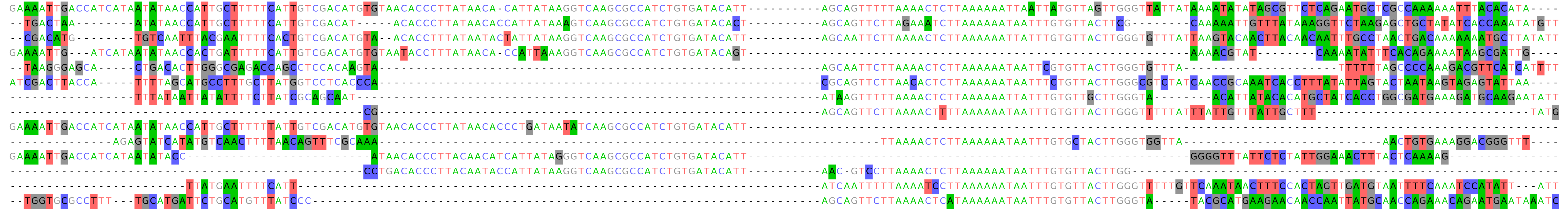


Start crop Point

End crop Point

MSA length = 949



Start crop Point

End crop Point

0

10

0

200

400

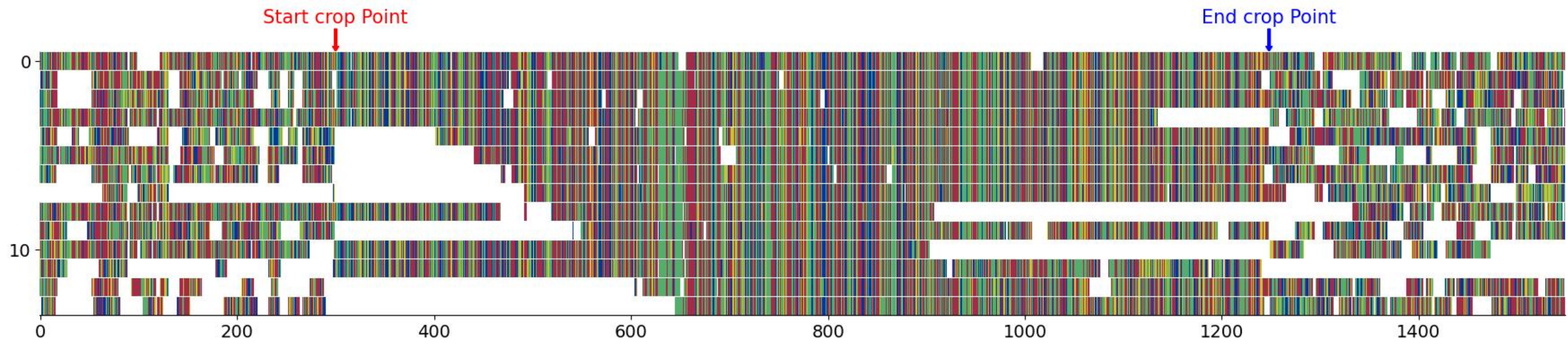
600

