

Start crop Point



End crop Point



MSA length = 1294



Start crop Point

End crop Point

0 -

10

0

250

500

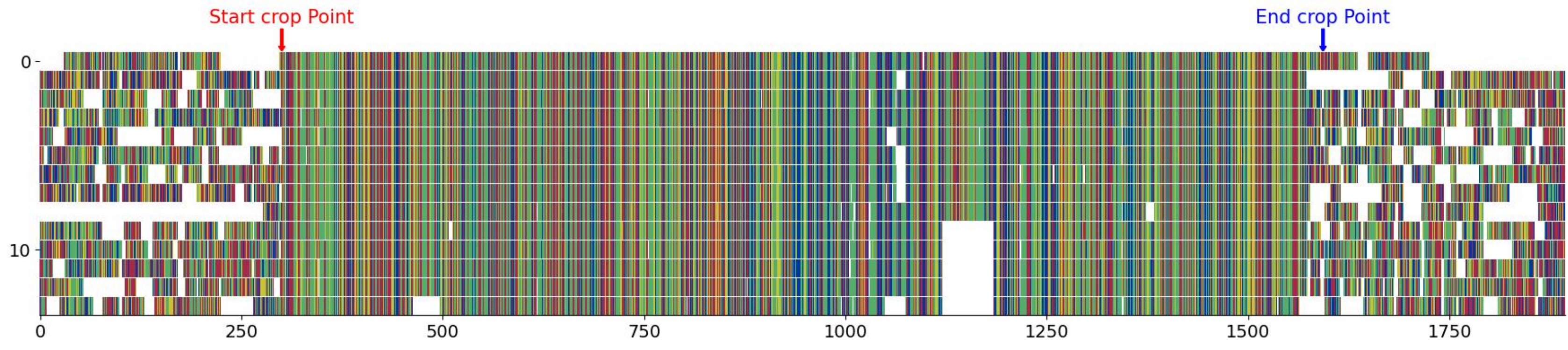
750

1000

1250

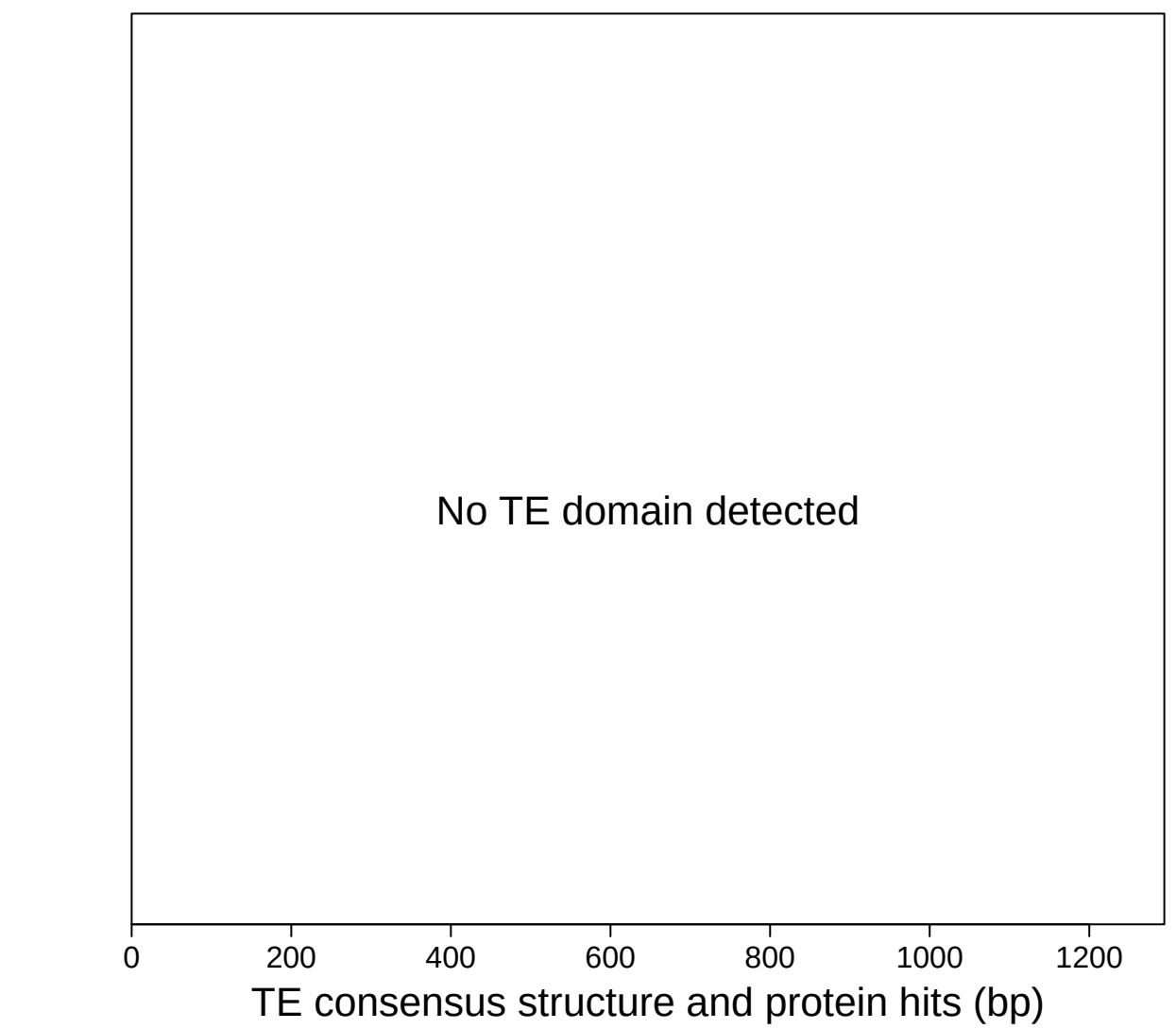
