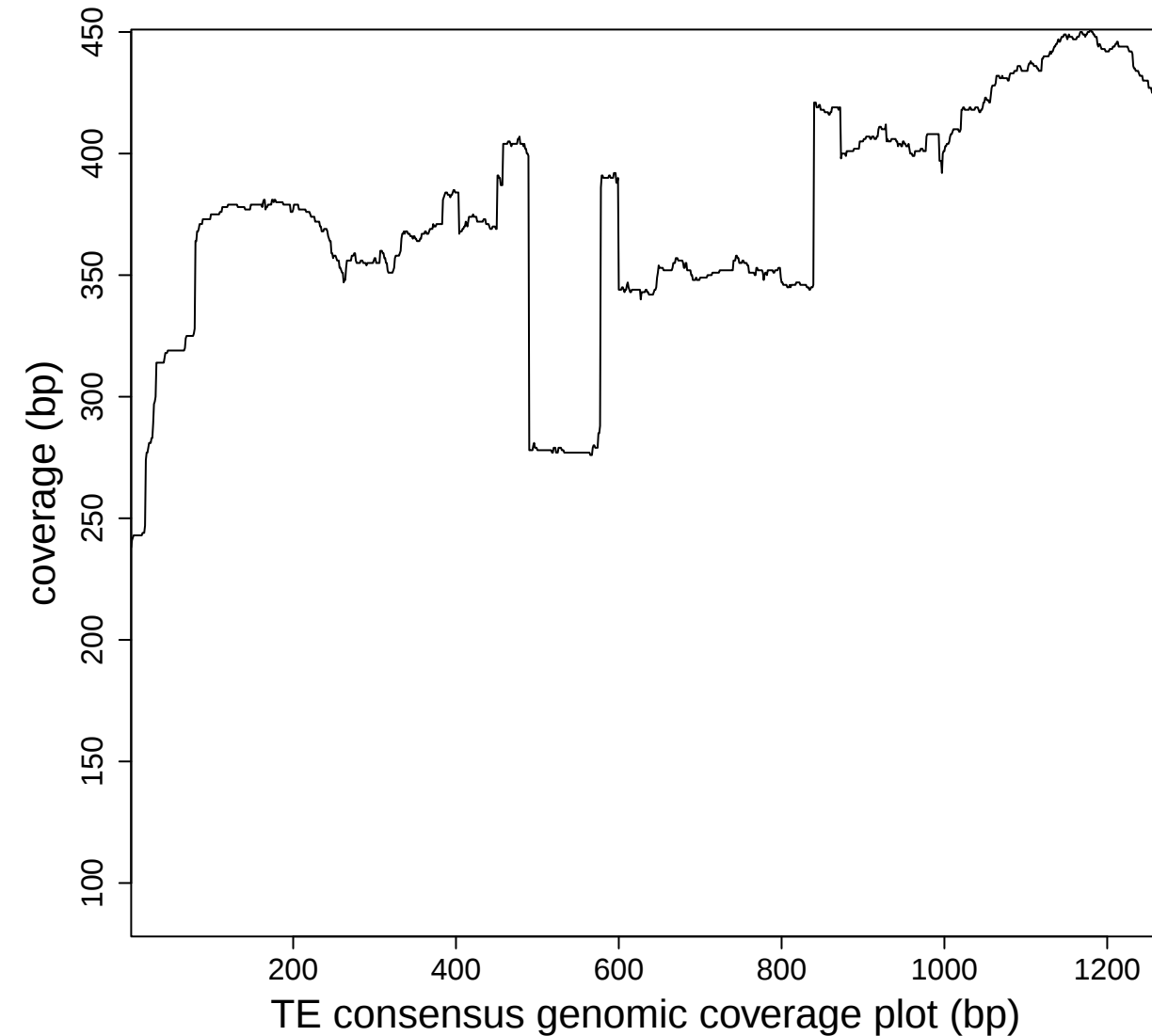
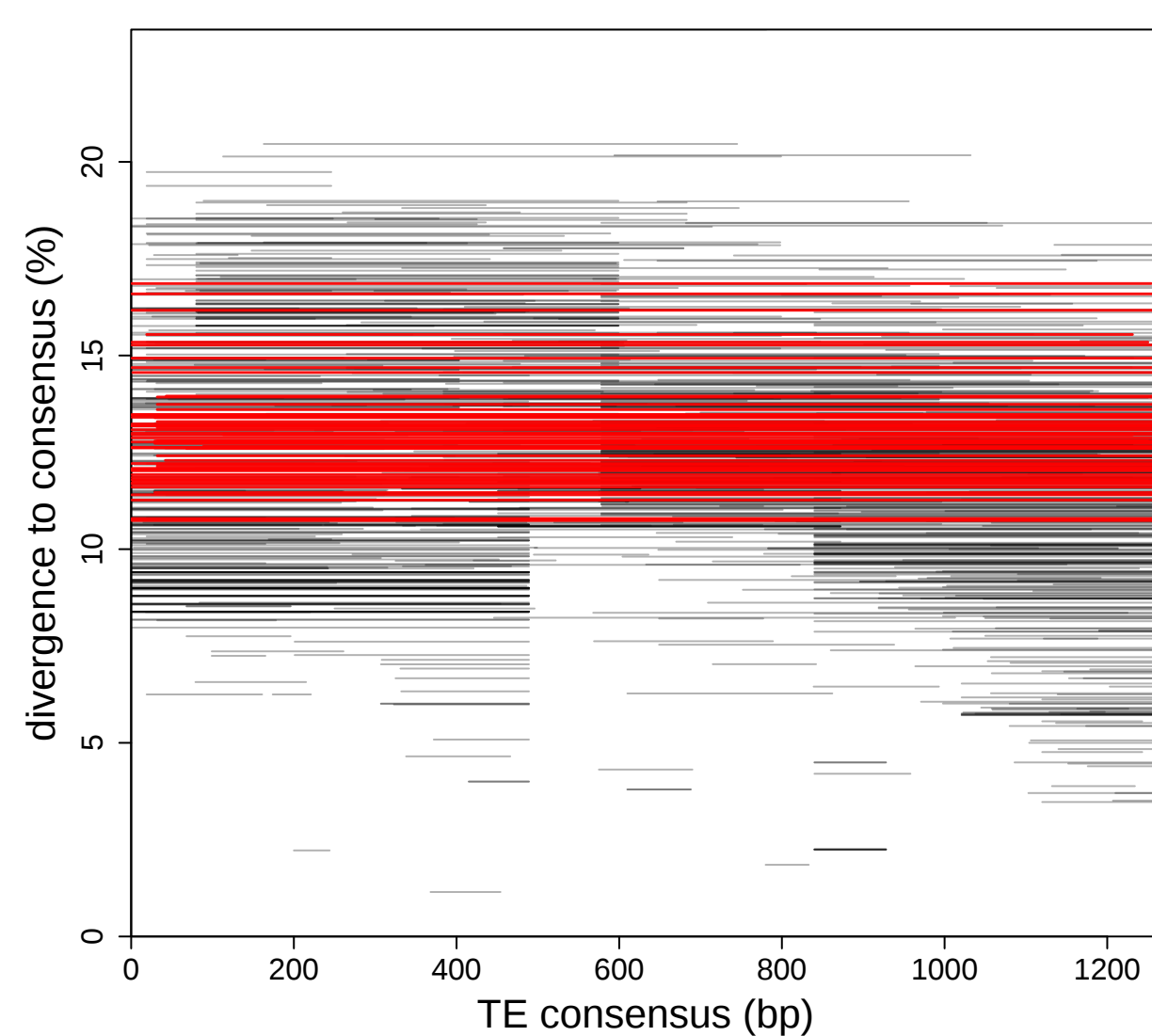


asta.b.bed_uf.bed_g_4.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_c
