

Start crop Point

End crop Point

MSA length = 1043



Start crop Point

End crop Point

0 -

10 -

0

200

400

600

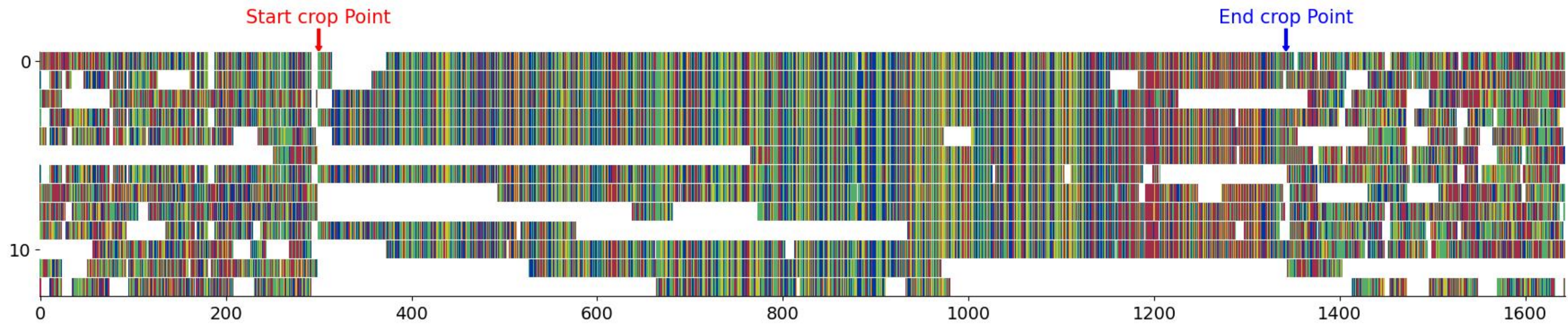
800

1000