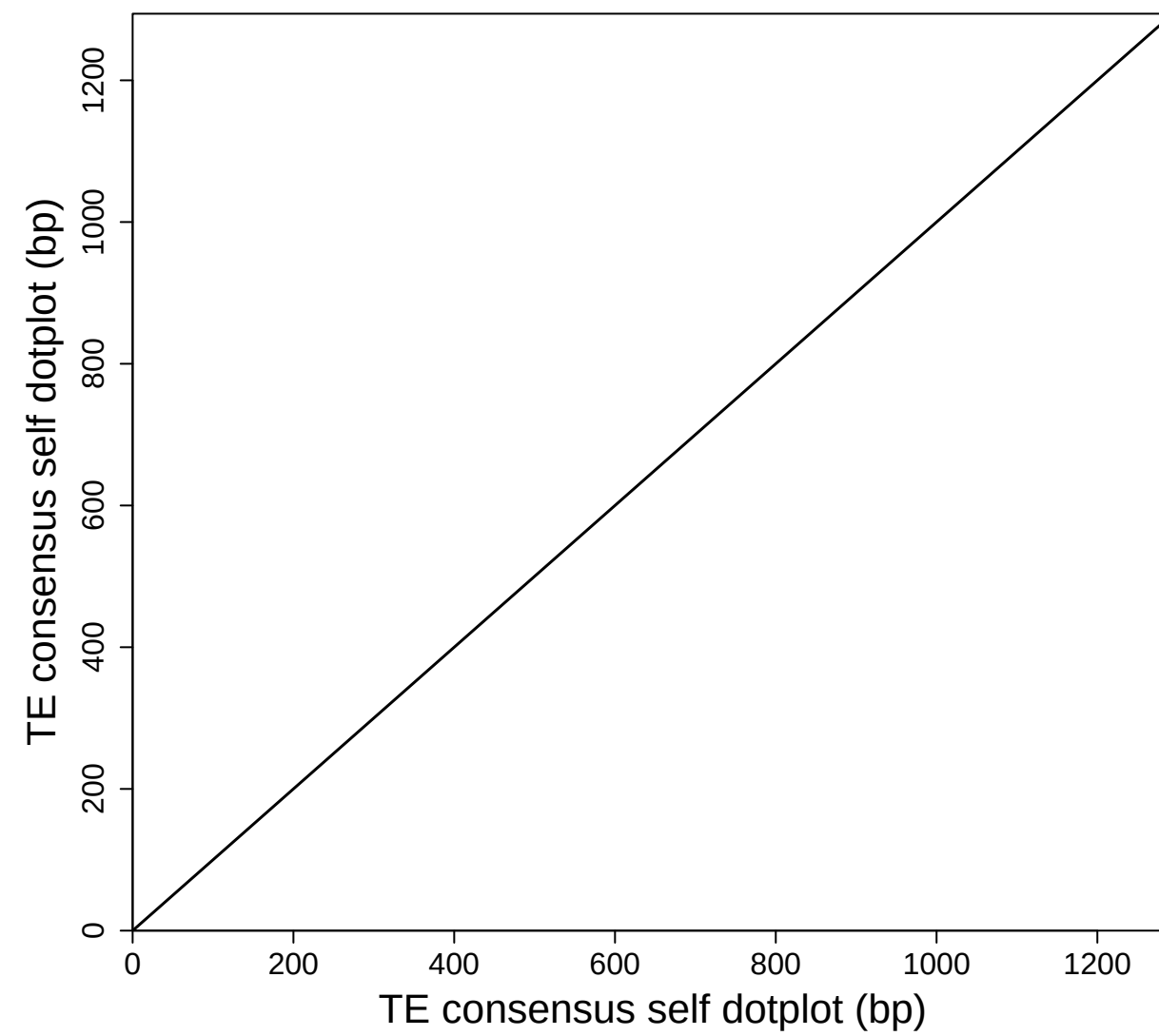
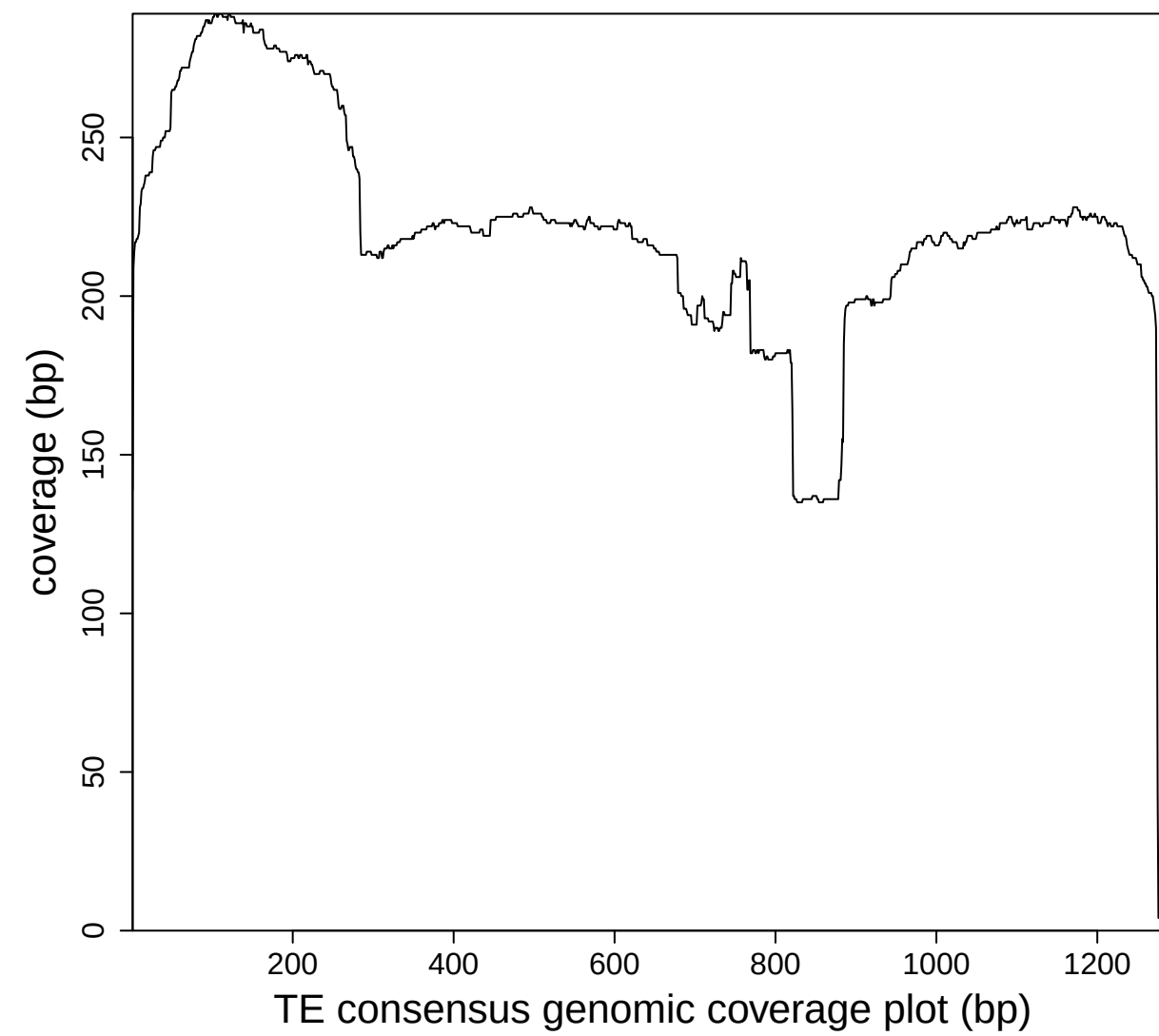
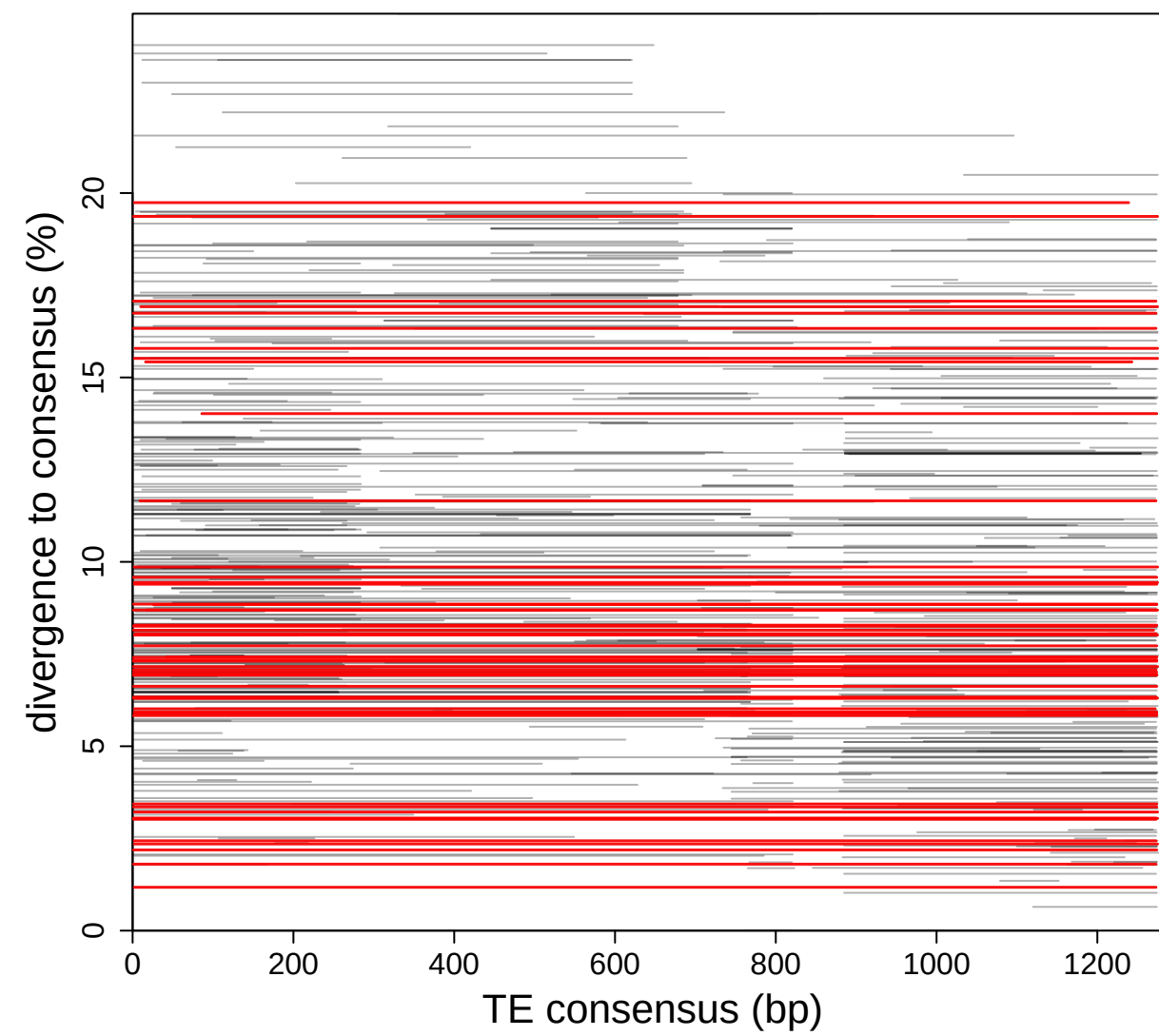
1500

