

Start crop Point

End crop Point

0

10

0

200

400

600

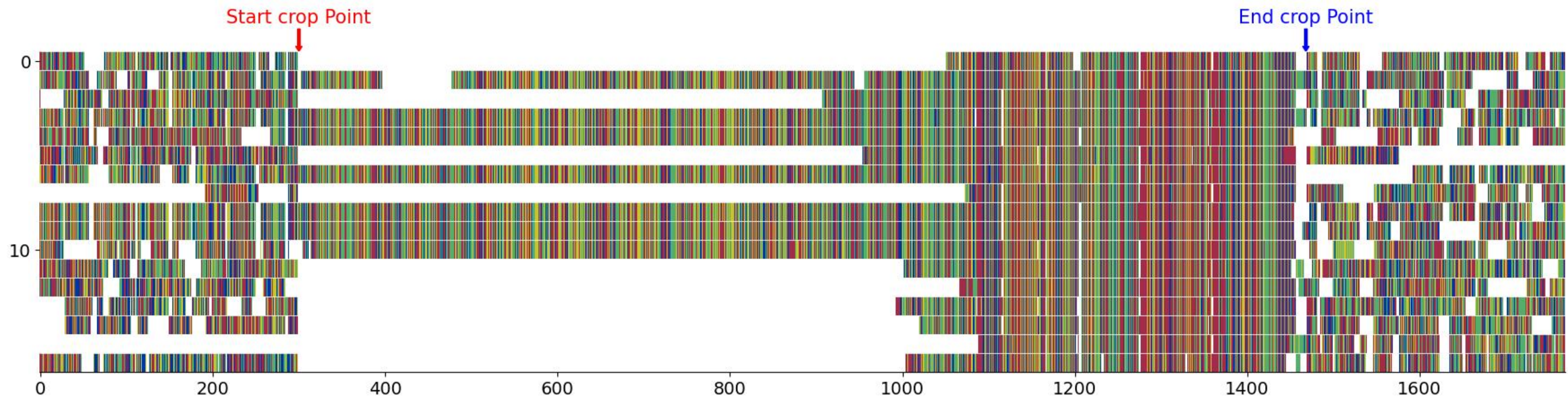