1200

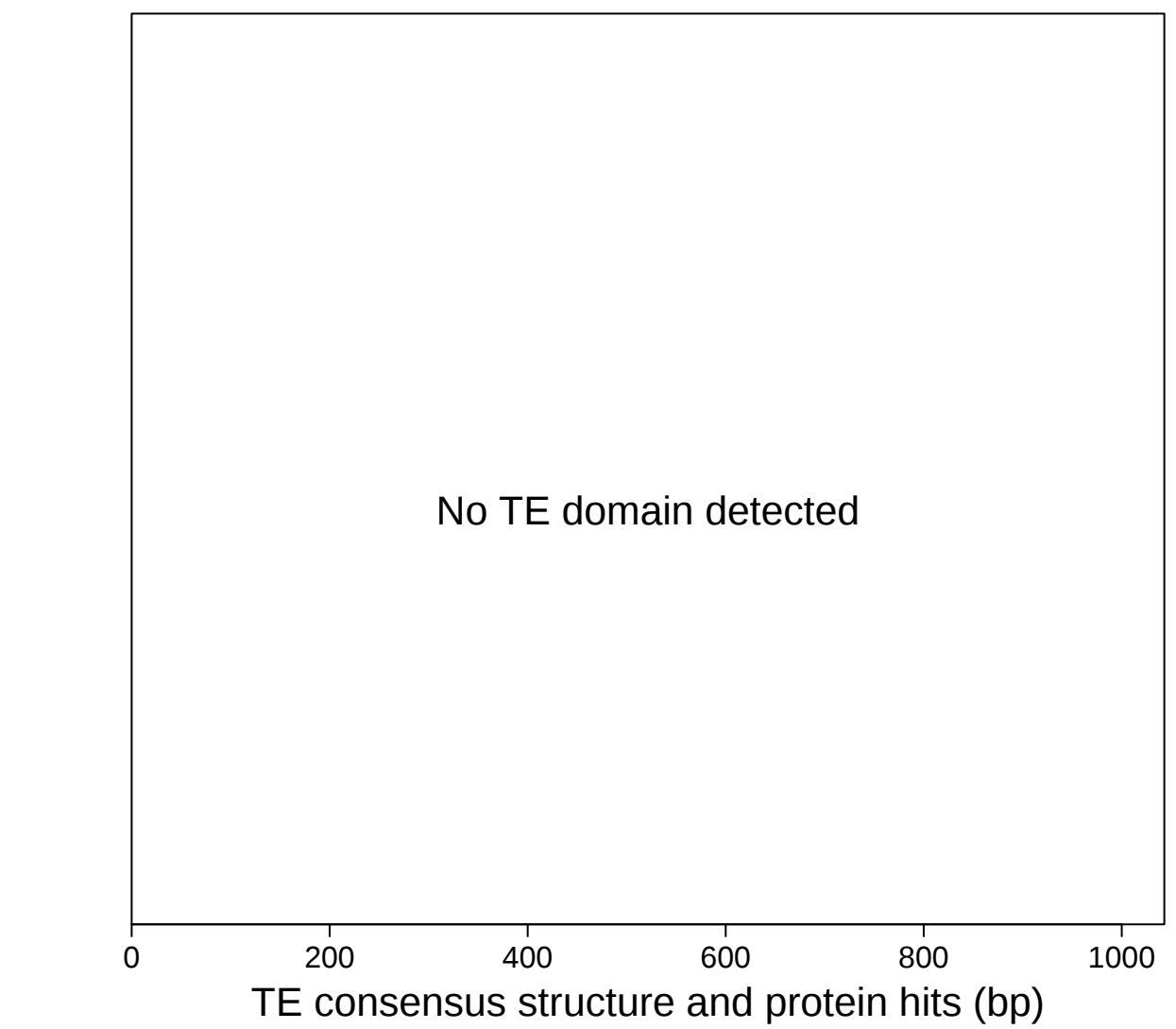
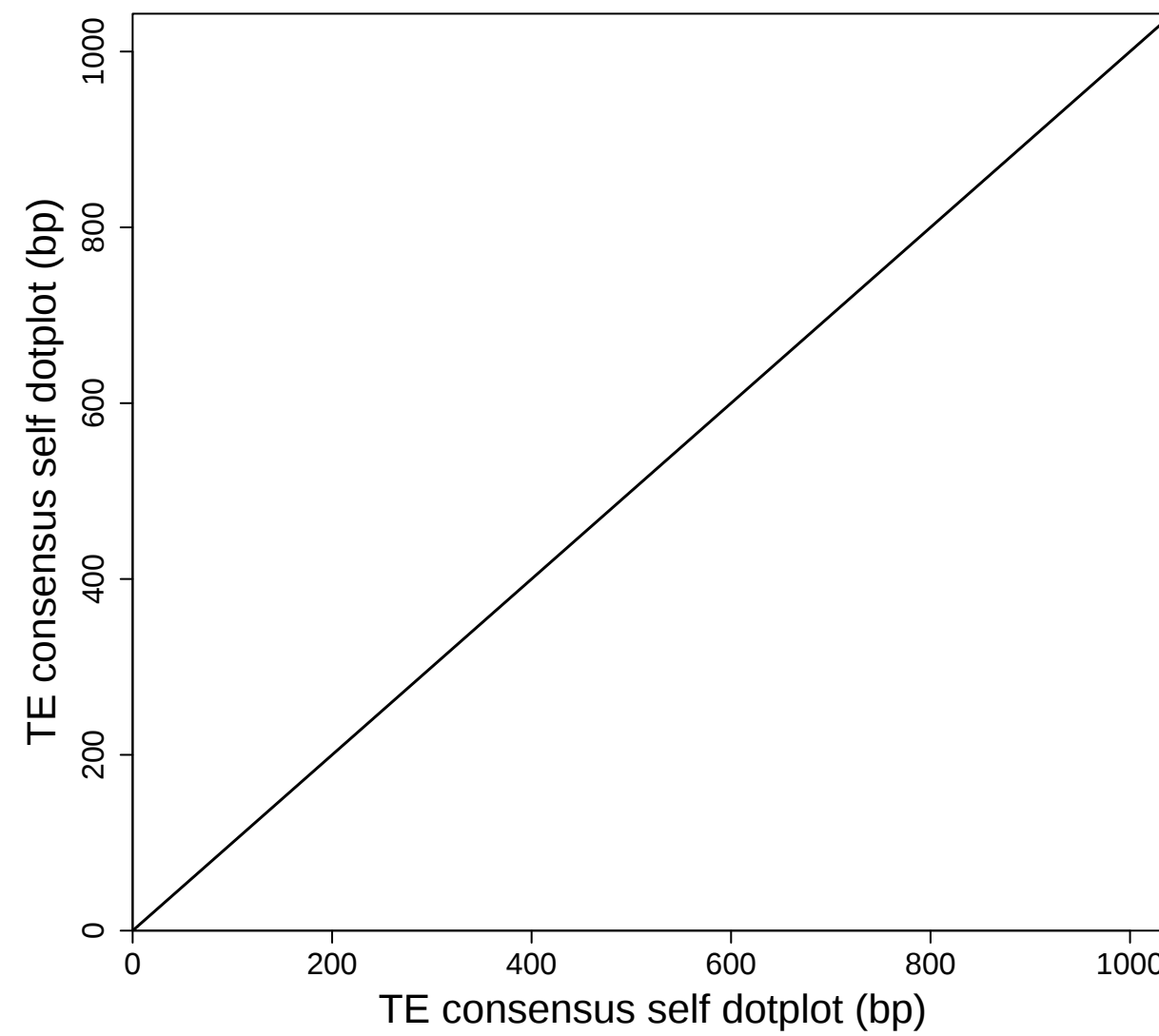
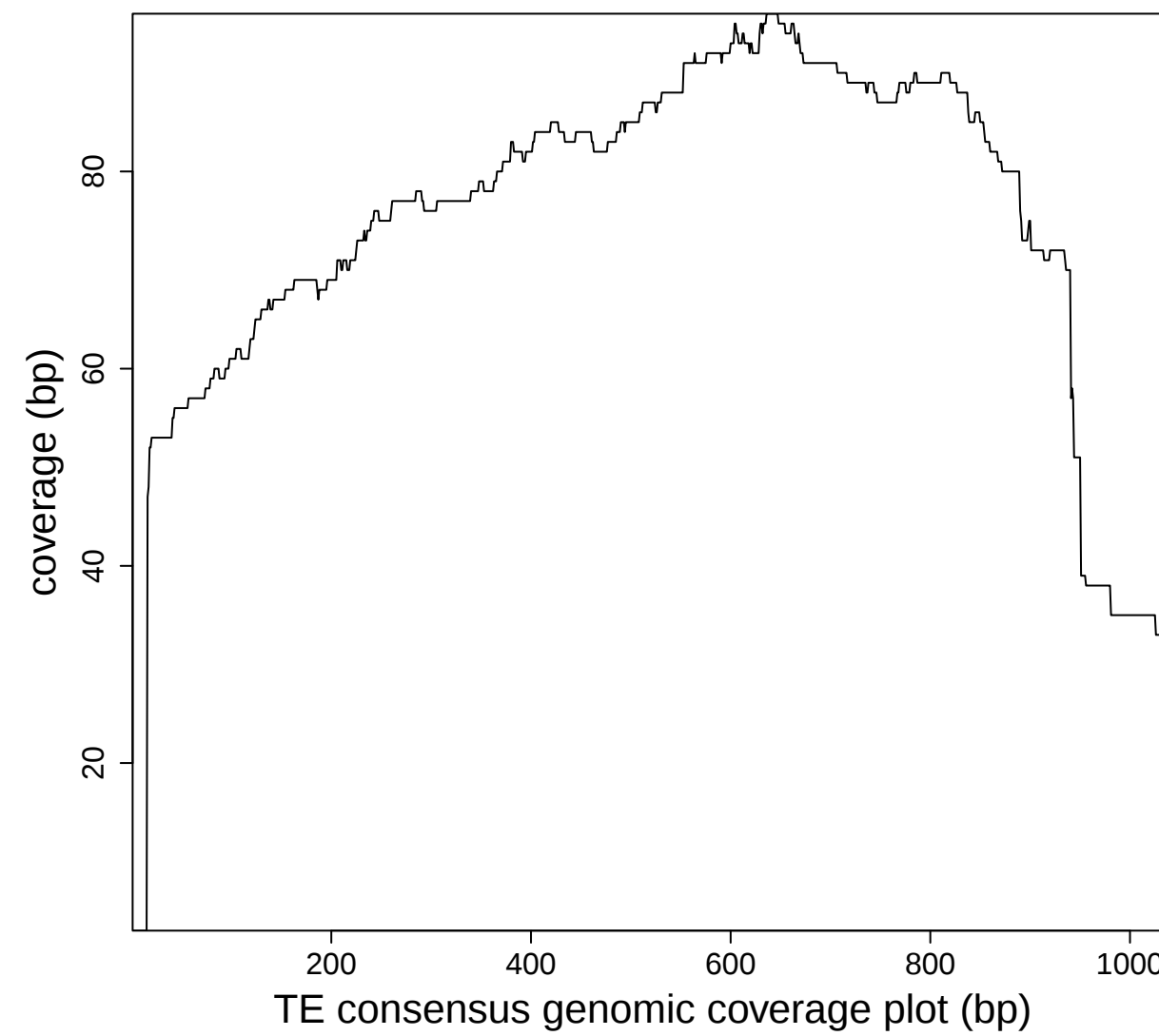
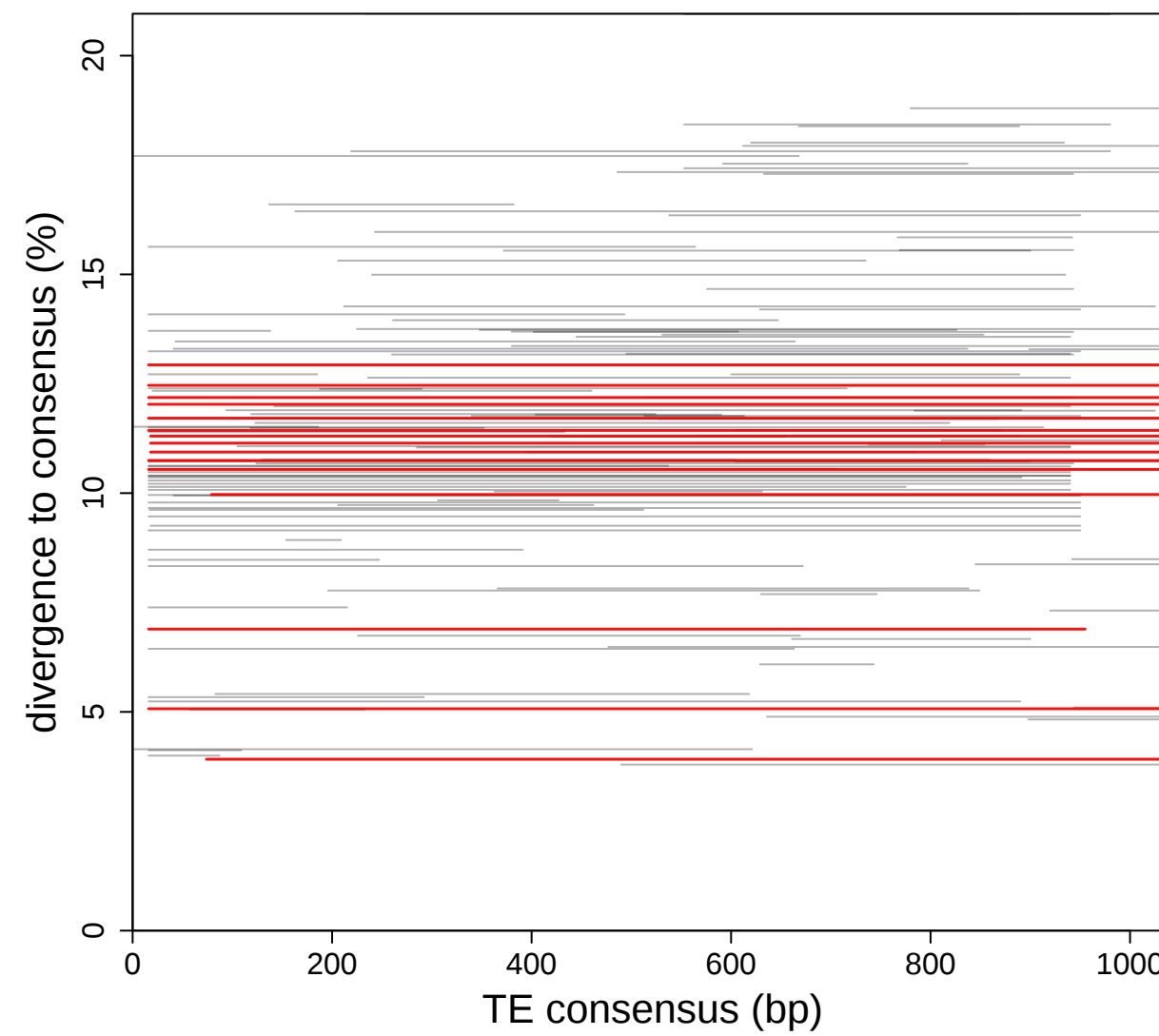
1400