1750



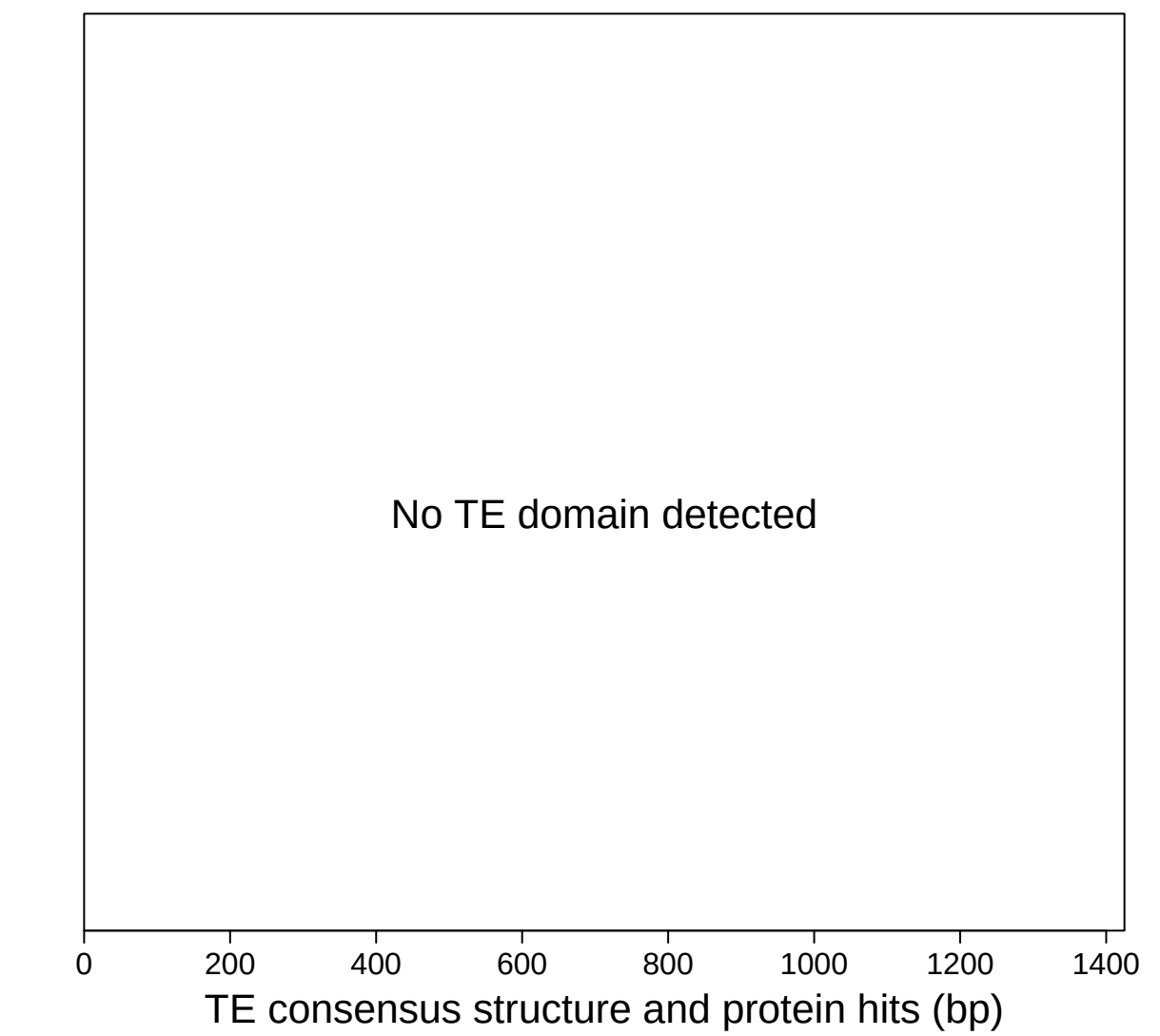
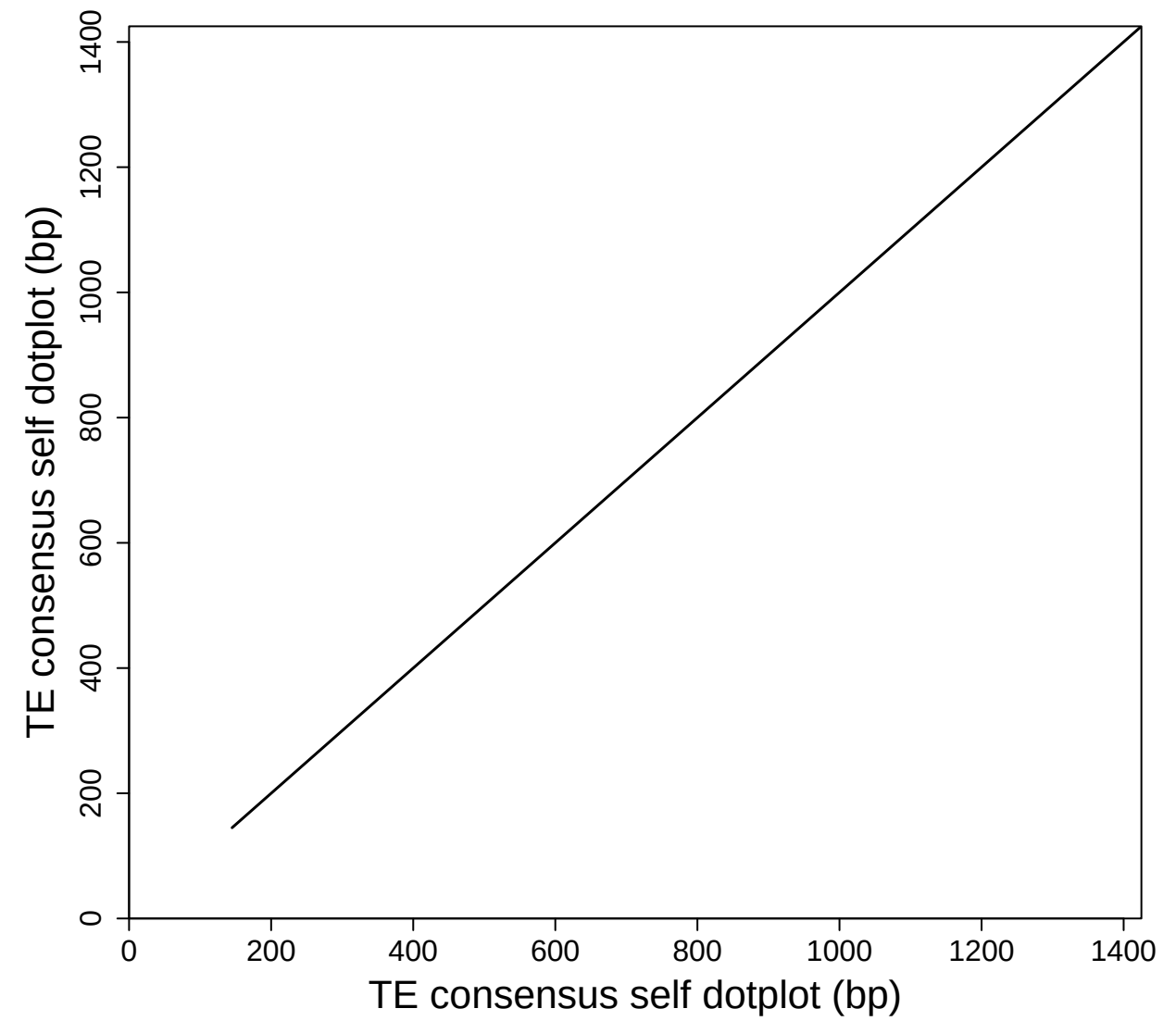
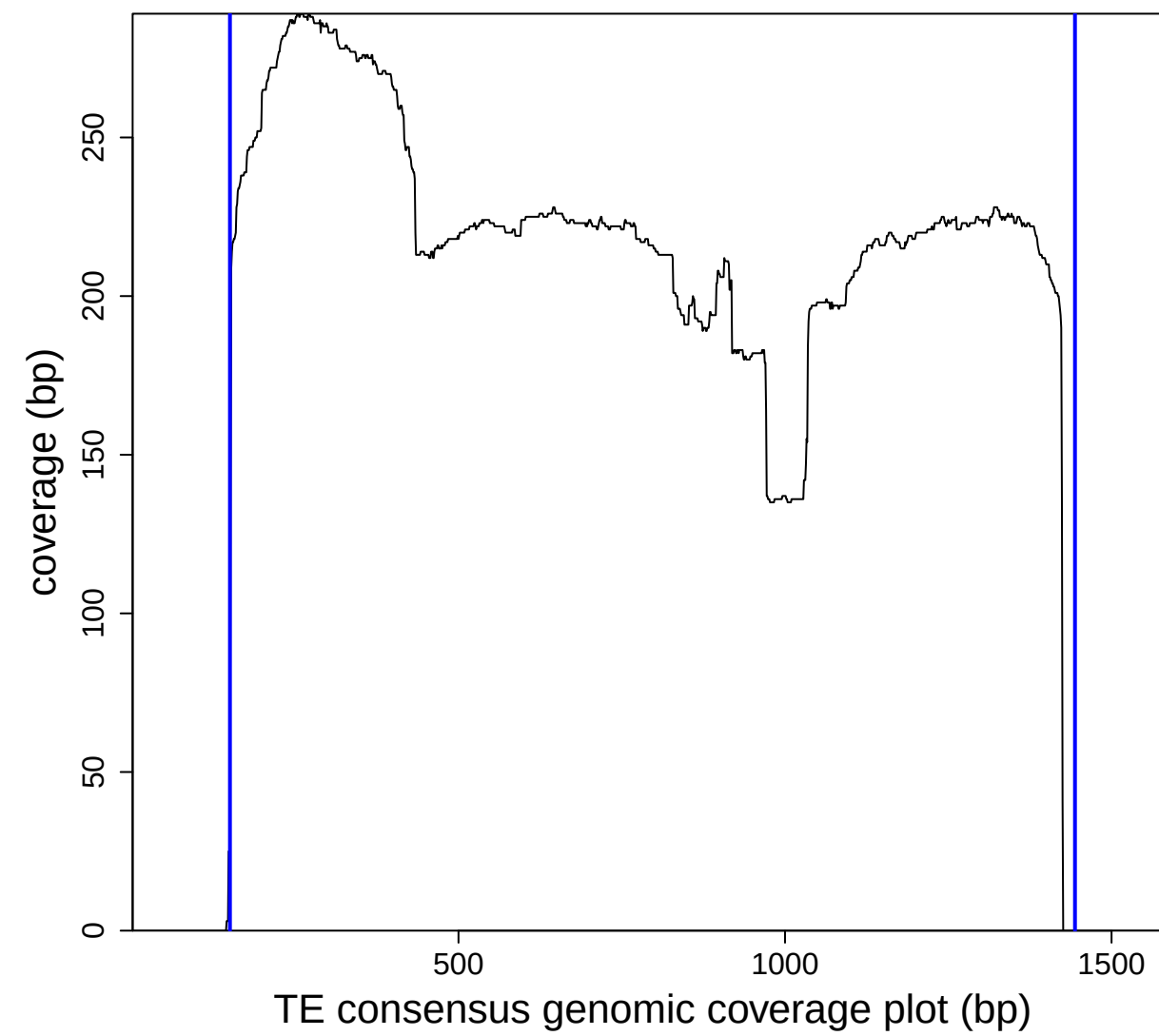