800

1000

1200

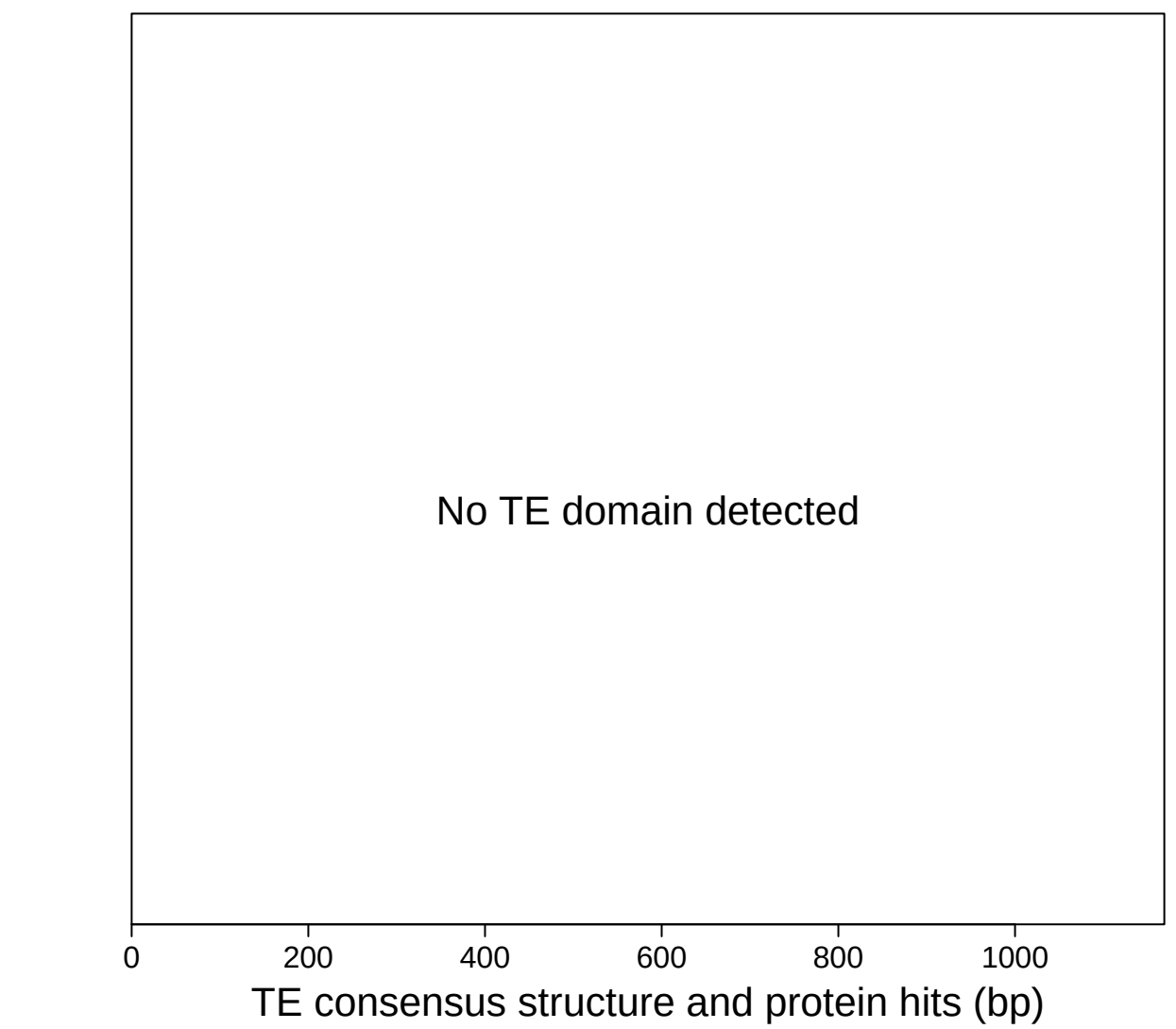
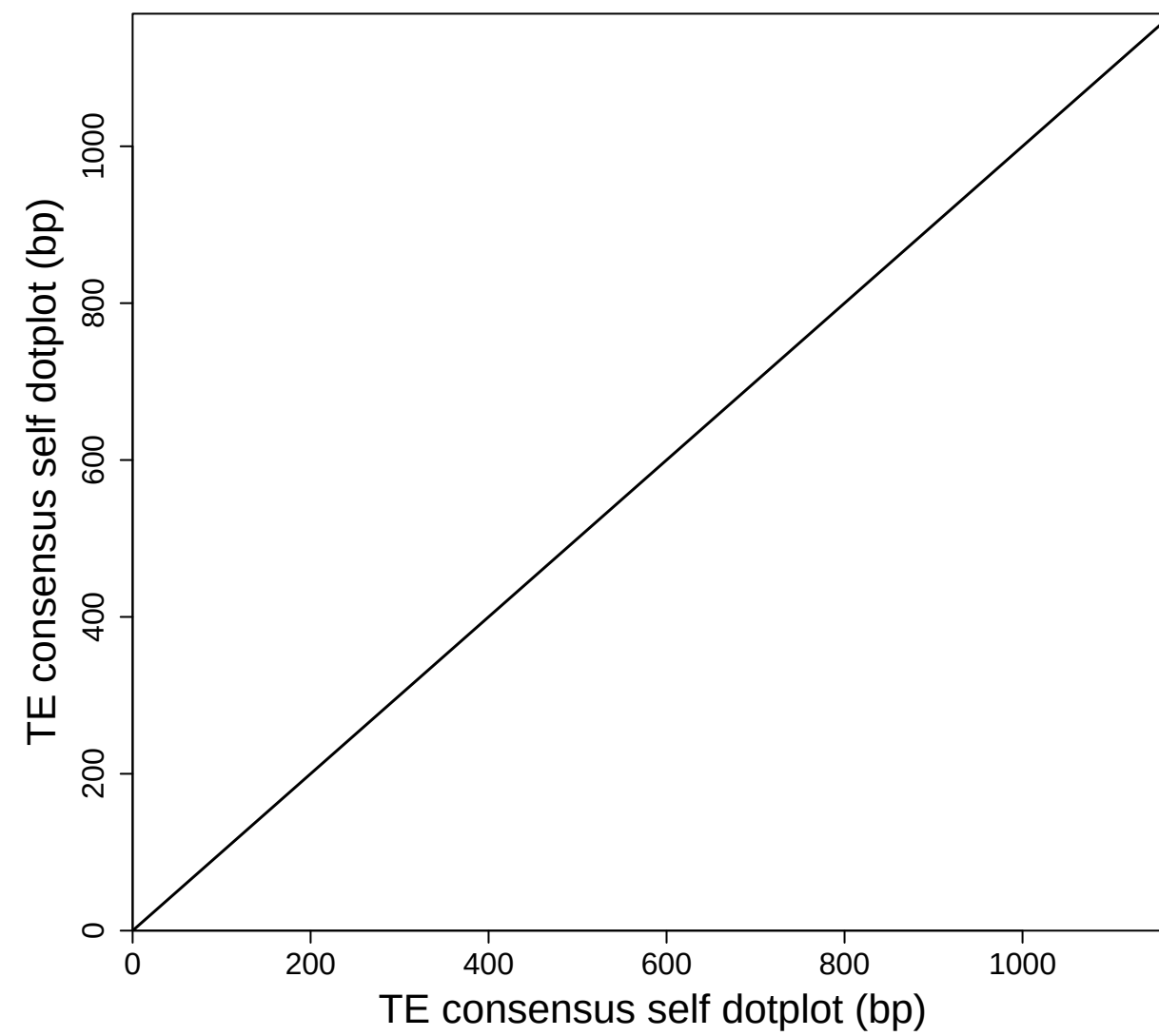