1600



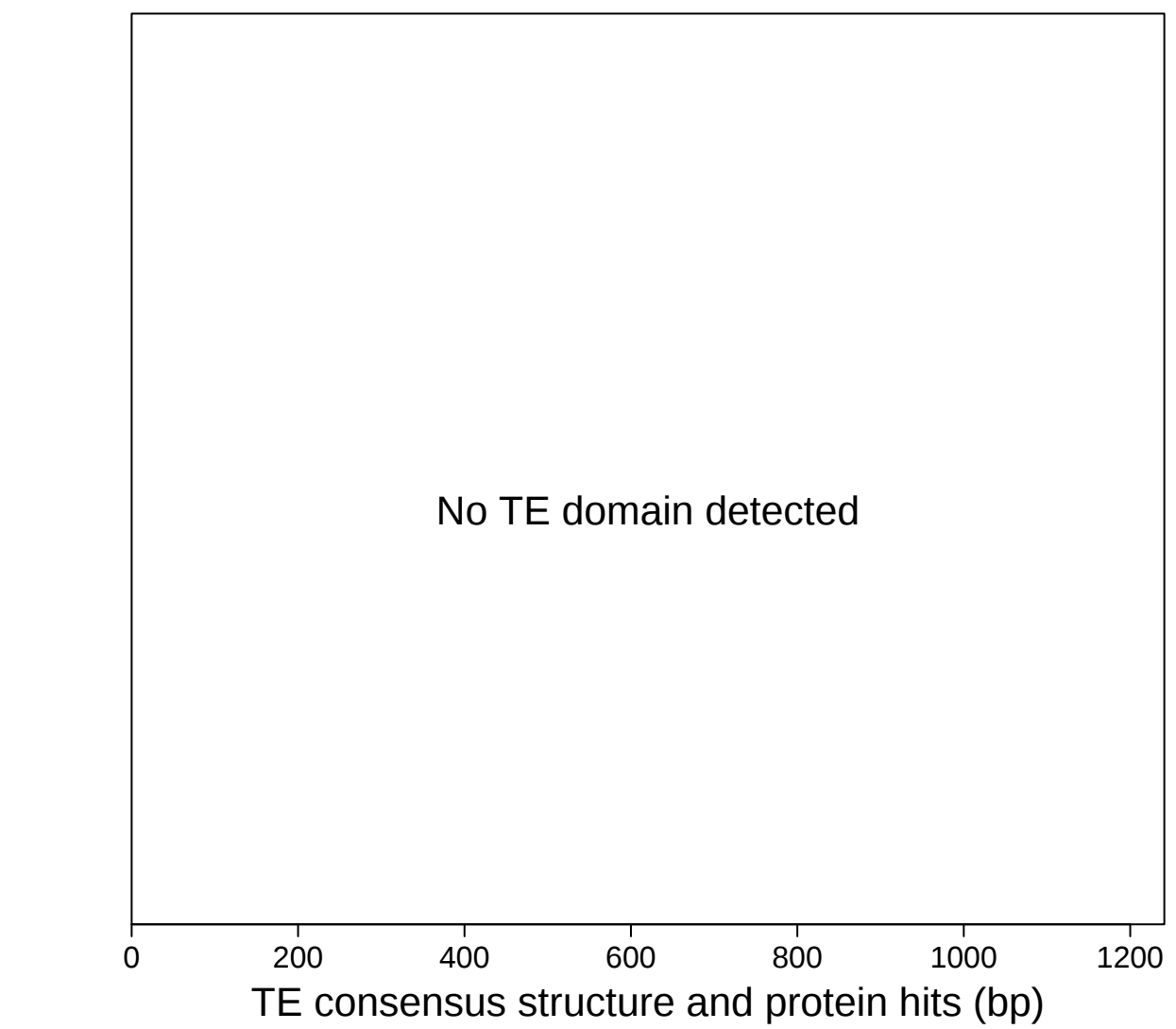
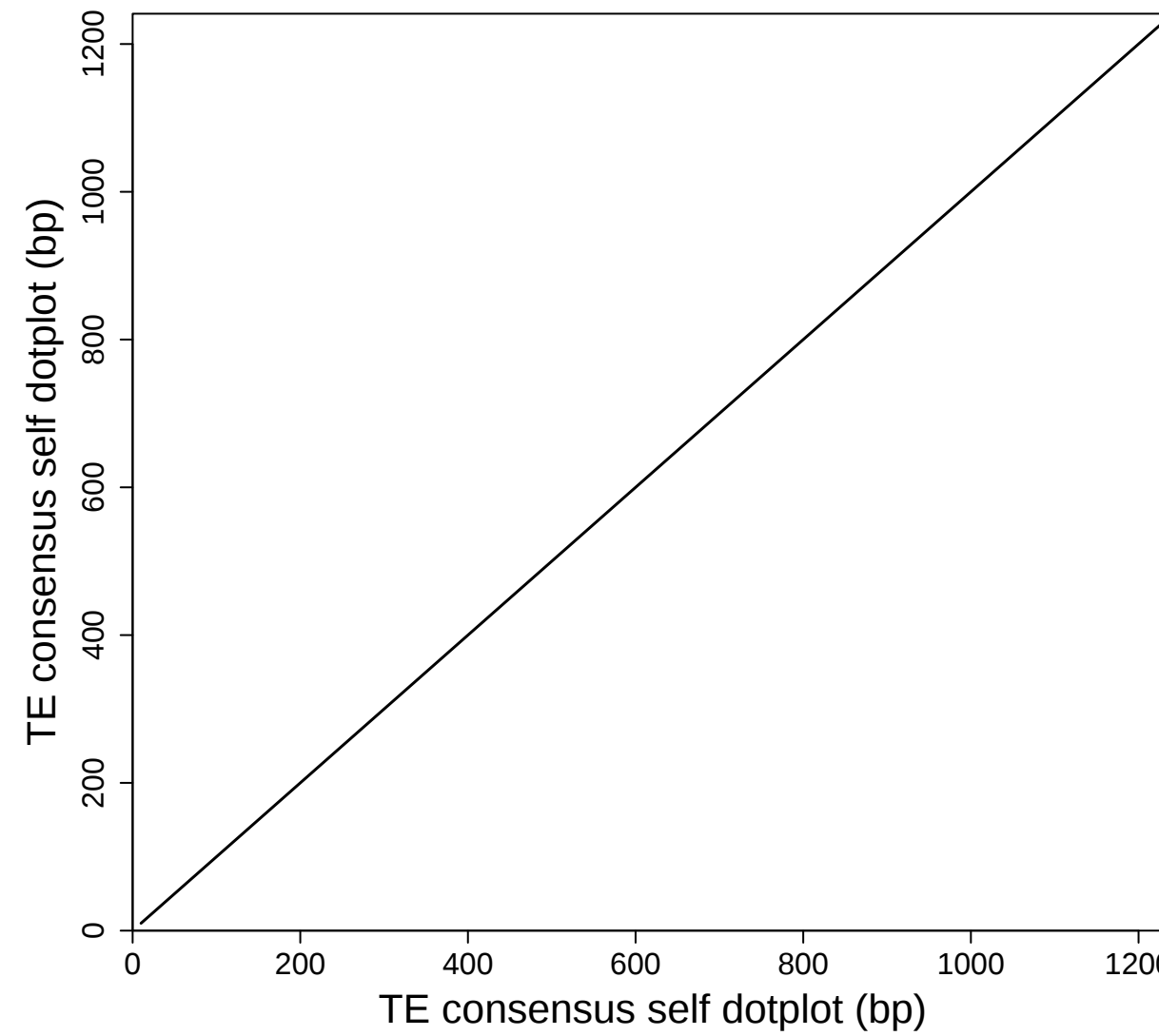
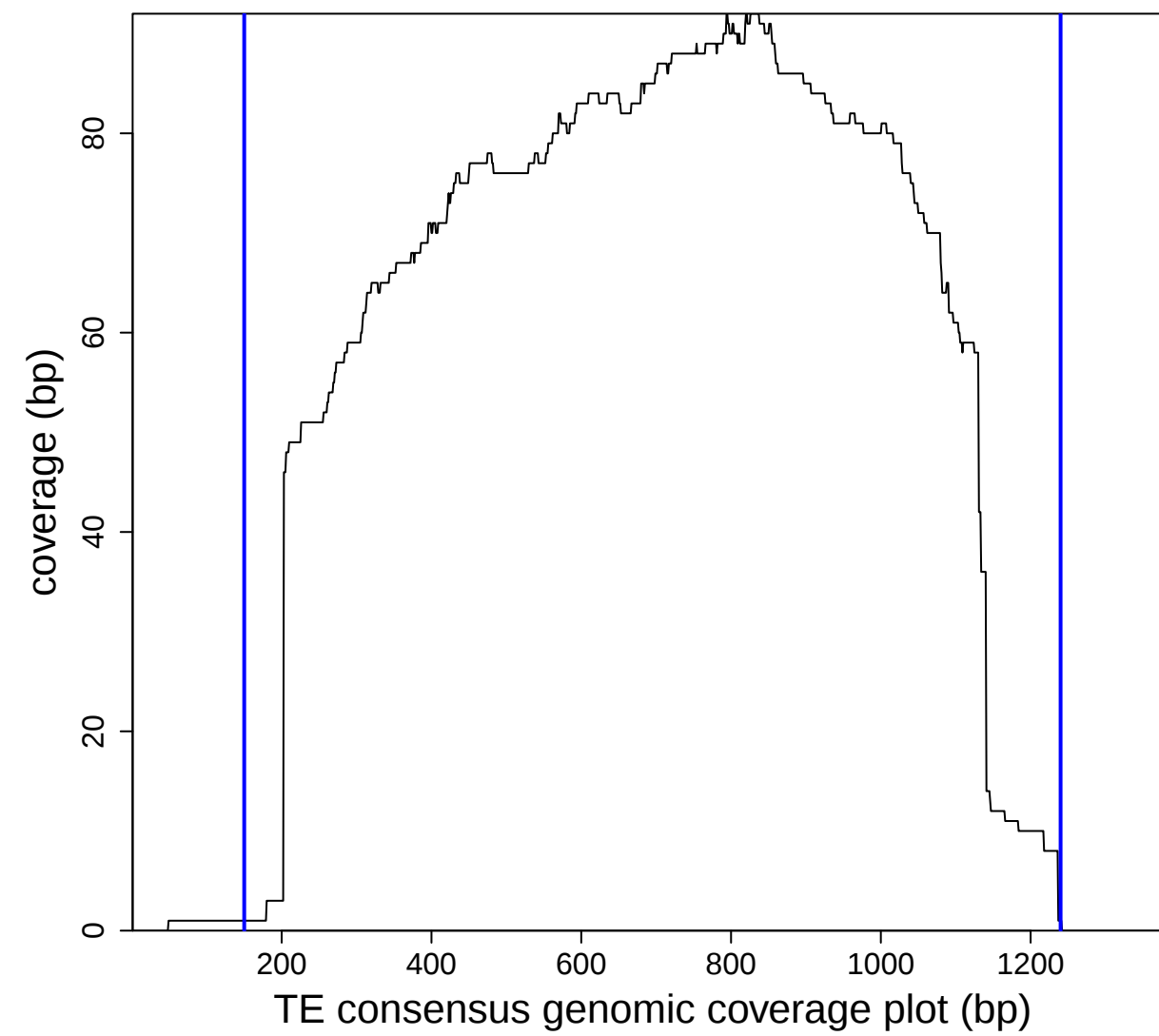
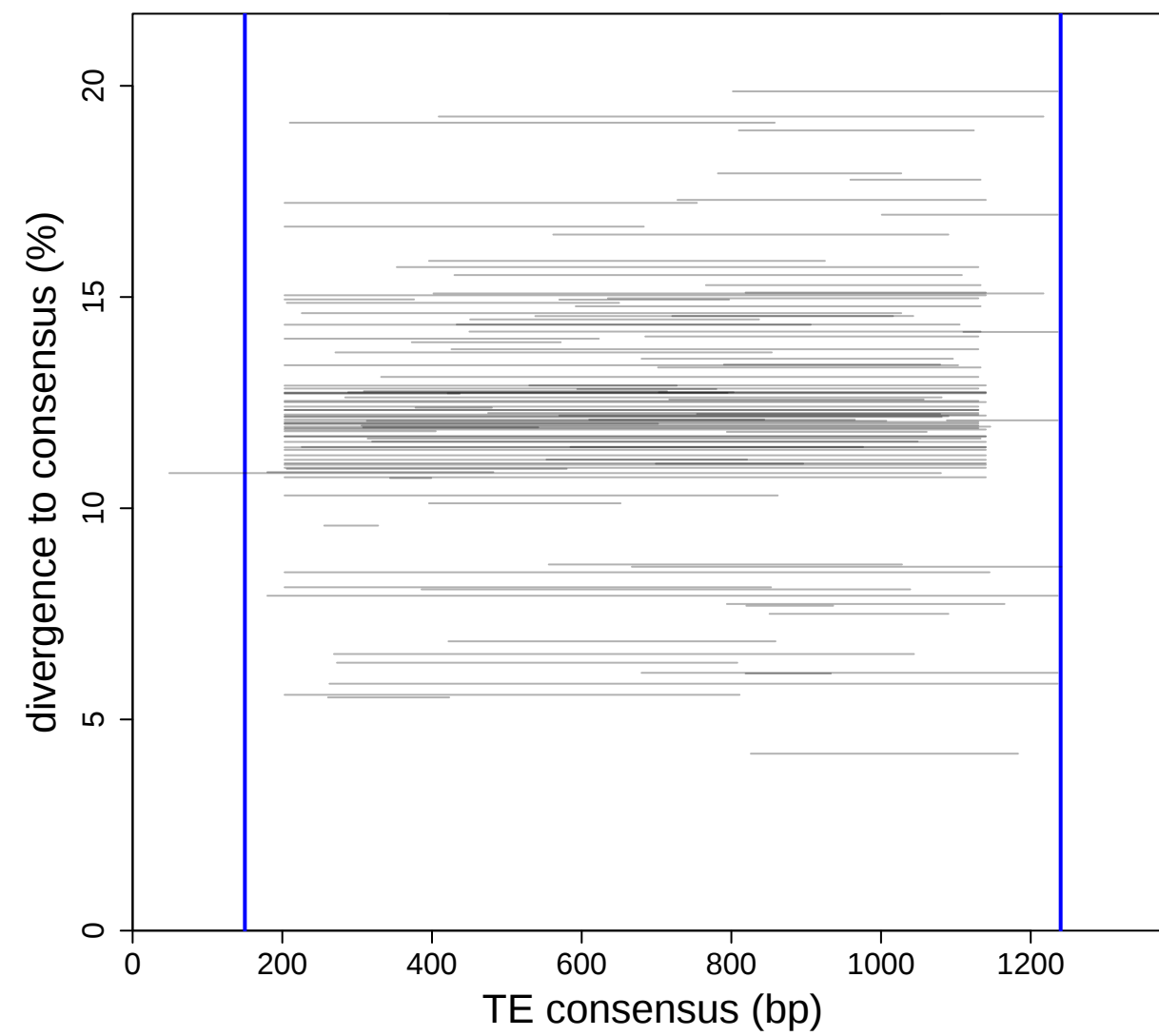
bed_uf.bed_g_2.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.fa_c
size: 1043bp; fragments: 140; full length: 15 (>=938.7bp)