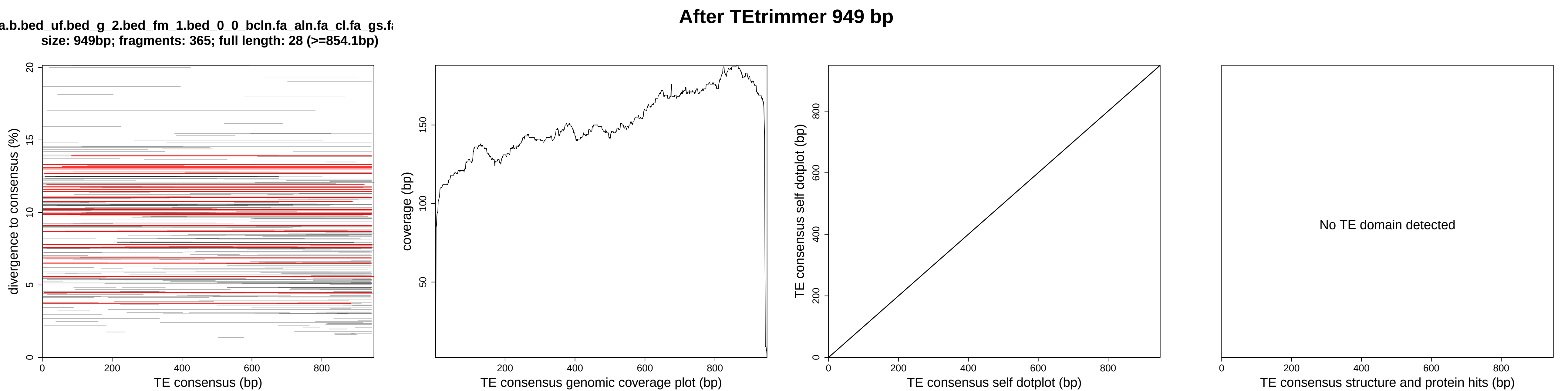
800

1000

1200

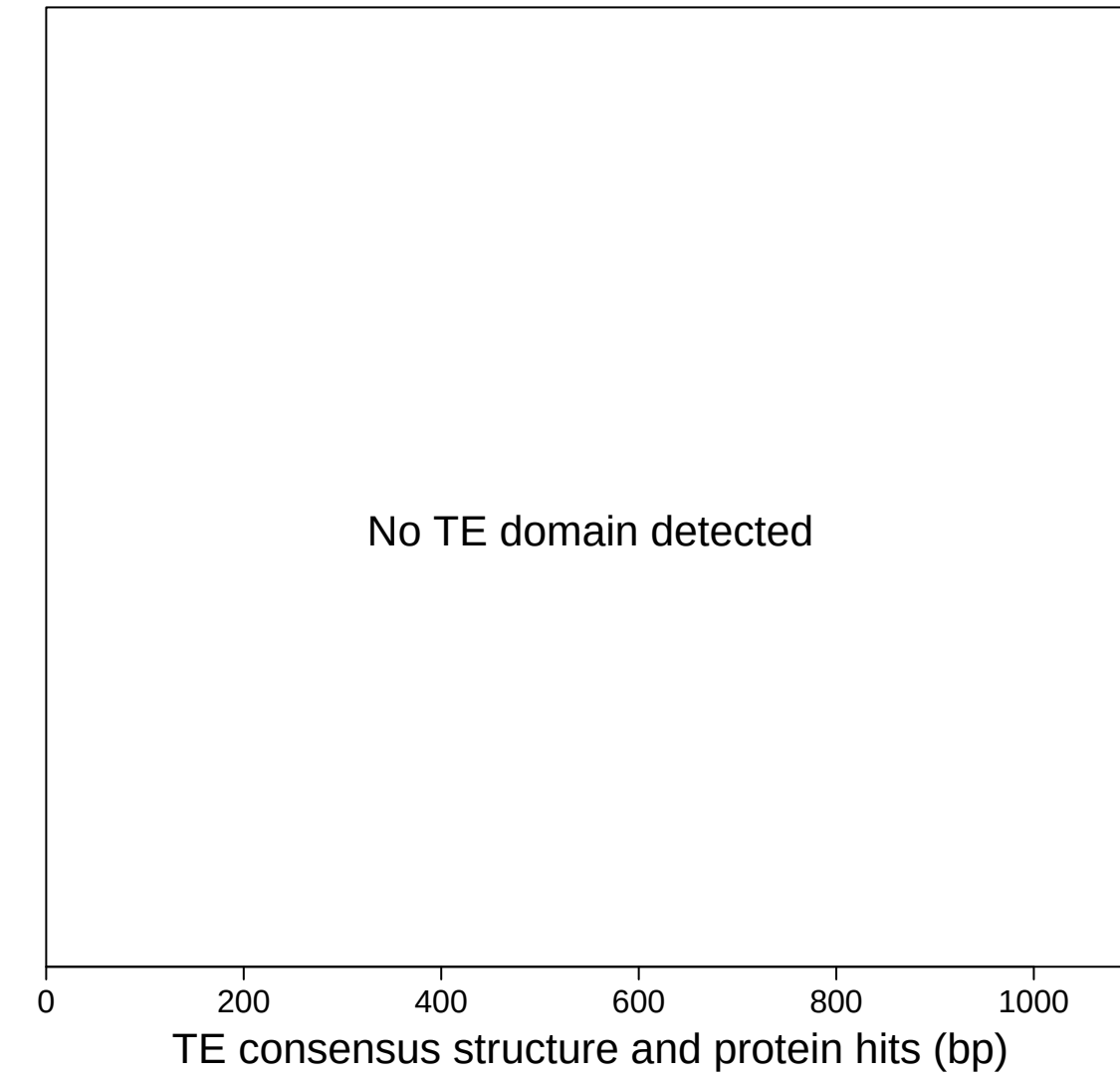
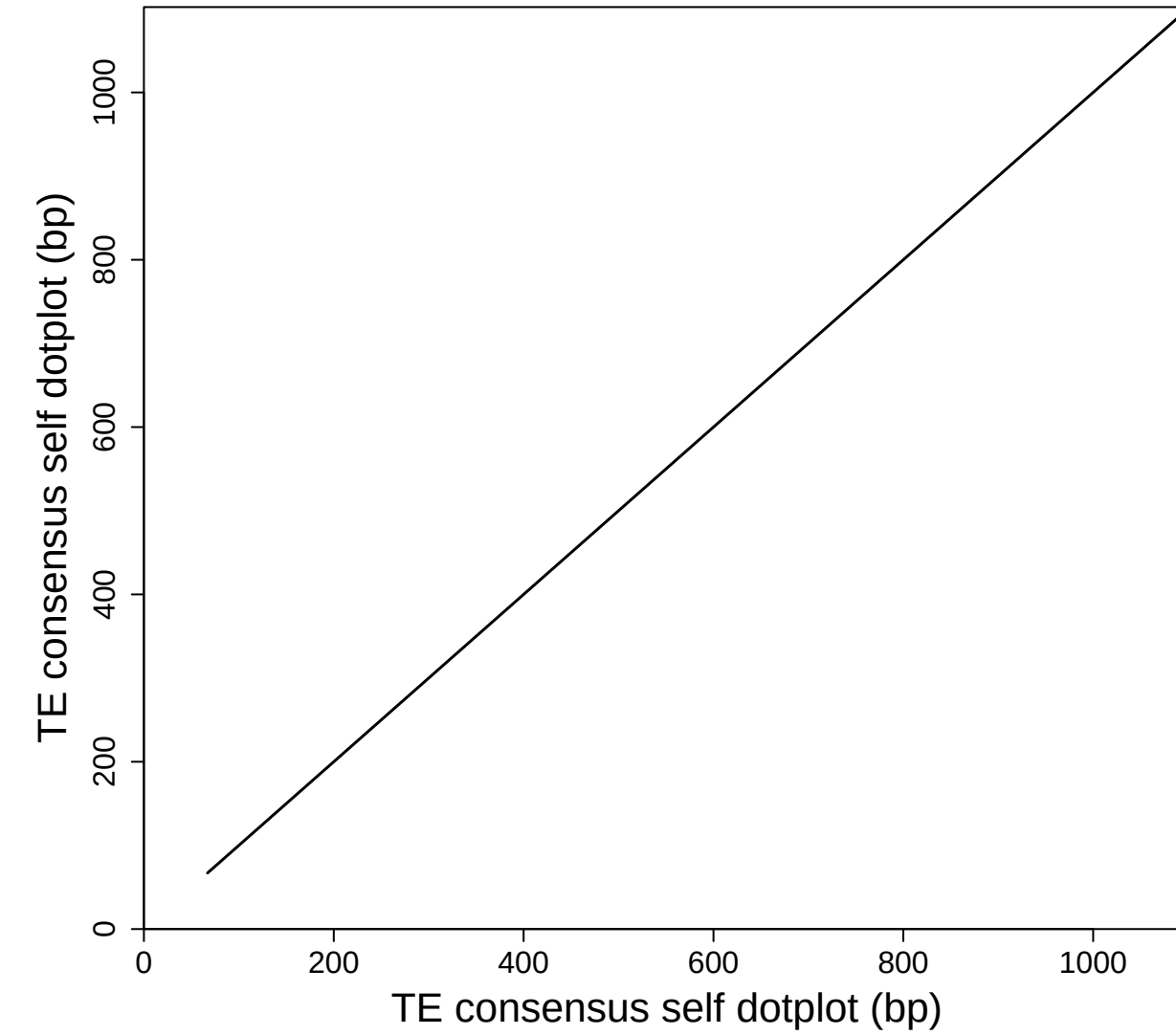