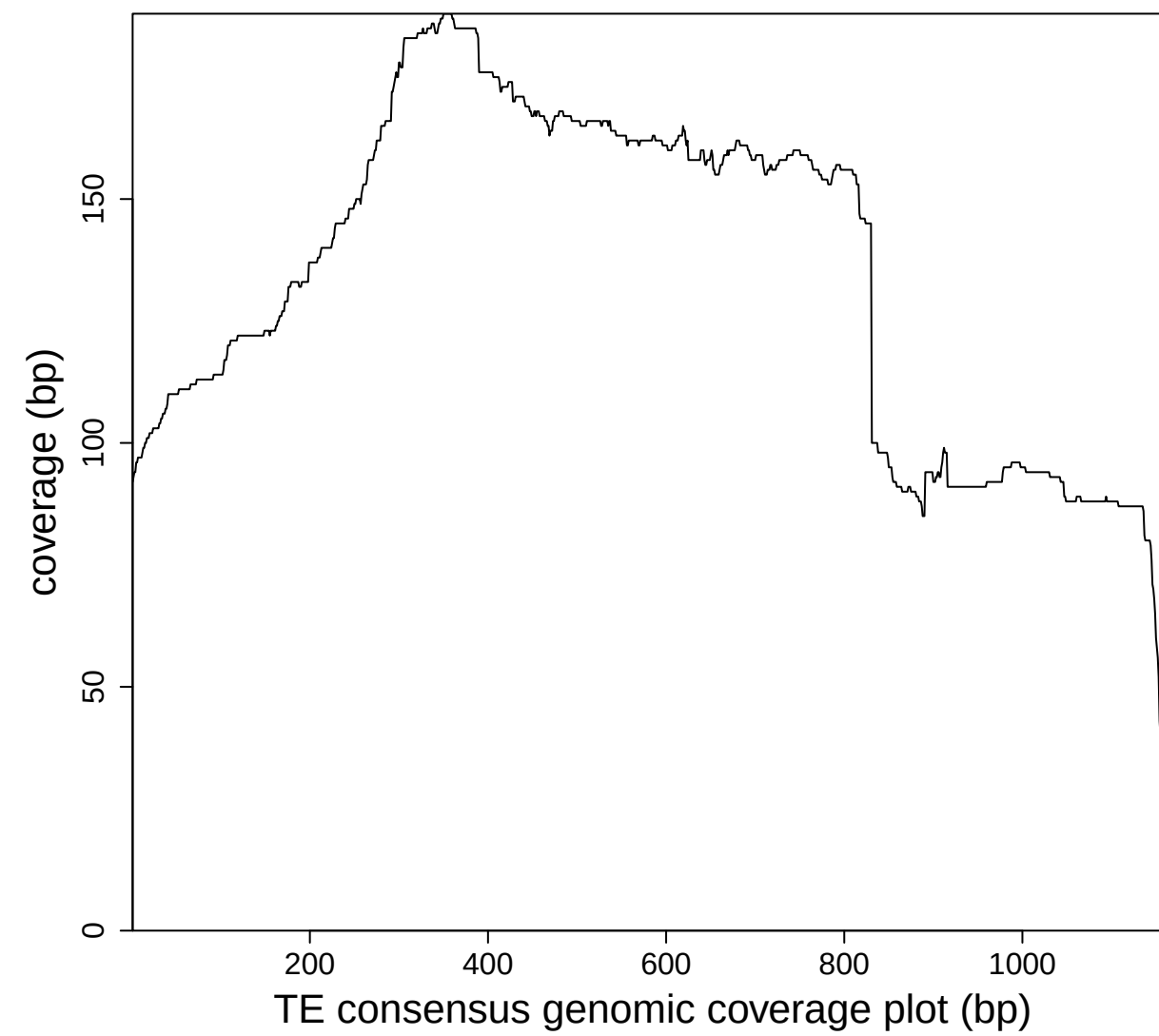
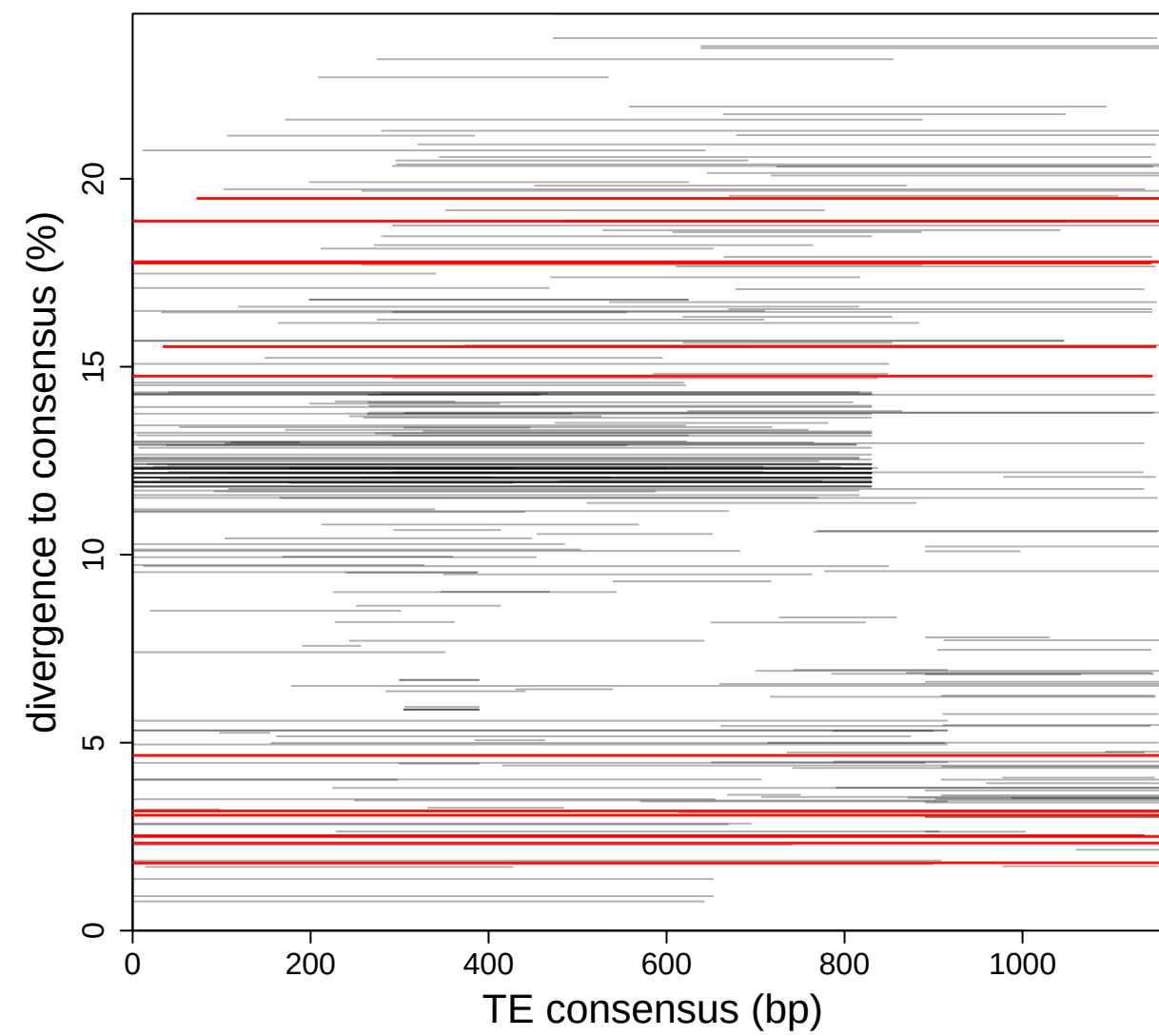
1400