After TEtrimmer 1043 bp

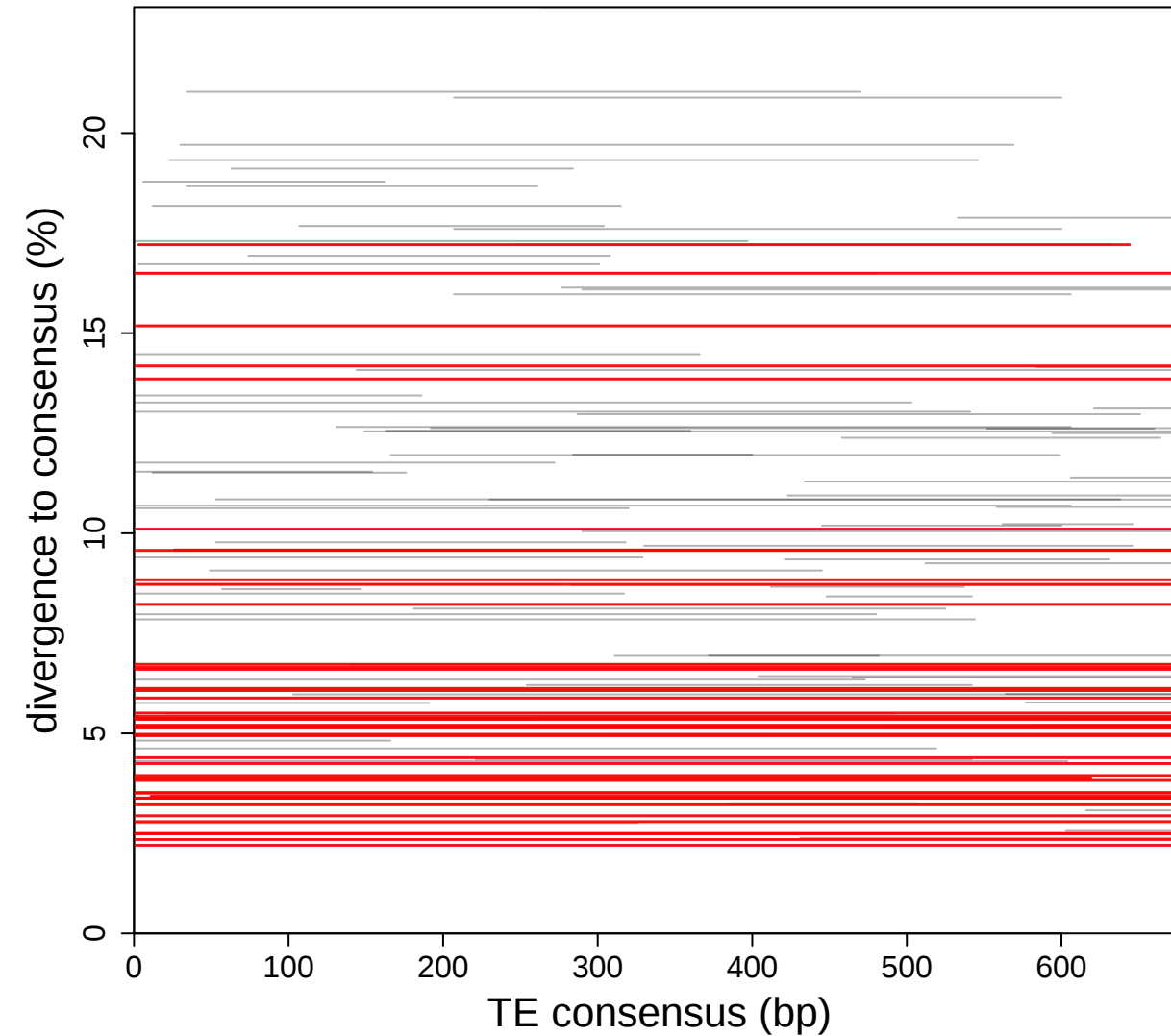


9.fasta.b.bed_uf.bed_g_2.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.f
size: 1390bp; fragments: 123; full length: 0 (>=1251bp)