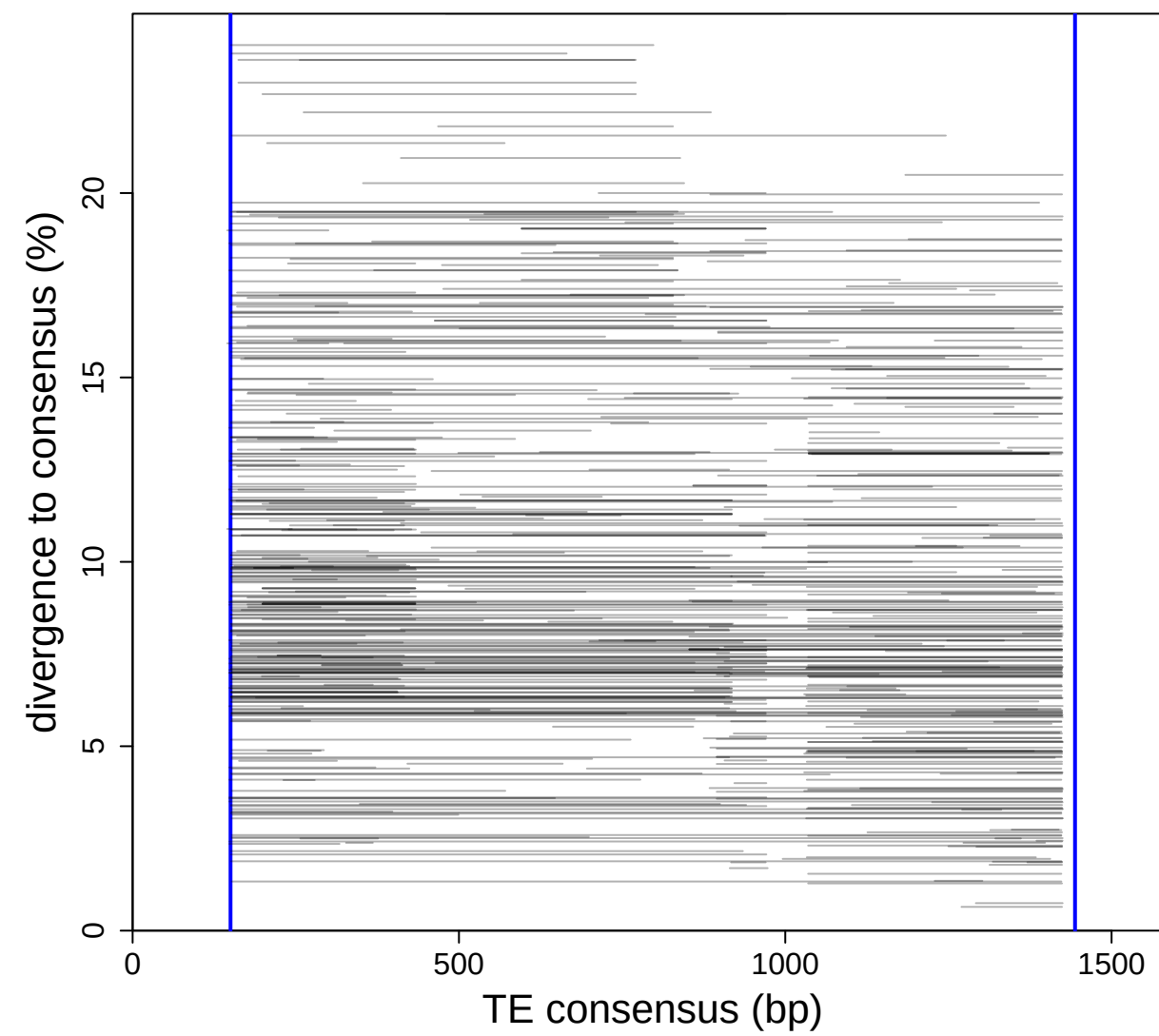
.l.b.bed uf.bed g_2.bed fm_1.bed 0_0_bcln.fa aln.fa cl.fa gs.fa
size: 1294bp; fragments: 565; full length: 53 (>=1164.6bp)