1600



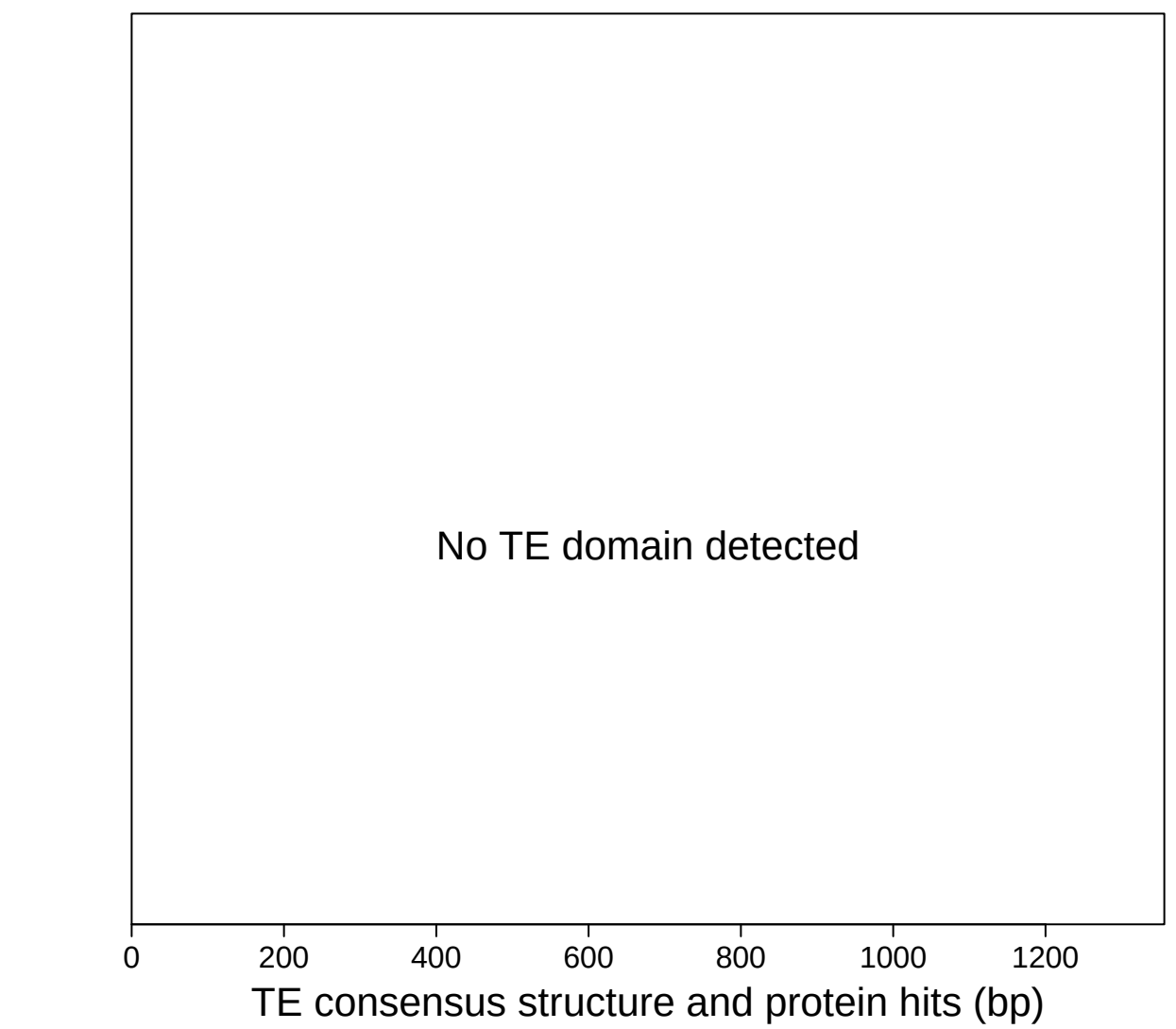
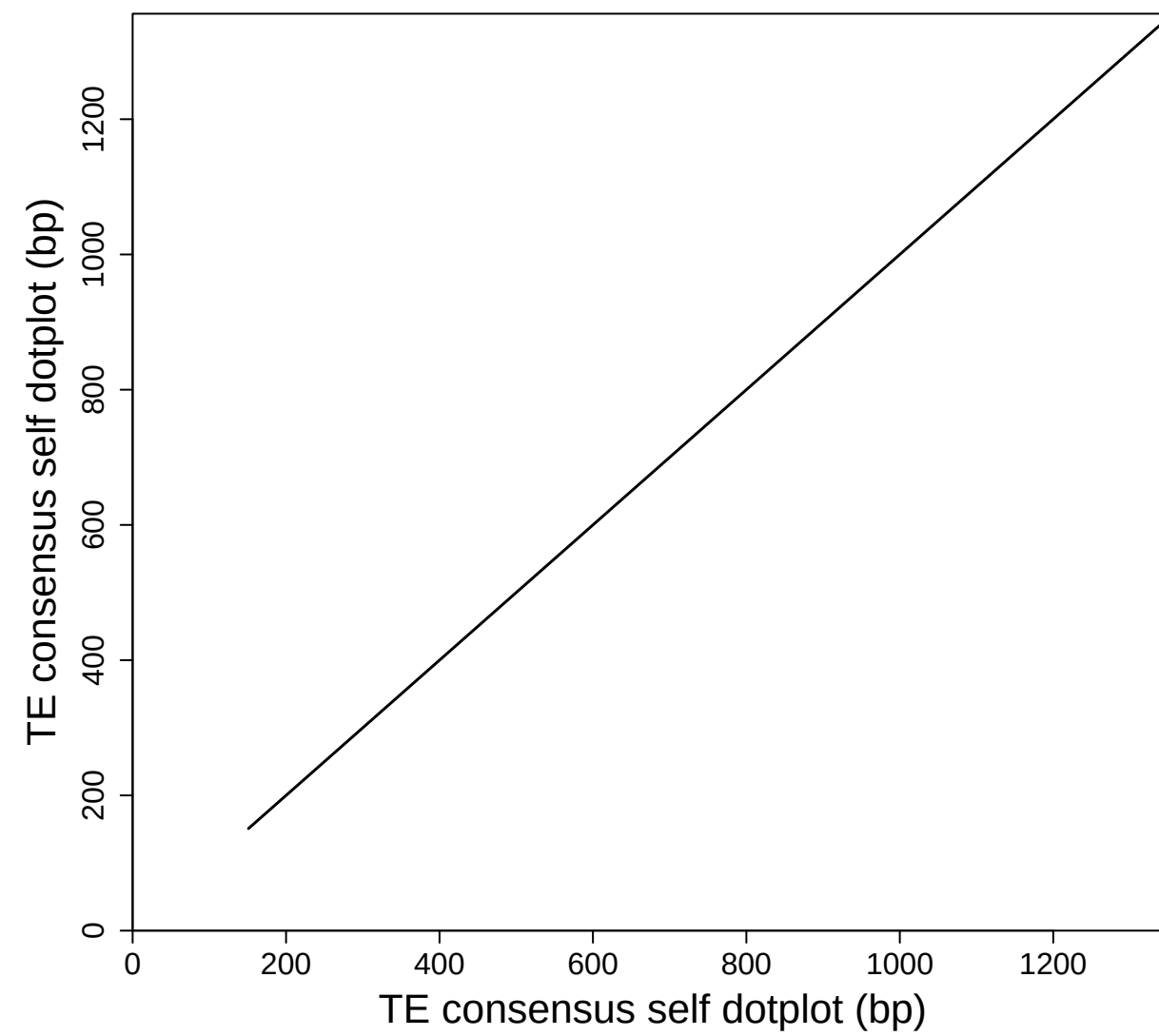
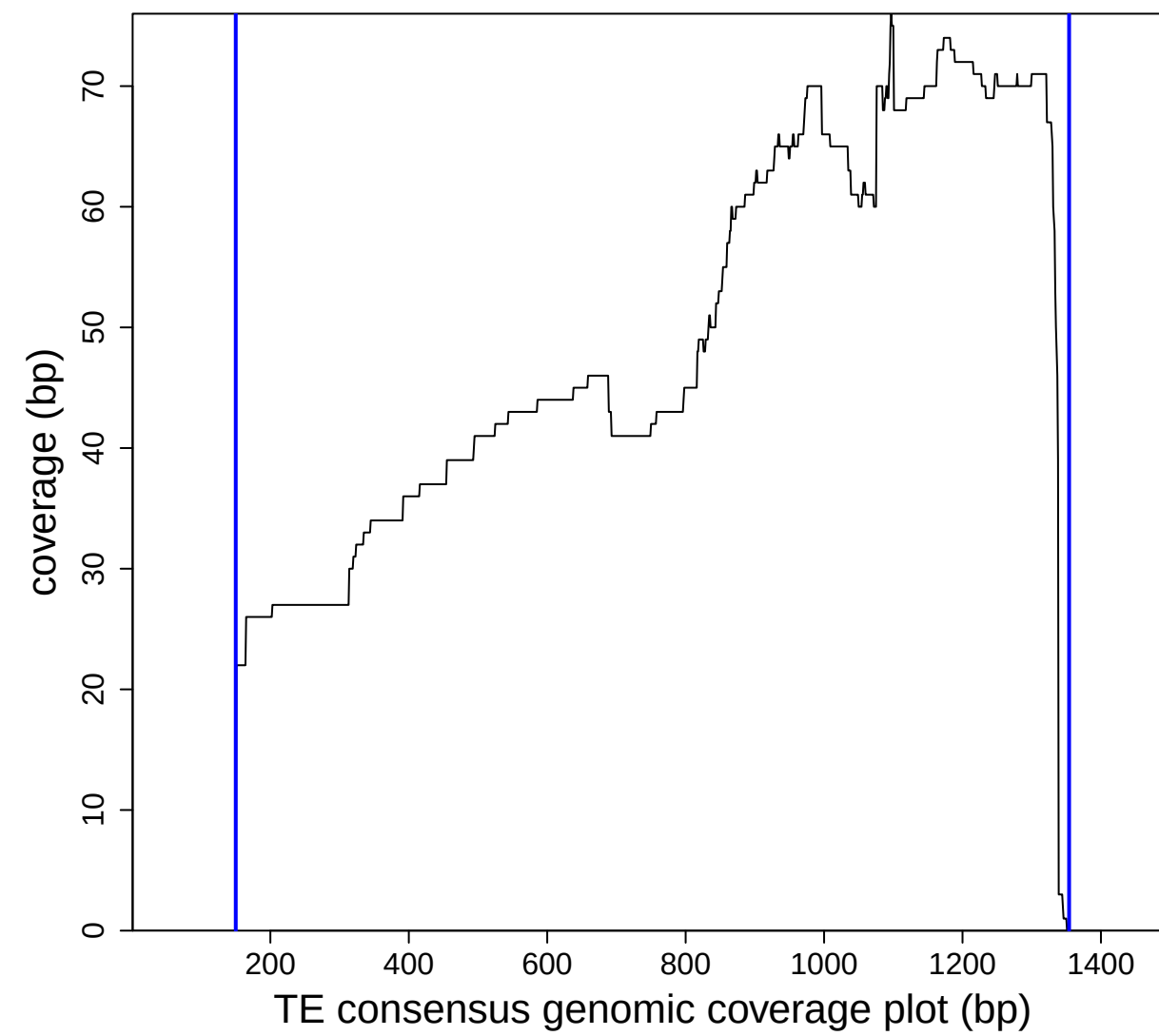