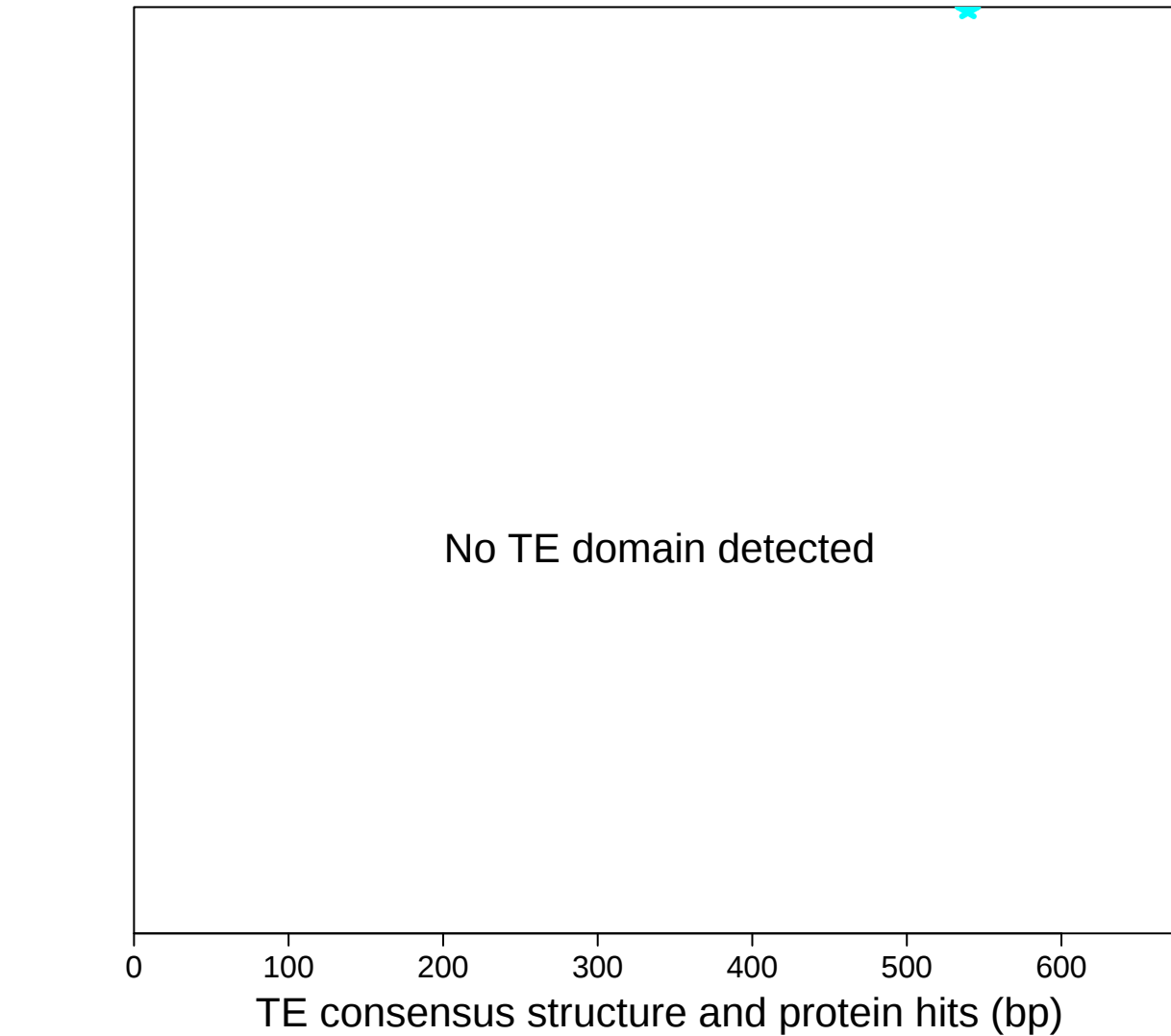
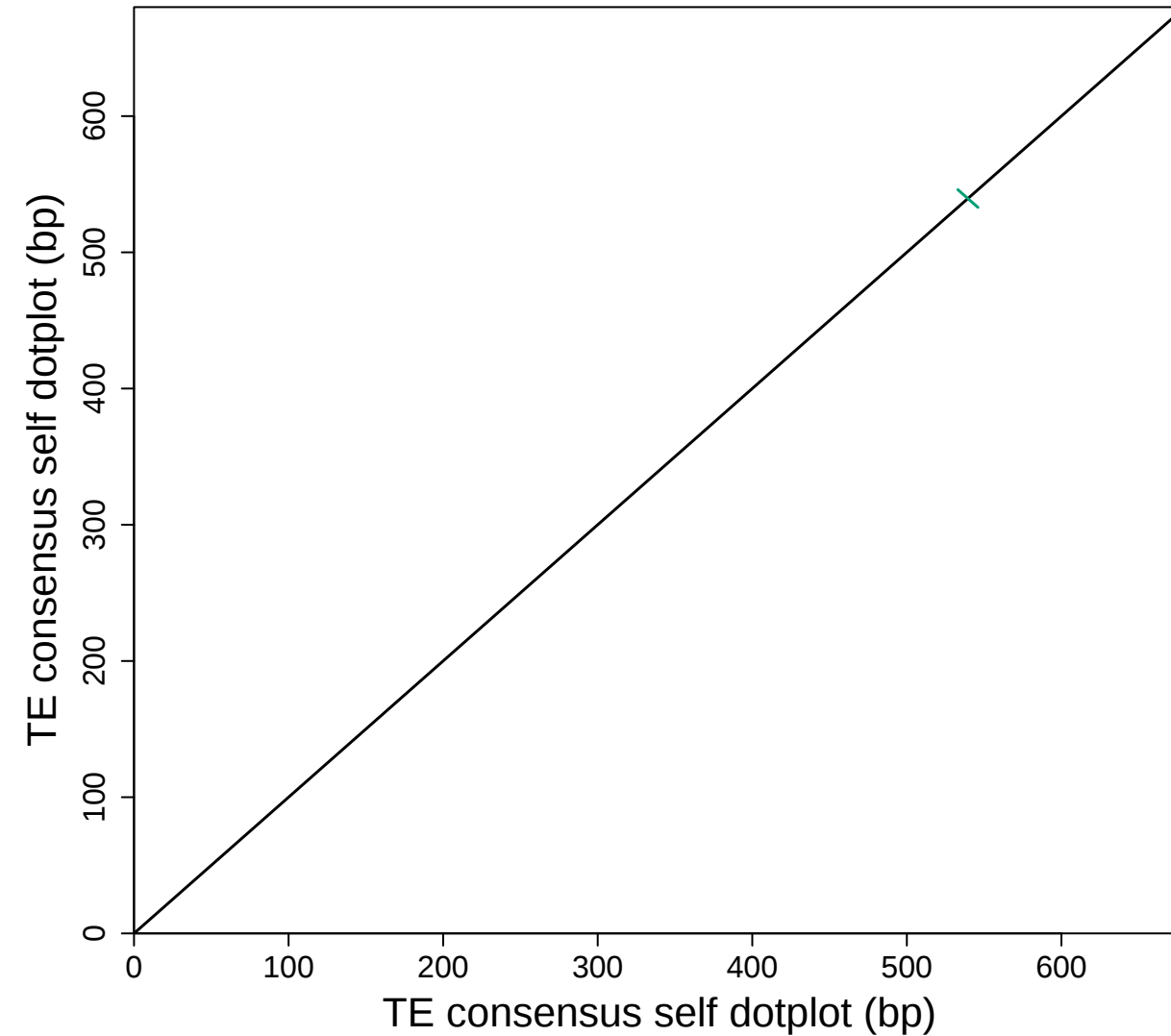
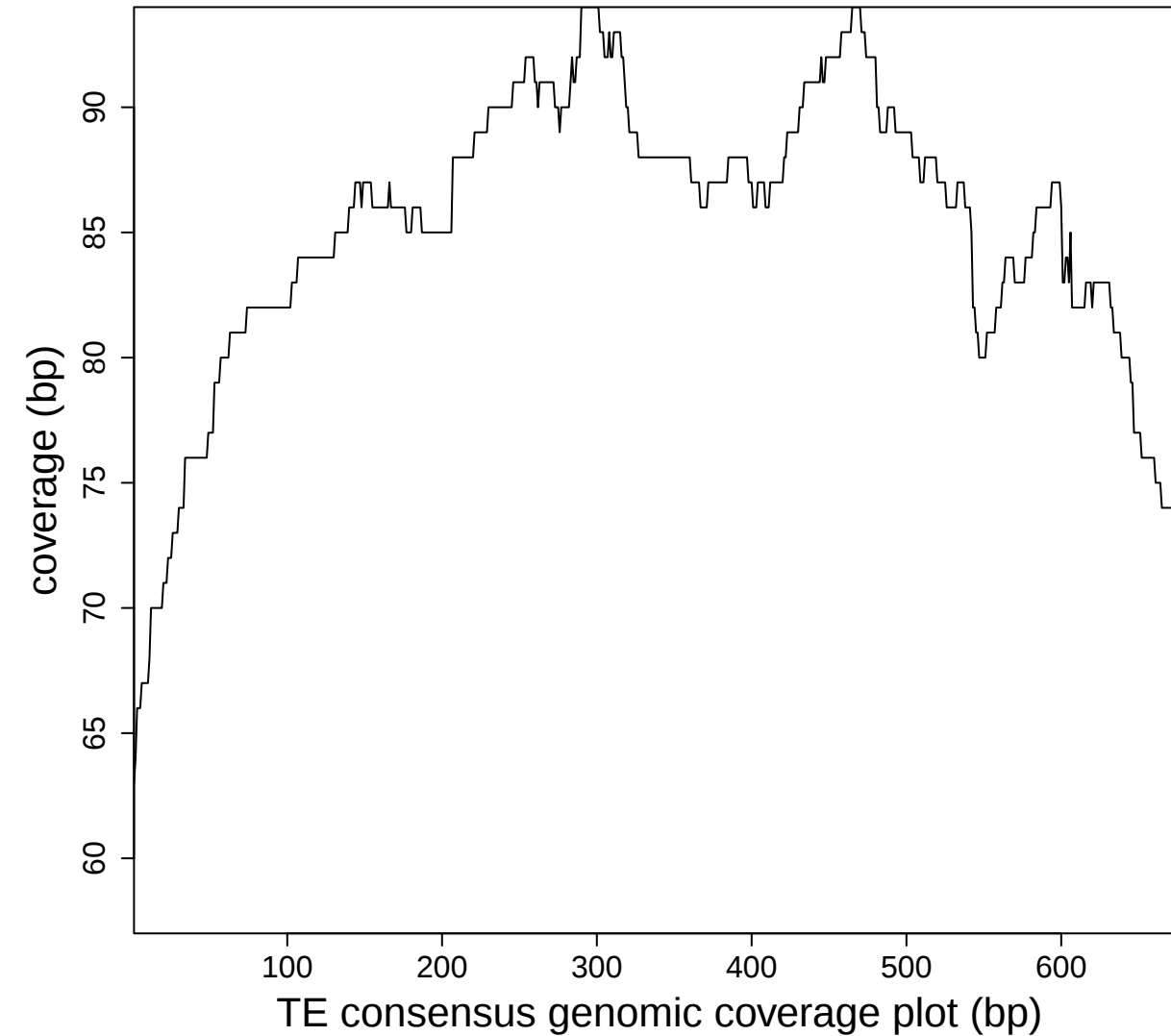
After TEtrimmer Extended plot Blue lines are boundaries