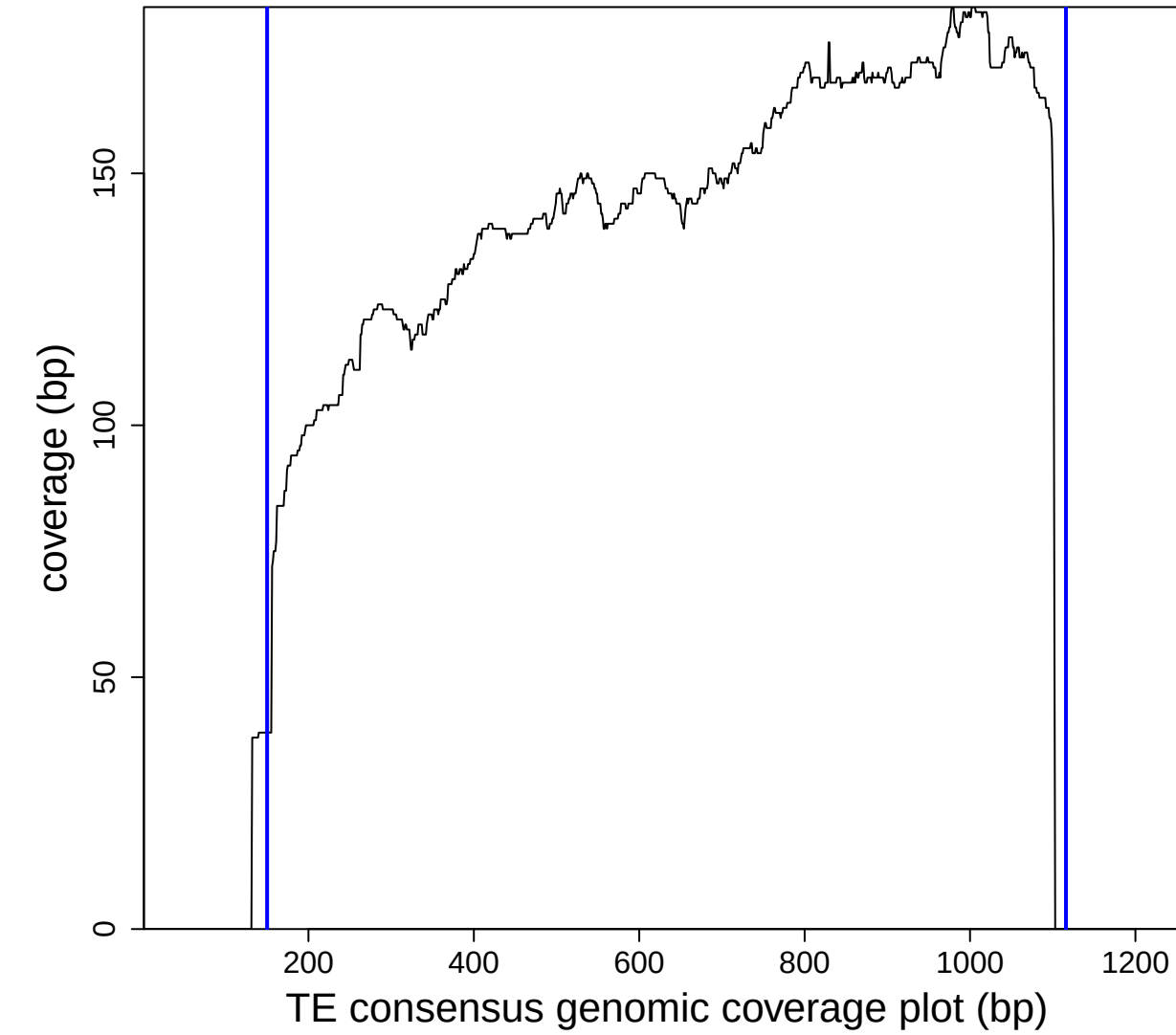
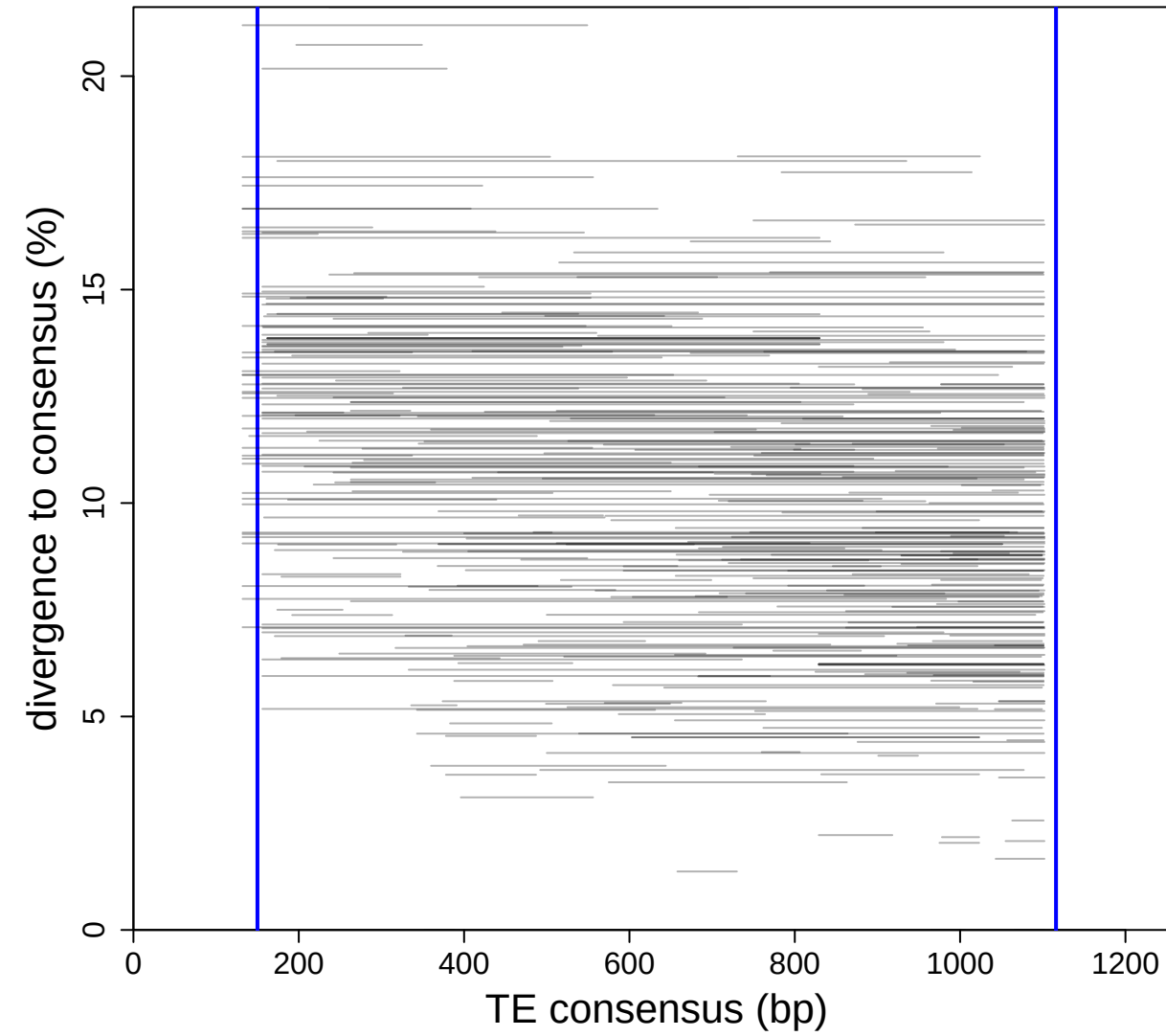
1400