size: 2406bp; fragments: 879; full length: 0 (>=2165.4bp)

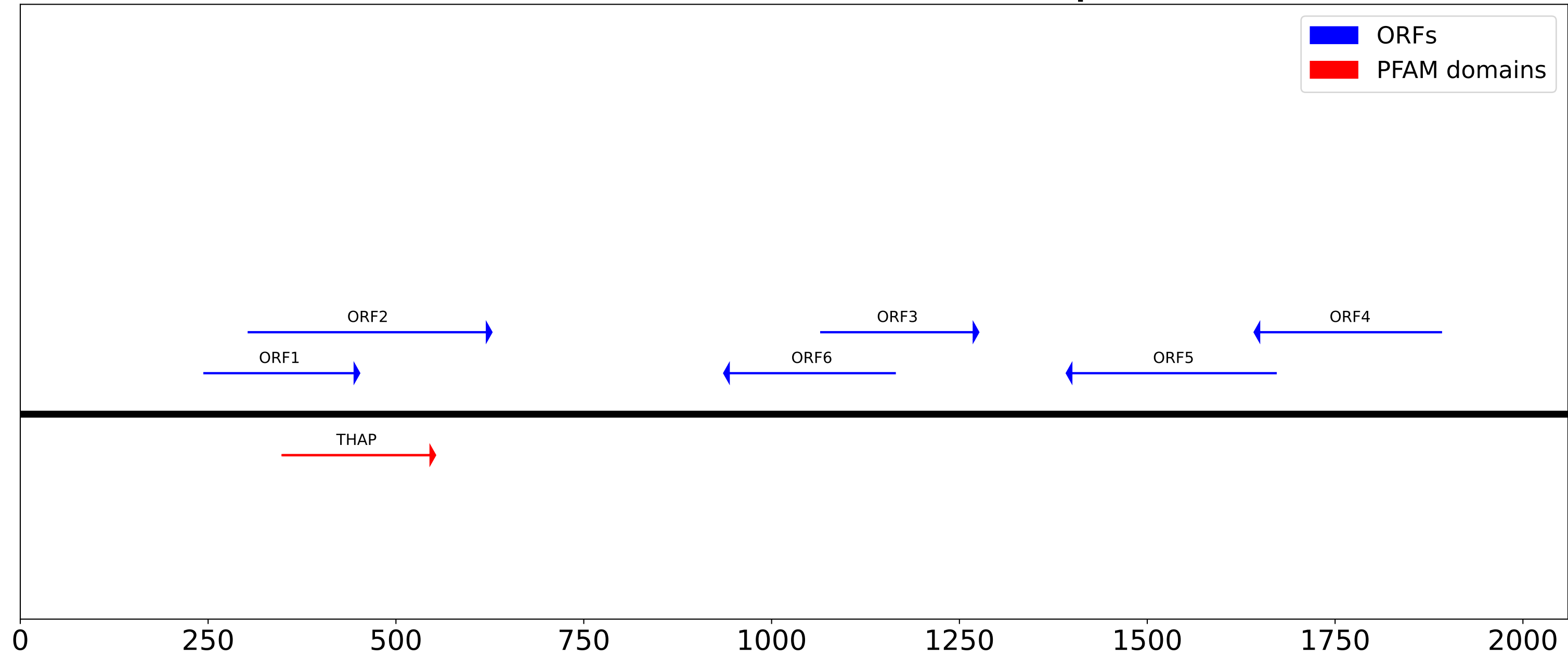
After TEtrimmer Extended plot