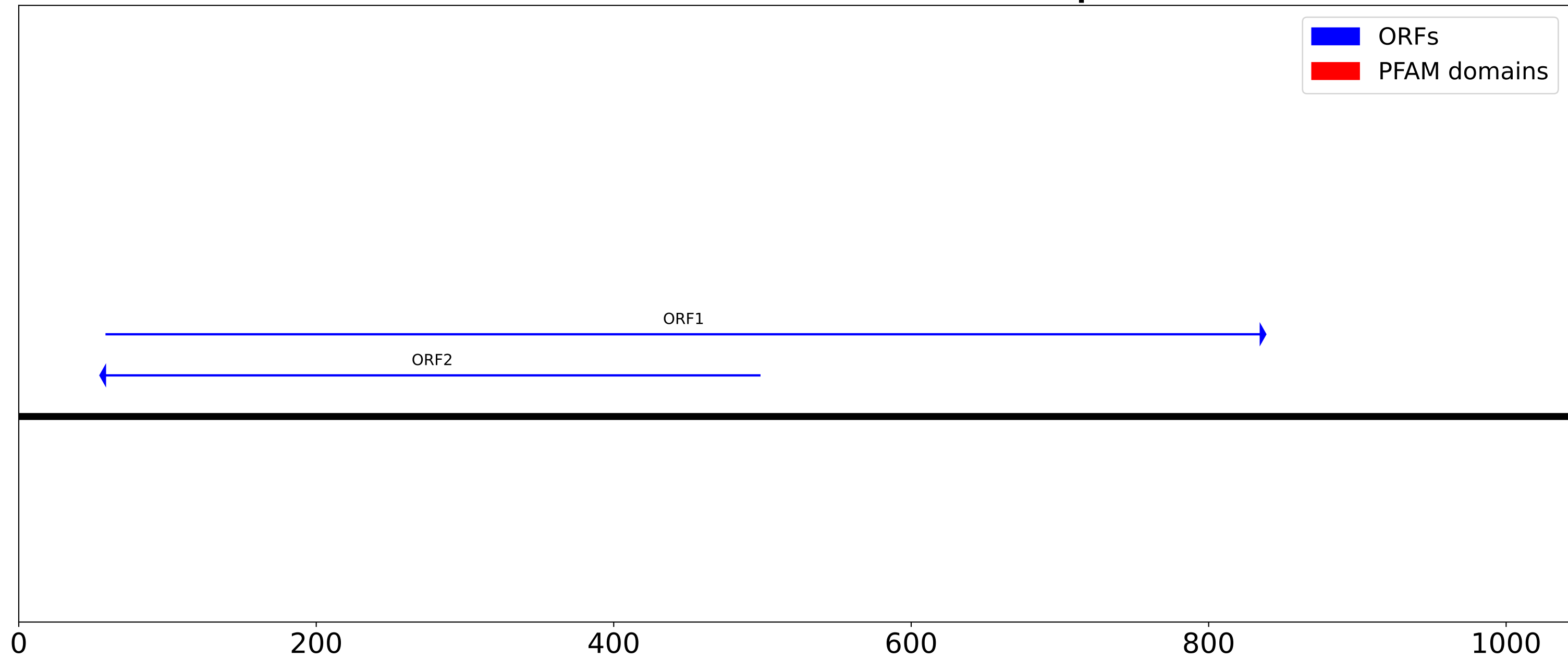
TE: rnd_5_family_8819
size: 680bp; fragments: 137; full length: 45 (>=612bp)