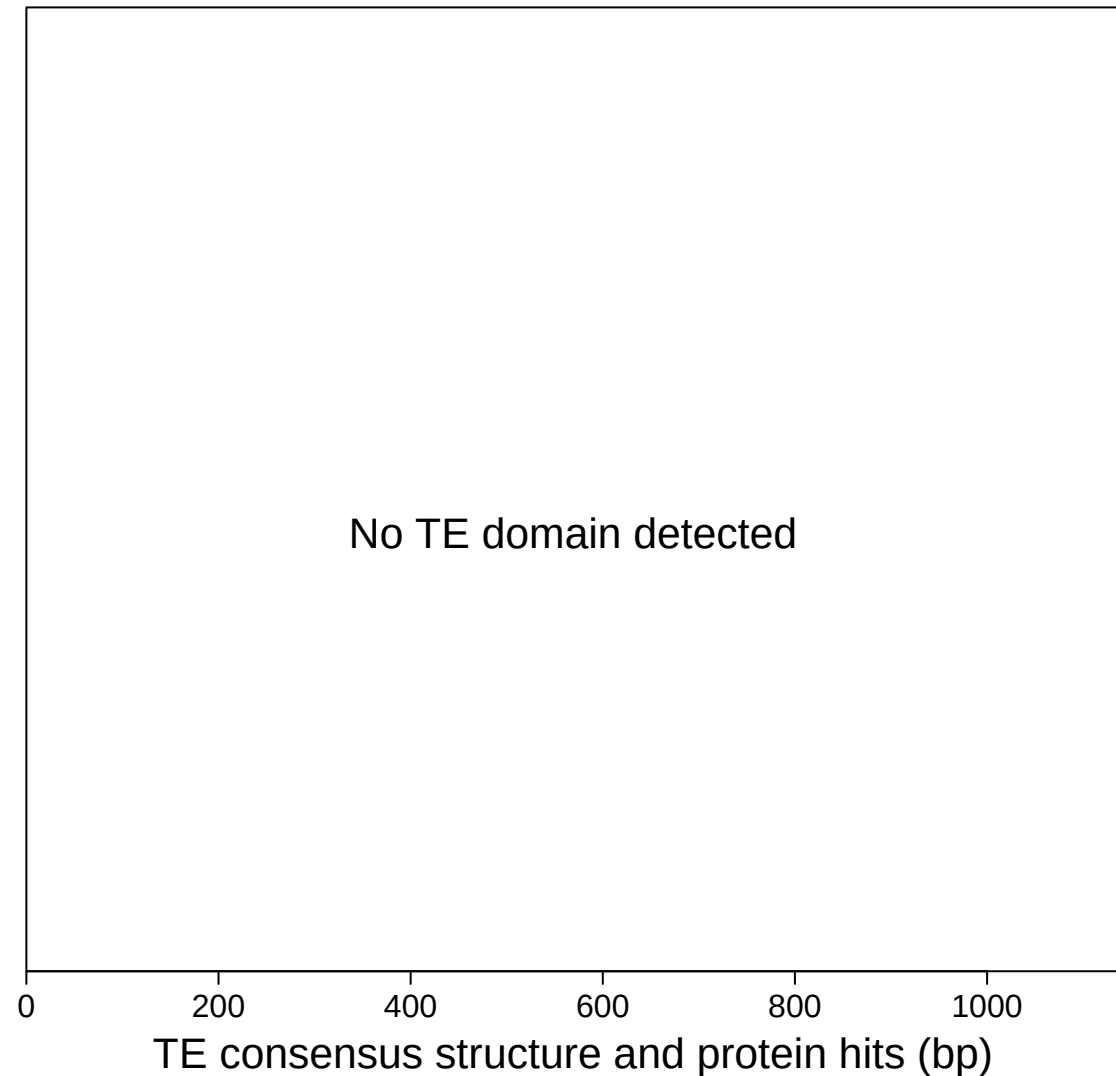
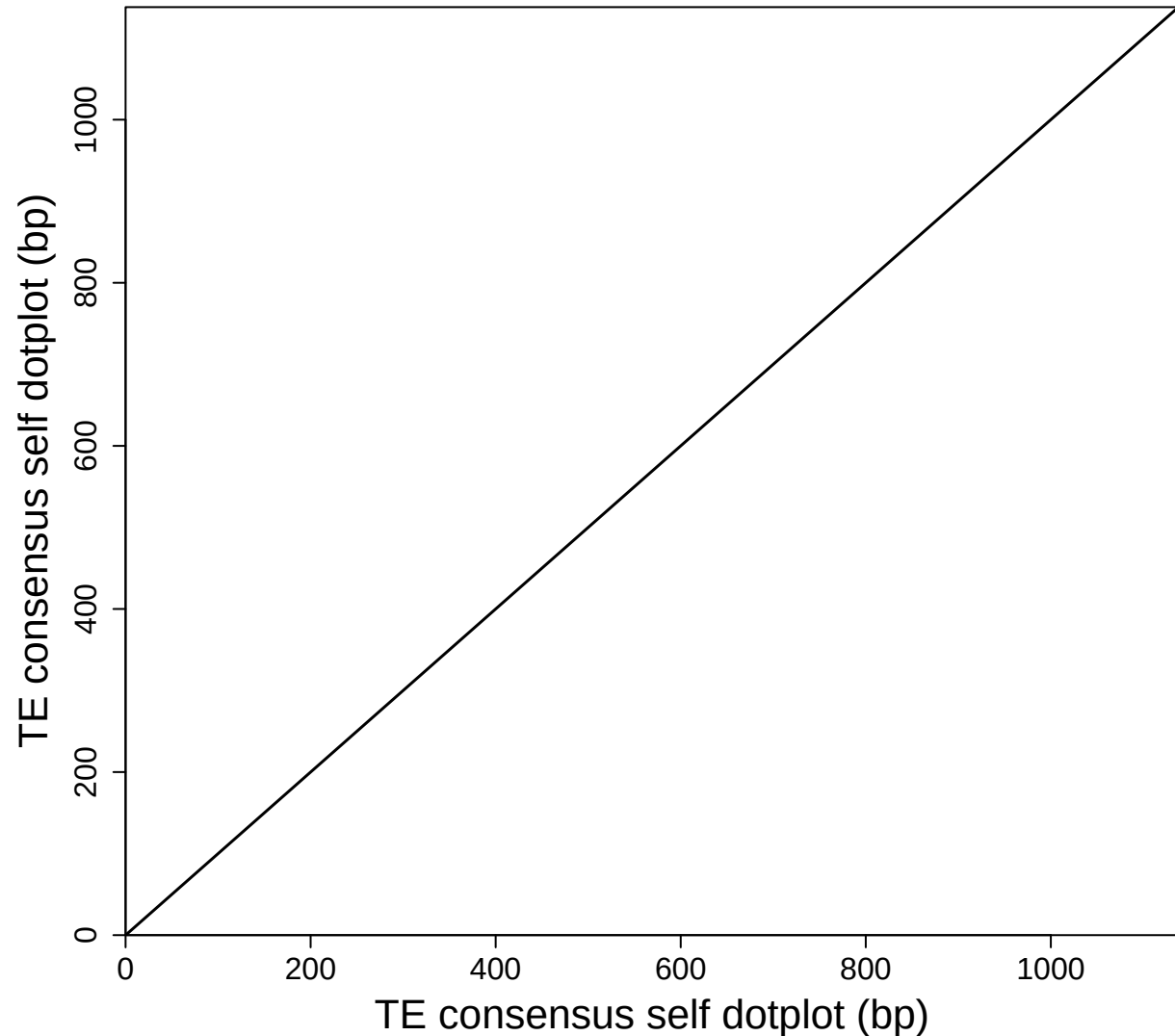