Blue lines are boundaries



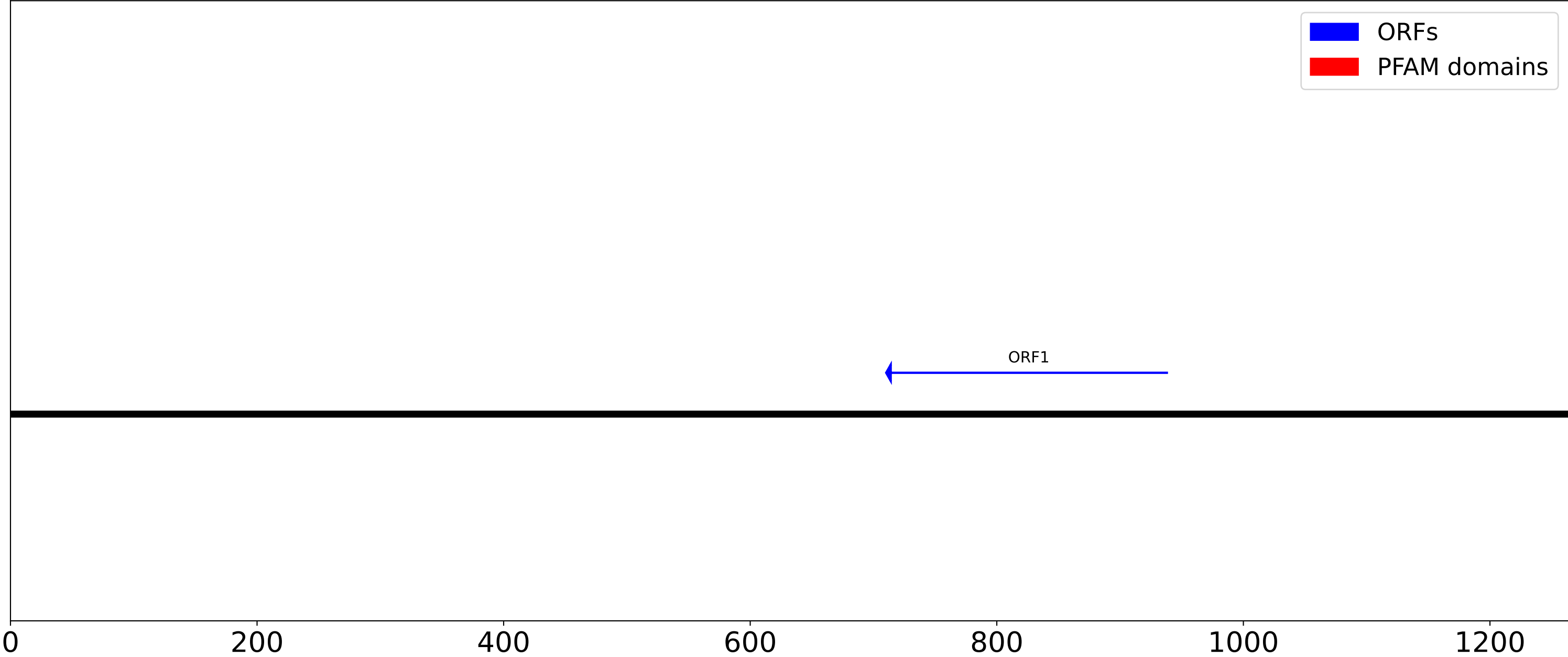
TE: rnd_4_family_1549
size: 1265bp; fragments: 950; full length: 59 (≥ 1138.5 bp)