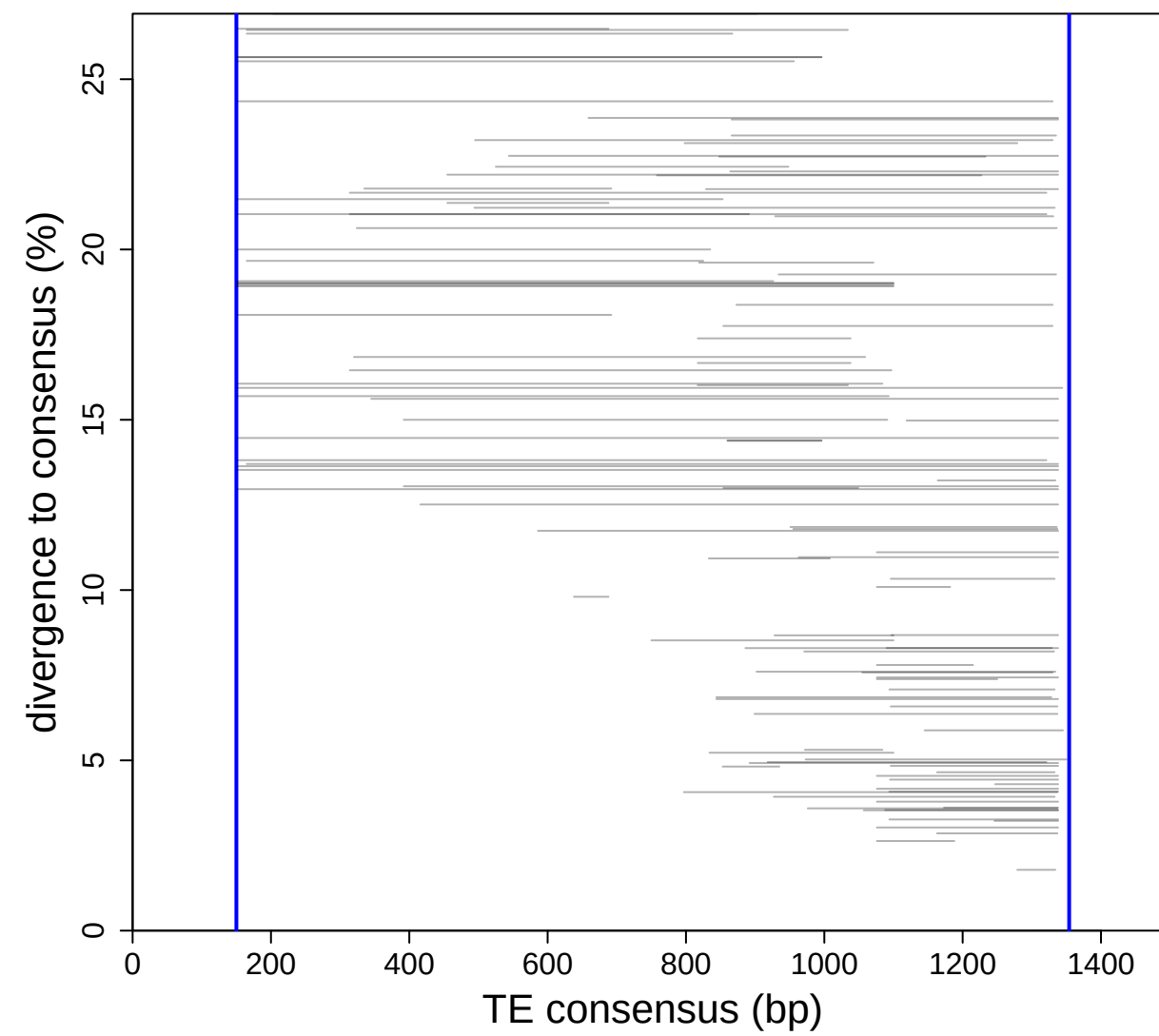
bed_uf.bed g_2.bed fm_1.bed 0_0_bcln.fa aln.fa cl.fa gs.fa c
size: 1169bp; fragments: 283; full length: 13 (>=1052.1bp)