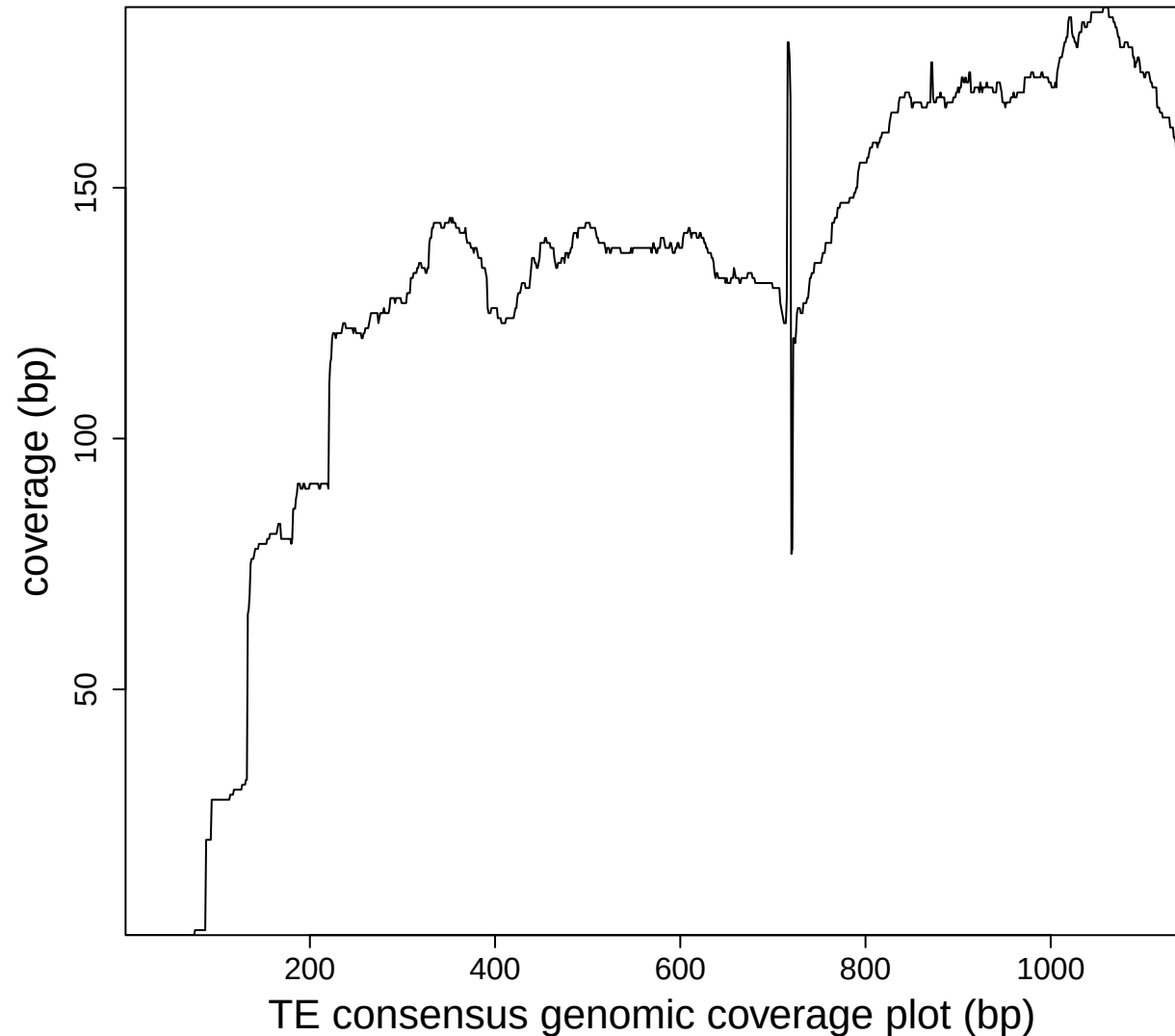
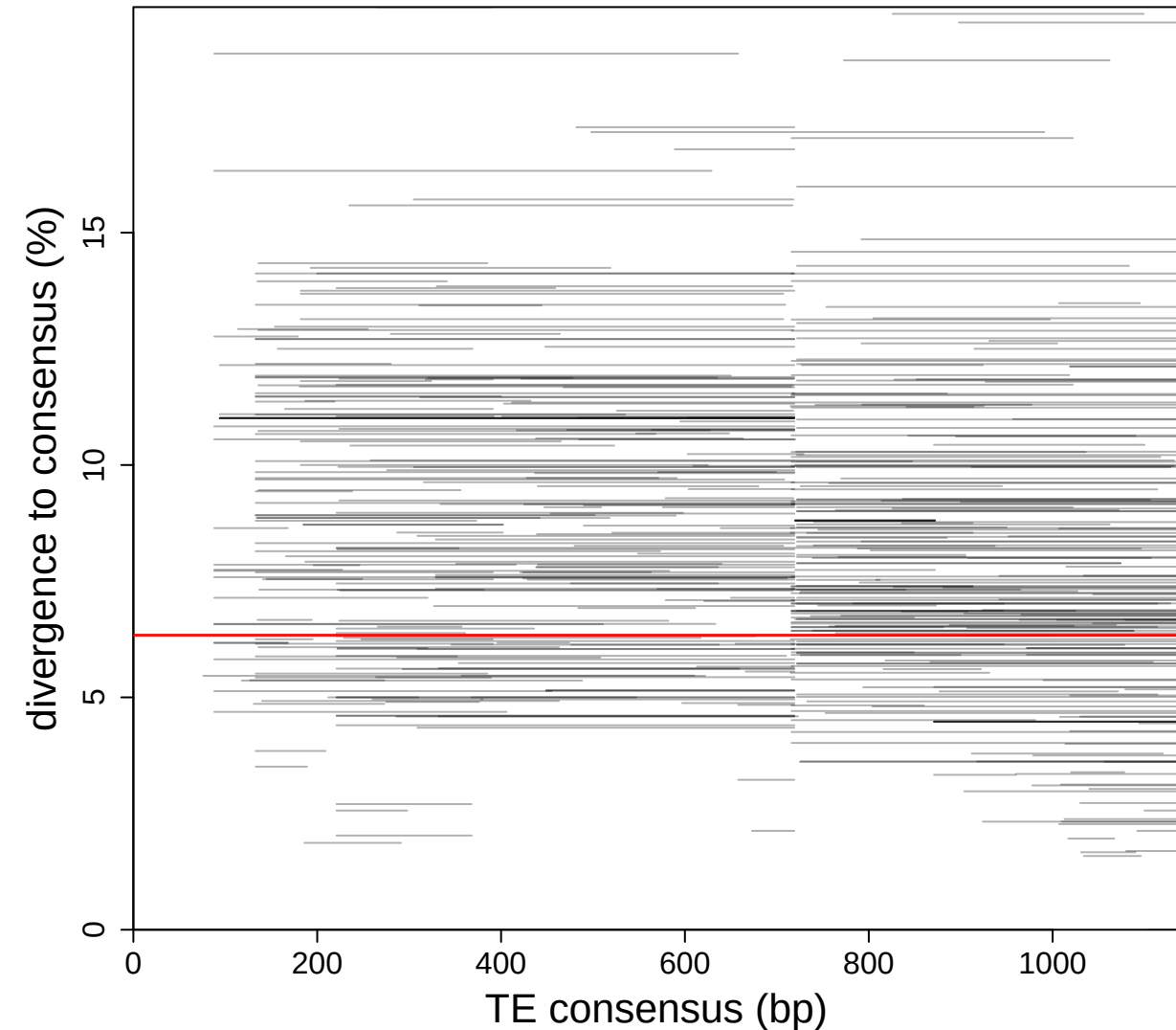
5.fasta.b.bed uf.bed g_2.bed fm_1.bed 0_0_bcln.fa aln.fa cl.f
size: 1266bp; fragments: 341; full length: 0 (>=1139.4bp)