After TEtrimmer 1294 bp



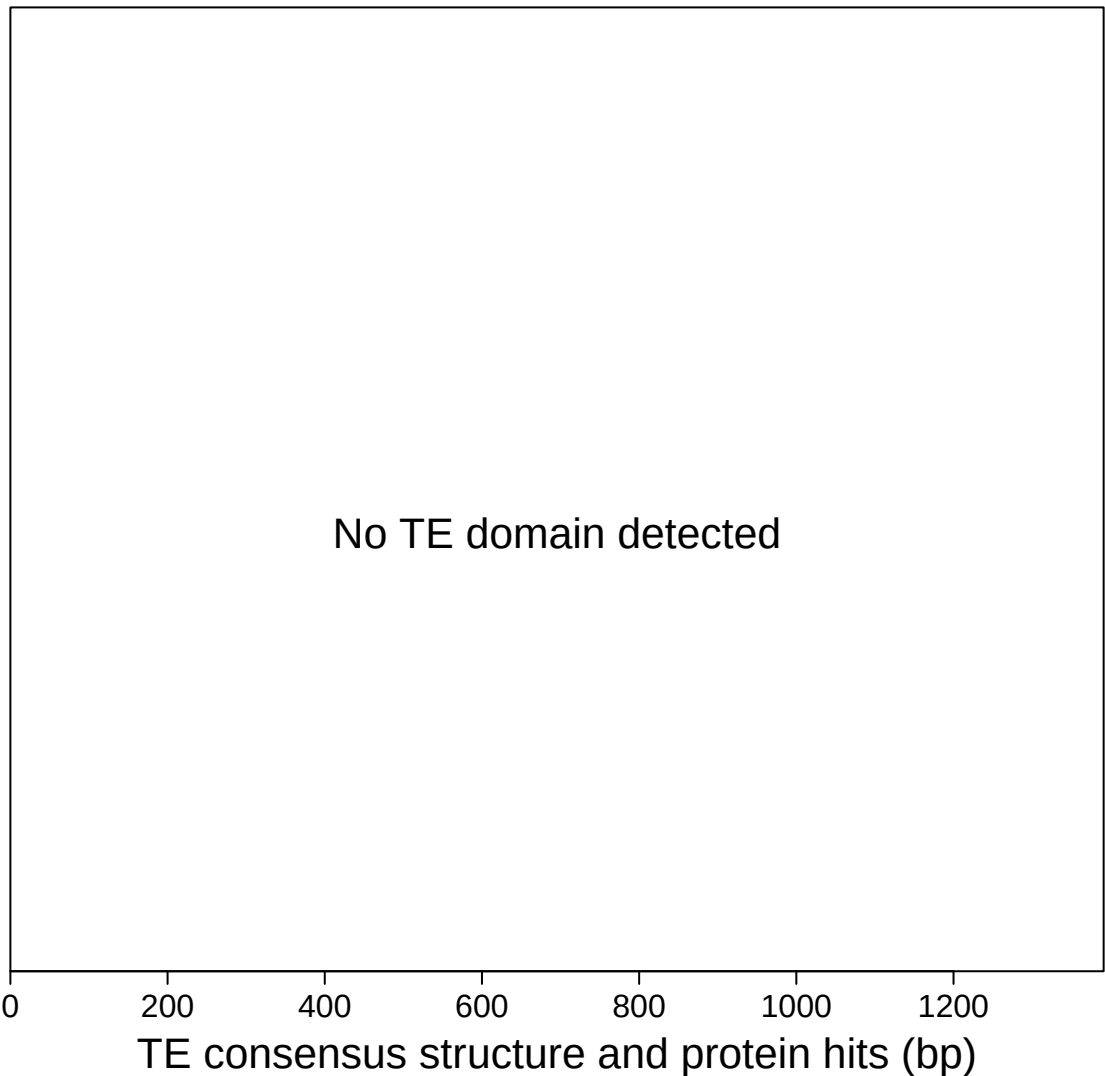
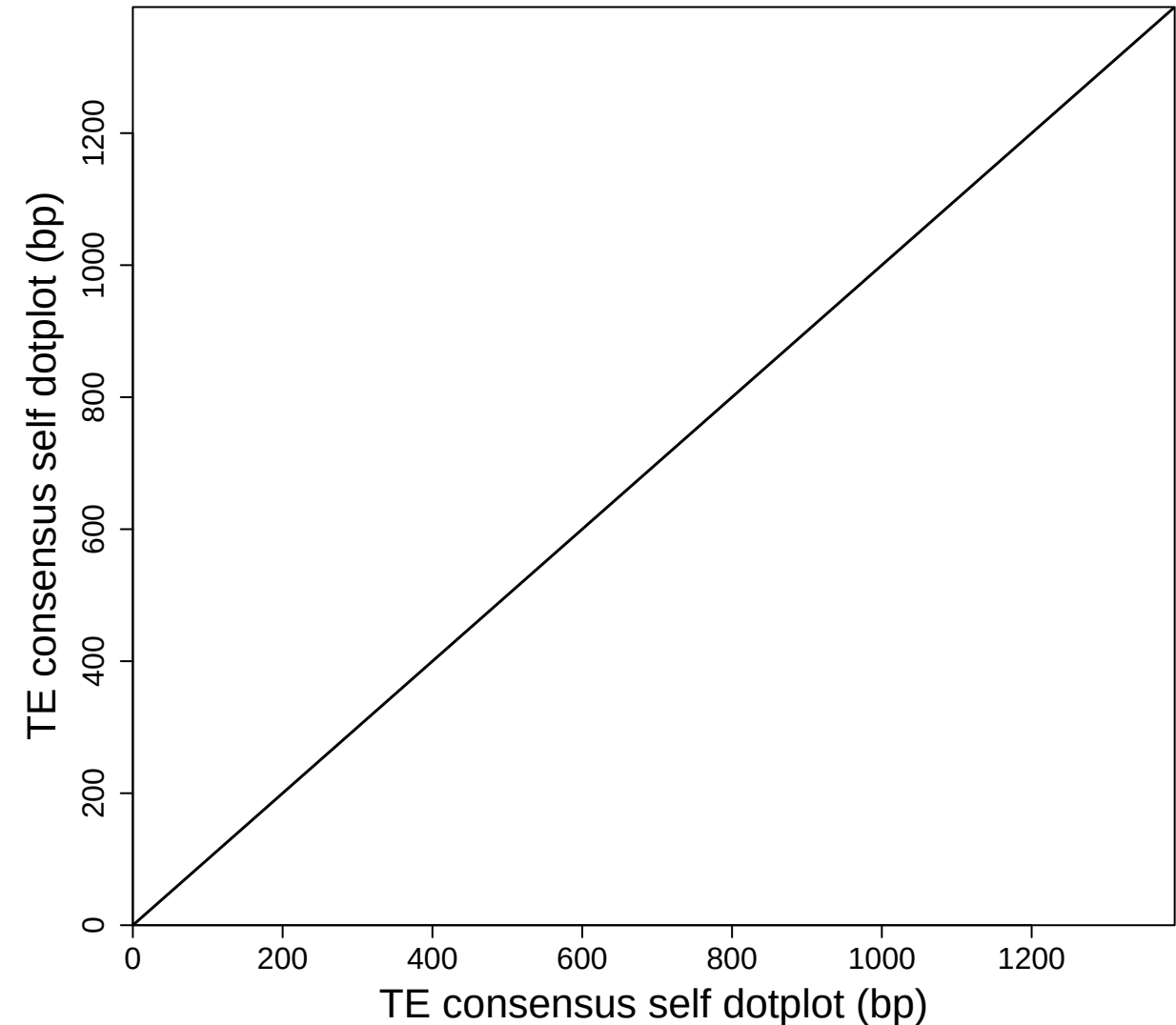
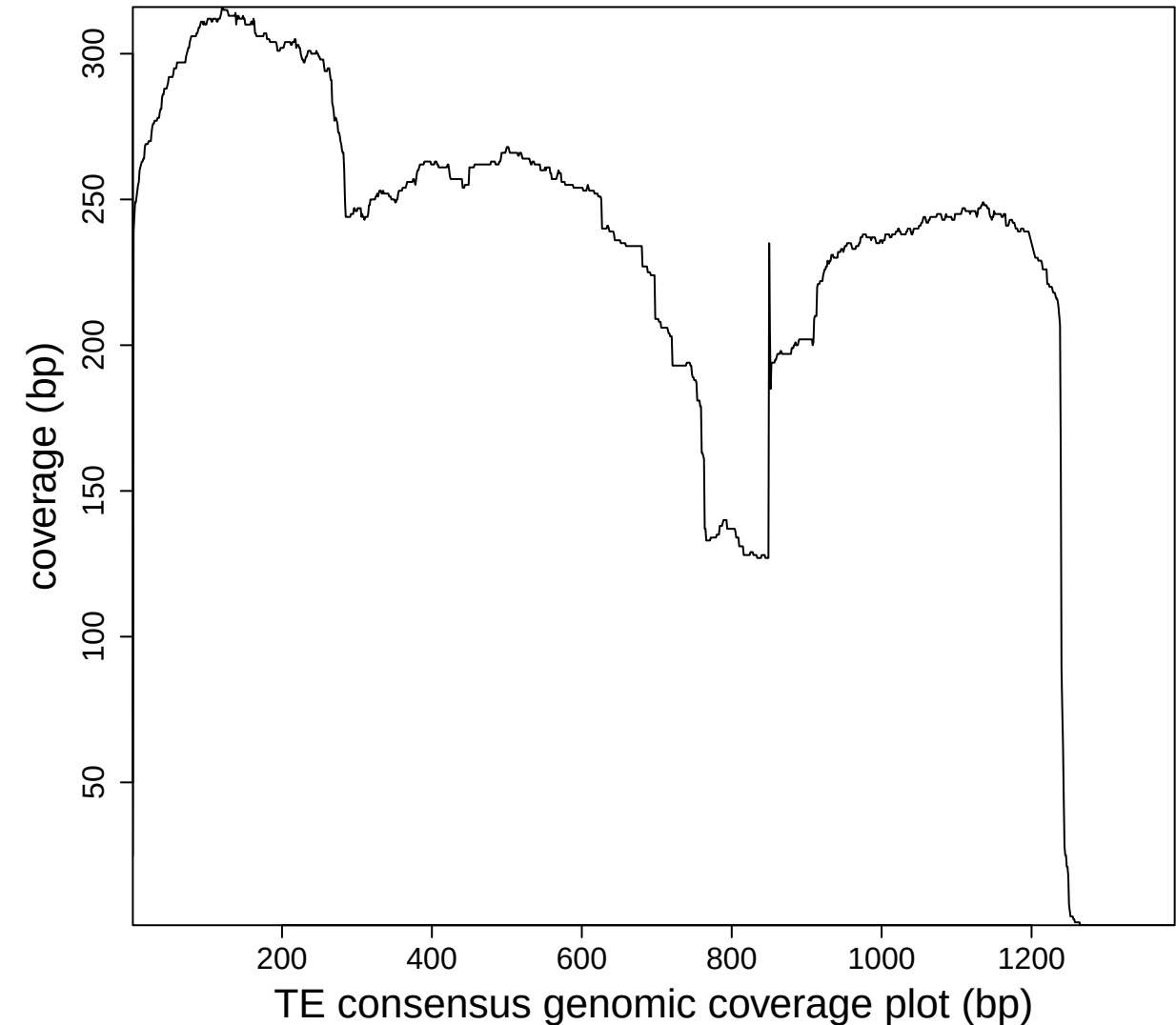