After TEtrimmer 1169 bp

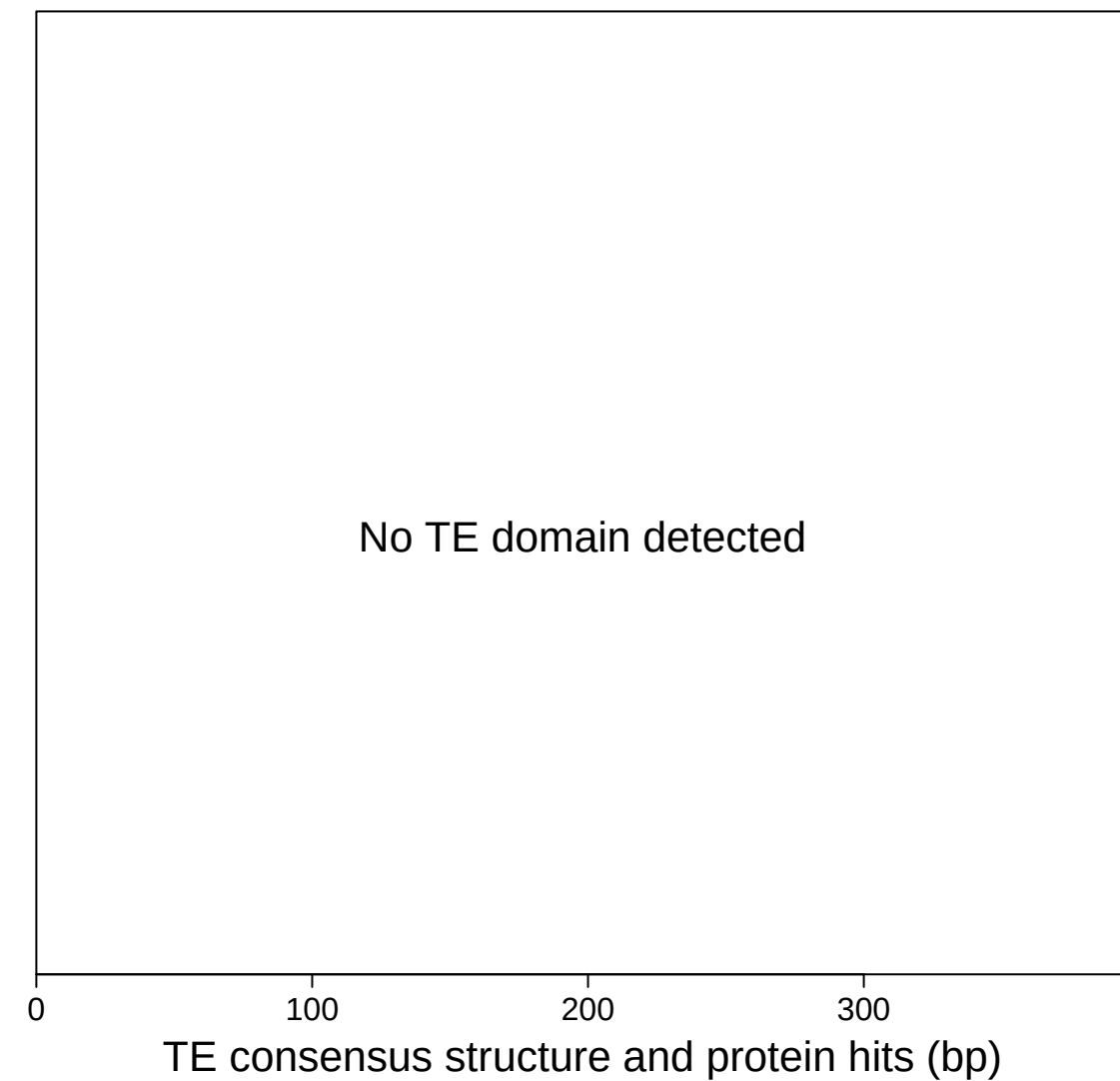
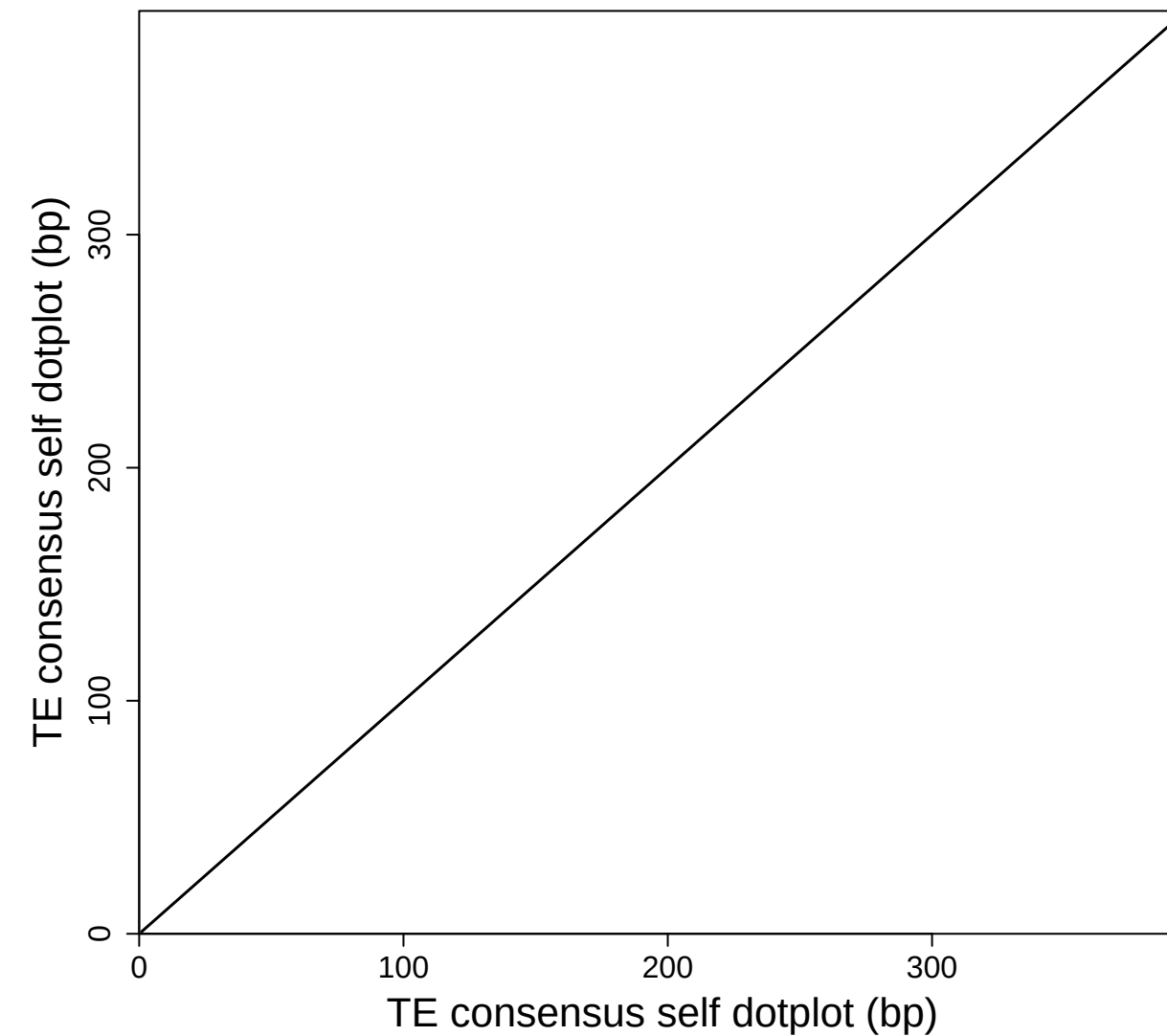