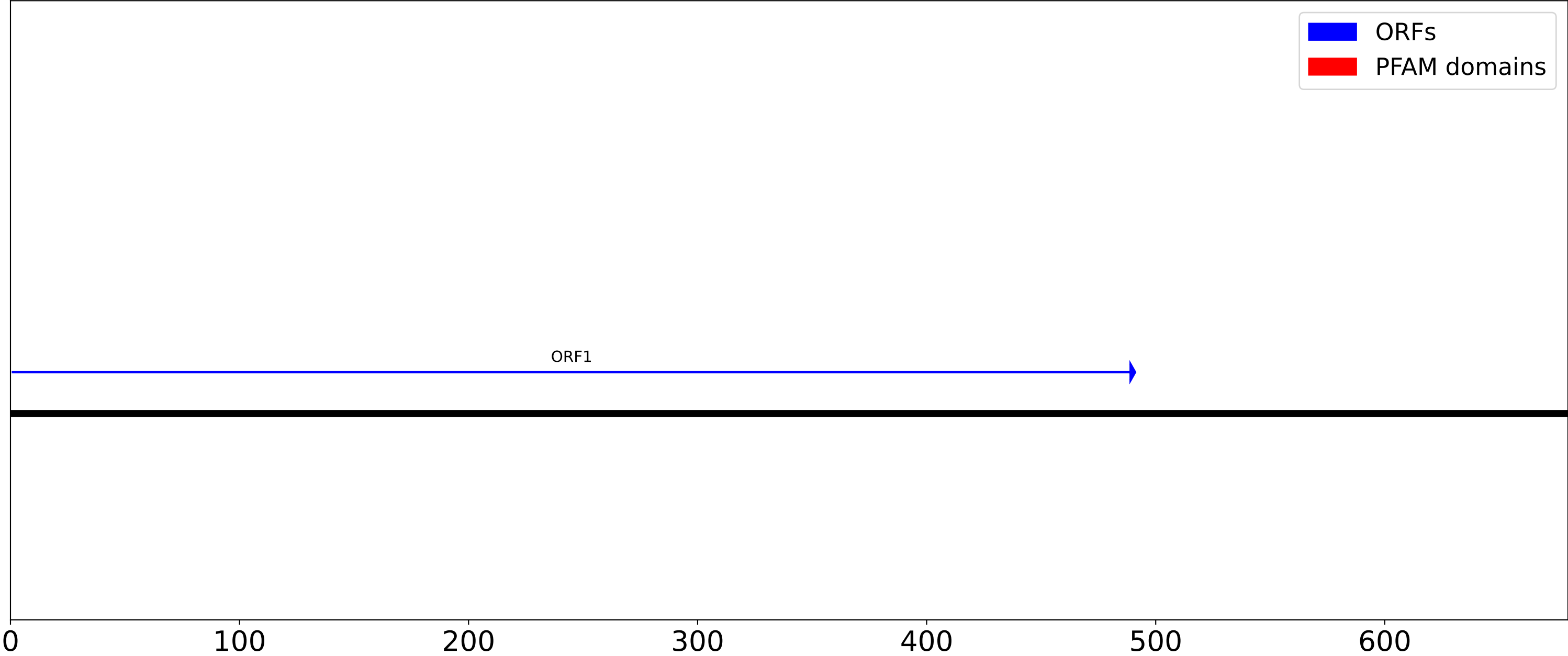
Before TEtrimmer 680 bp



After TEtrimmer ORF and PFAM domain plot



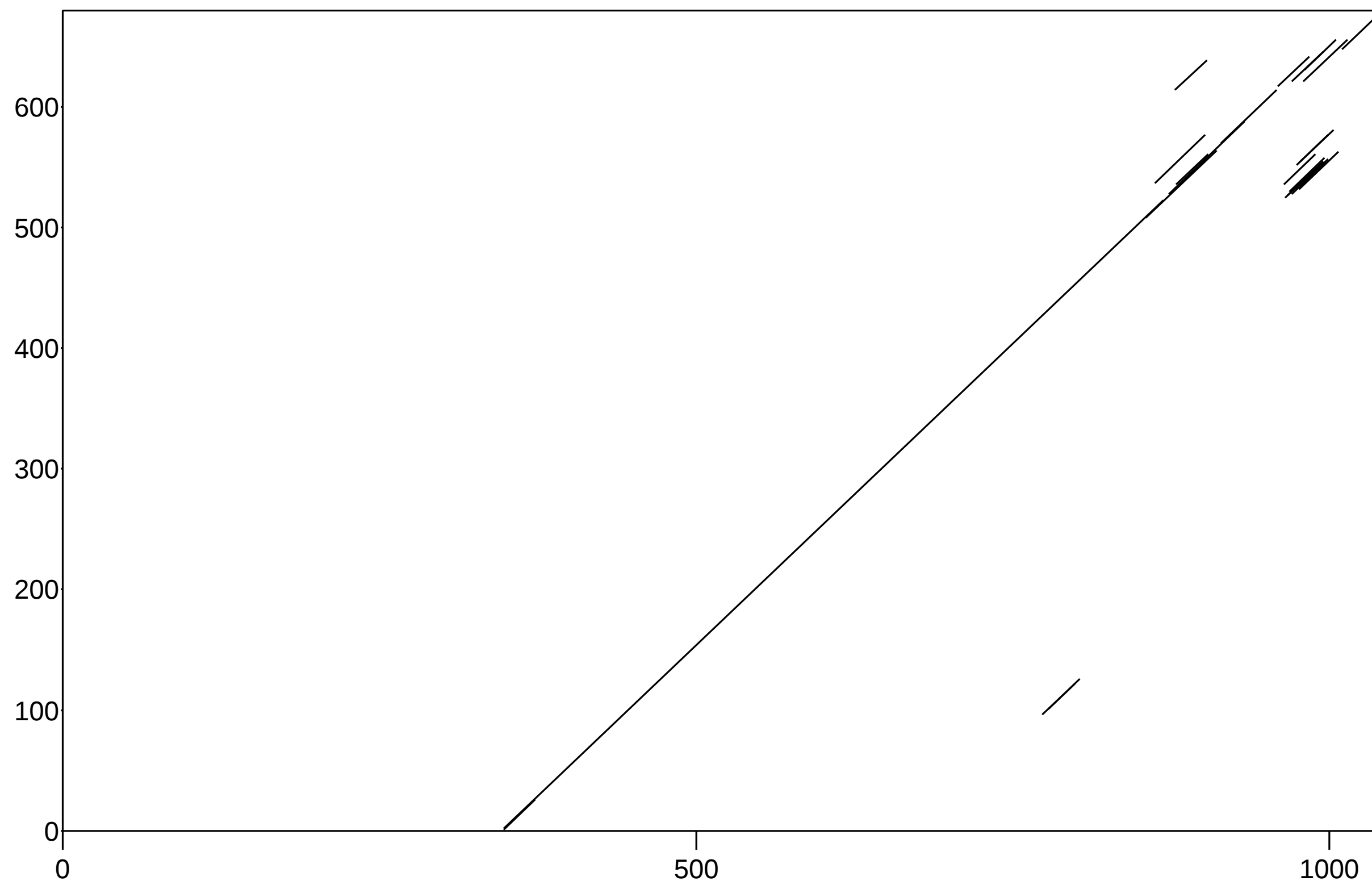
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)