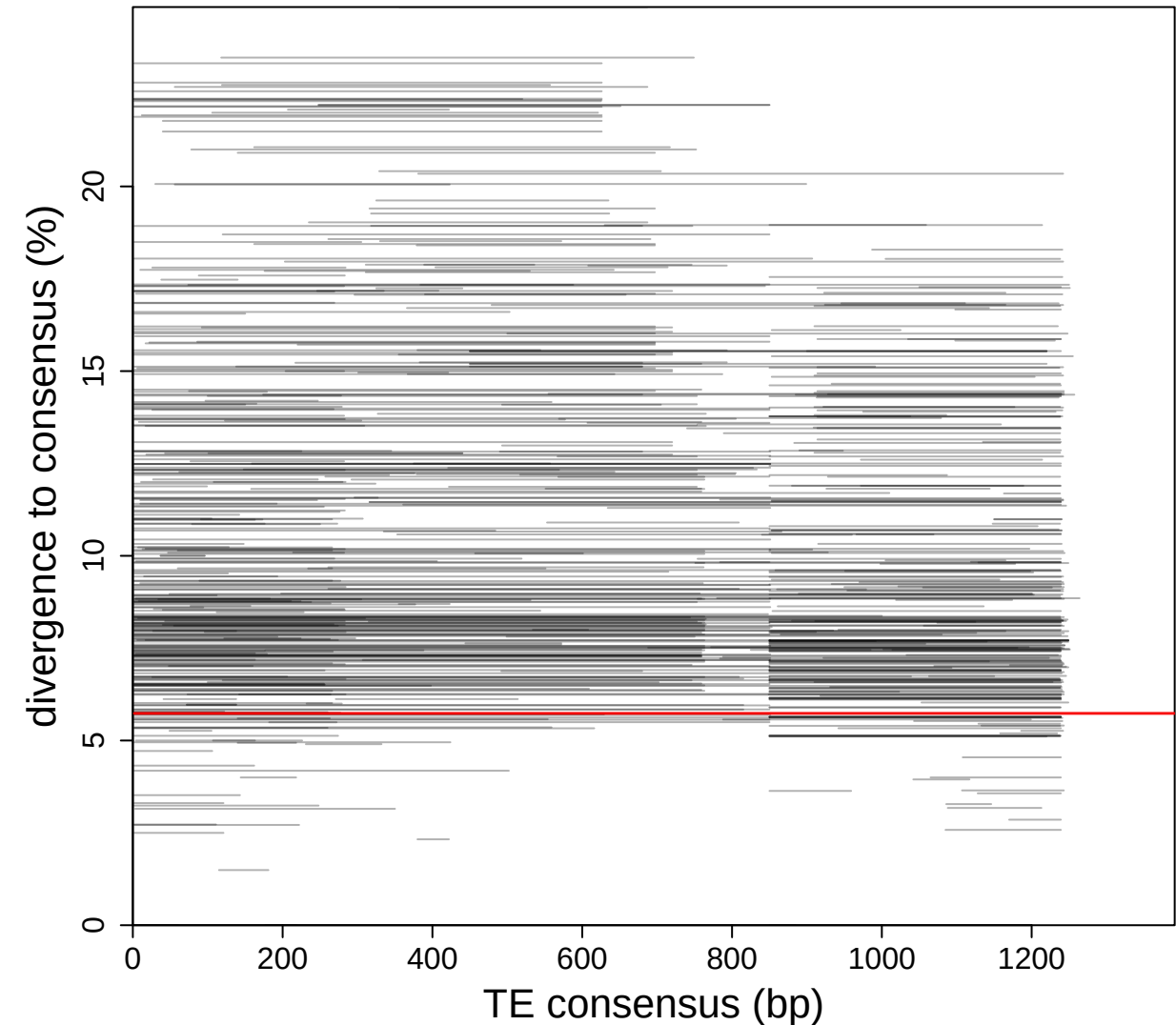
0.fasta.b.bed uf.bed g 2.bed fm 1.bed 0 0 bcln.fa aln.fa cl.fa
size: 1594bp; fragments: 565; full length: 0 (>=1434.6bp)