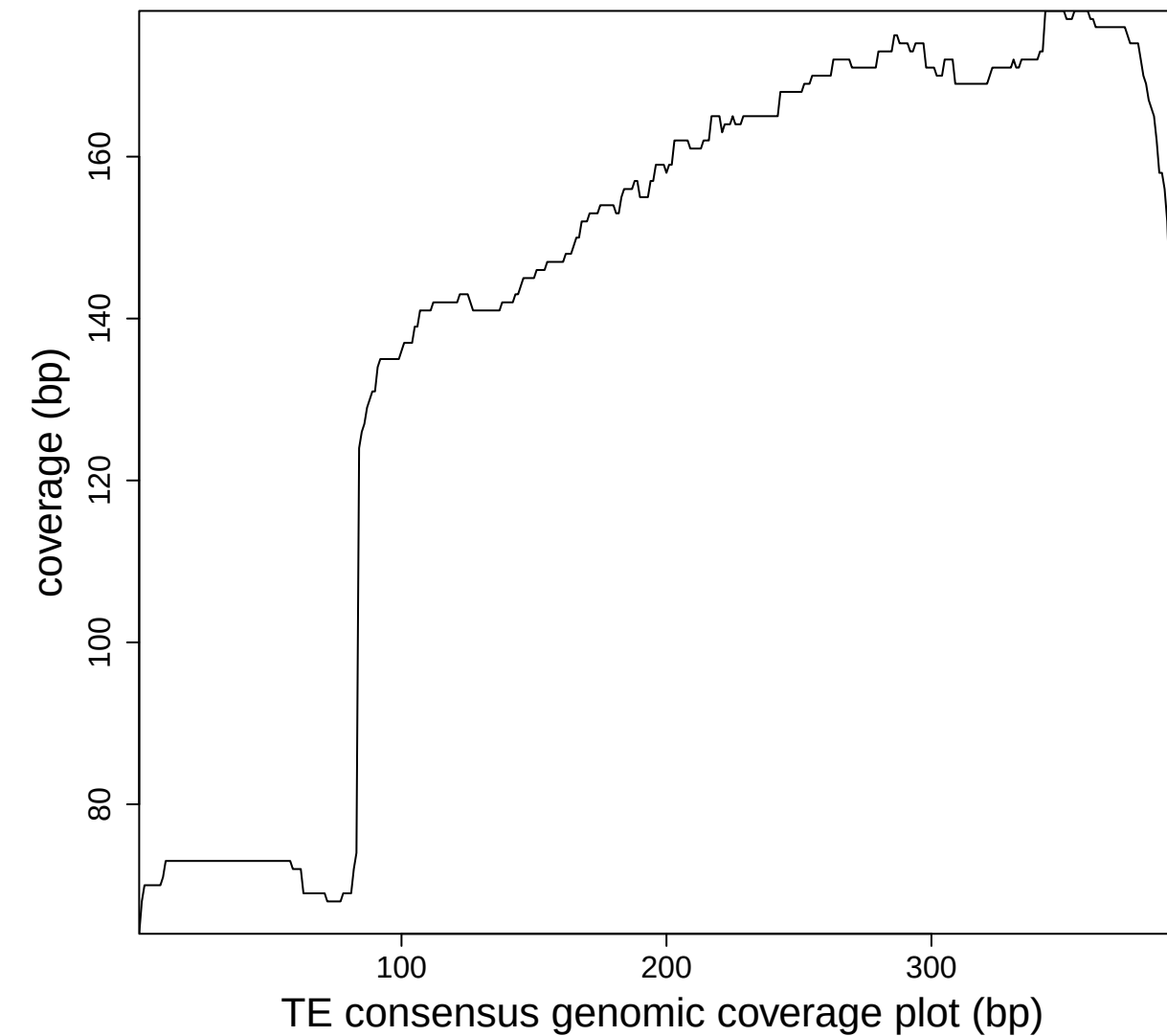
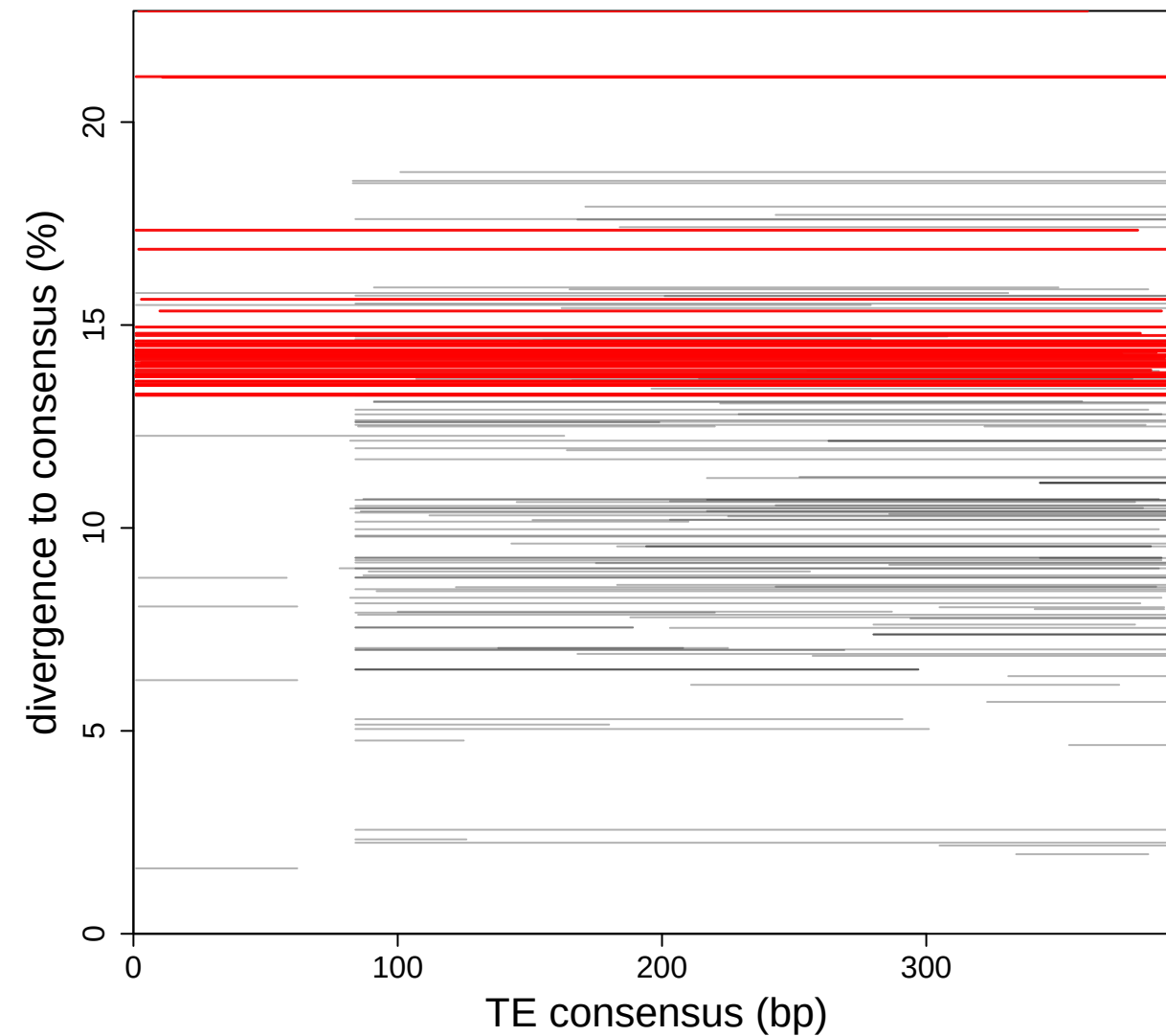