After TEtrimmer ORF and PFAM domain plot



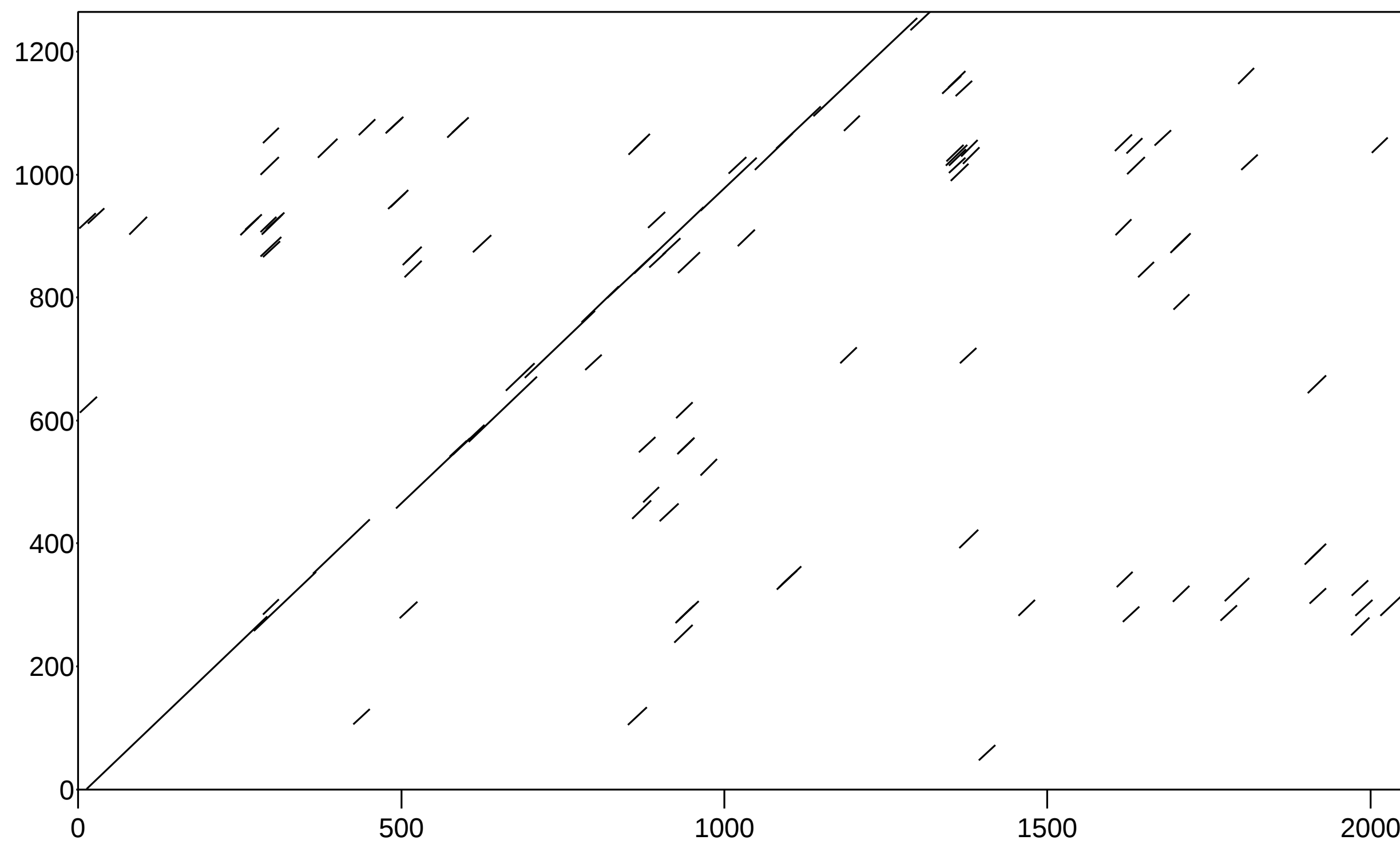
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)