After TEtrimmer Extended plot Blue lines are boundaries

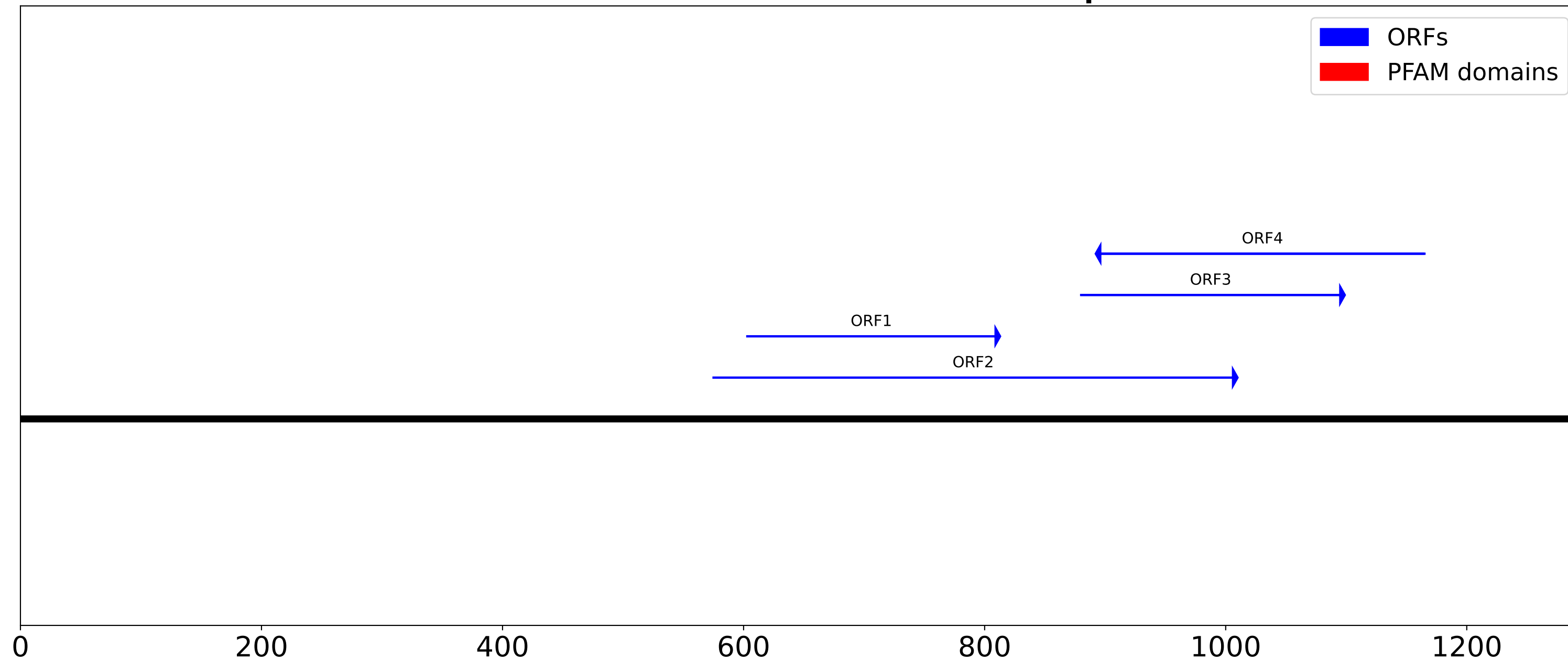


TE: rnd_5_family_330
size: 1391bp; fragments: 662; full length: 1 (>=1251.9bp)

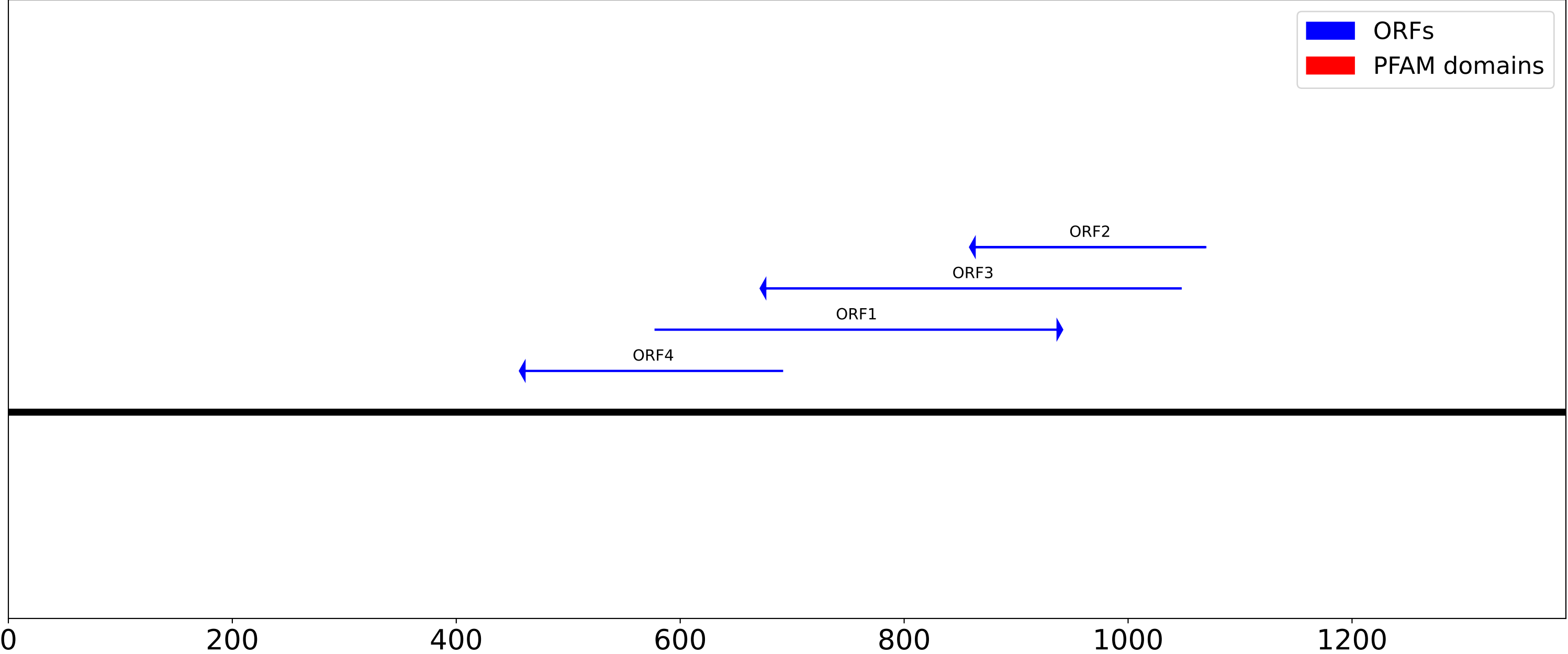
Before TEtrimmer 1391 bp



After TEtrimmer ORF and PFAM domain plot



Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