6.fasta.b.bed uf.bed g_2.bed fm_1.bed 0_0_bcln.fa aln.fa cl.f
size: 1504bp; fragments: 117; full length: 0 (>=1353.6bp)

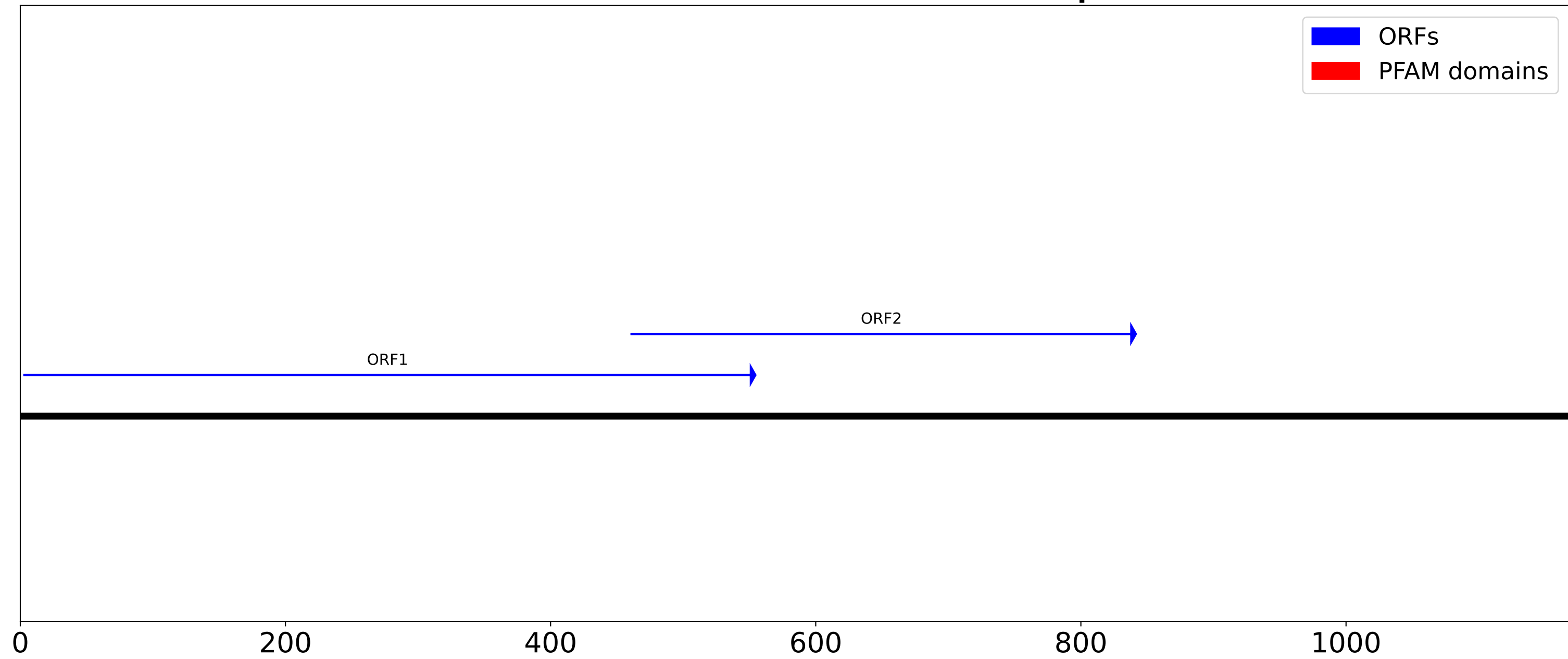
After TEtrimmer Extended plot Blue lines are boundaries



TE: rnd_5_family_7356
size: 396bp; fragments: 210; full length: 65 (≥ 356.4 bp)



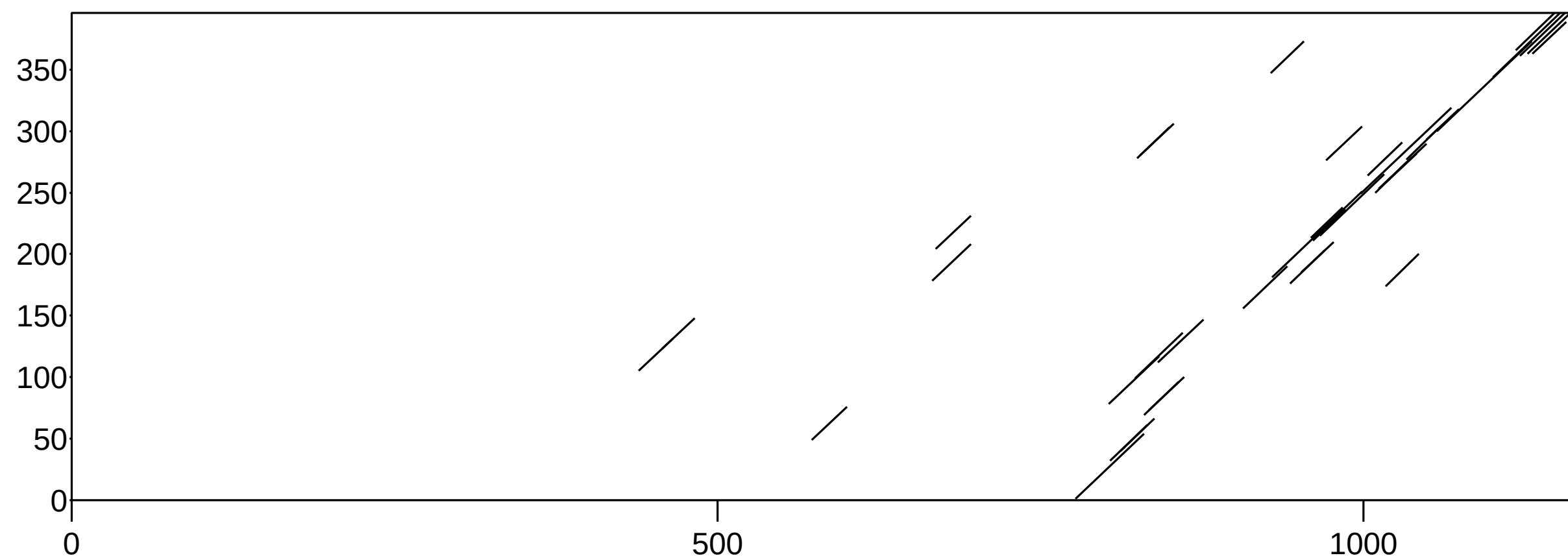
After TEtrimmer ORF and PFAM domain plot



Dotplot

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

E consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)