After TEtrimmer Extended plot Blue lines are boundaries

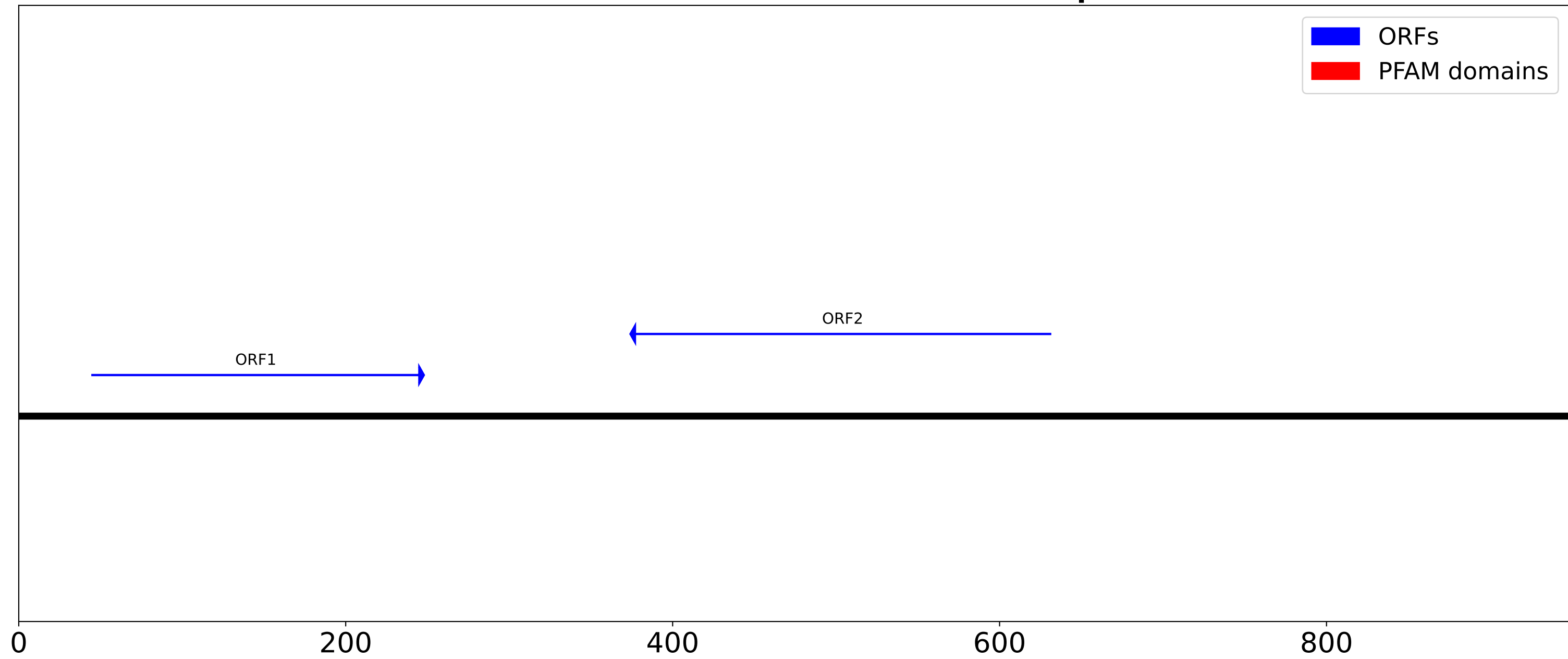


TE: rnd_5_family_3335
size: 1138bp; fragments: 473; full length: 1 (>=1024.2bp)

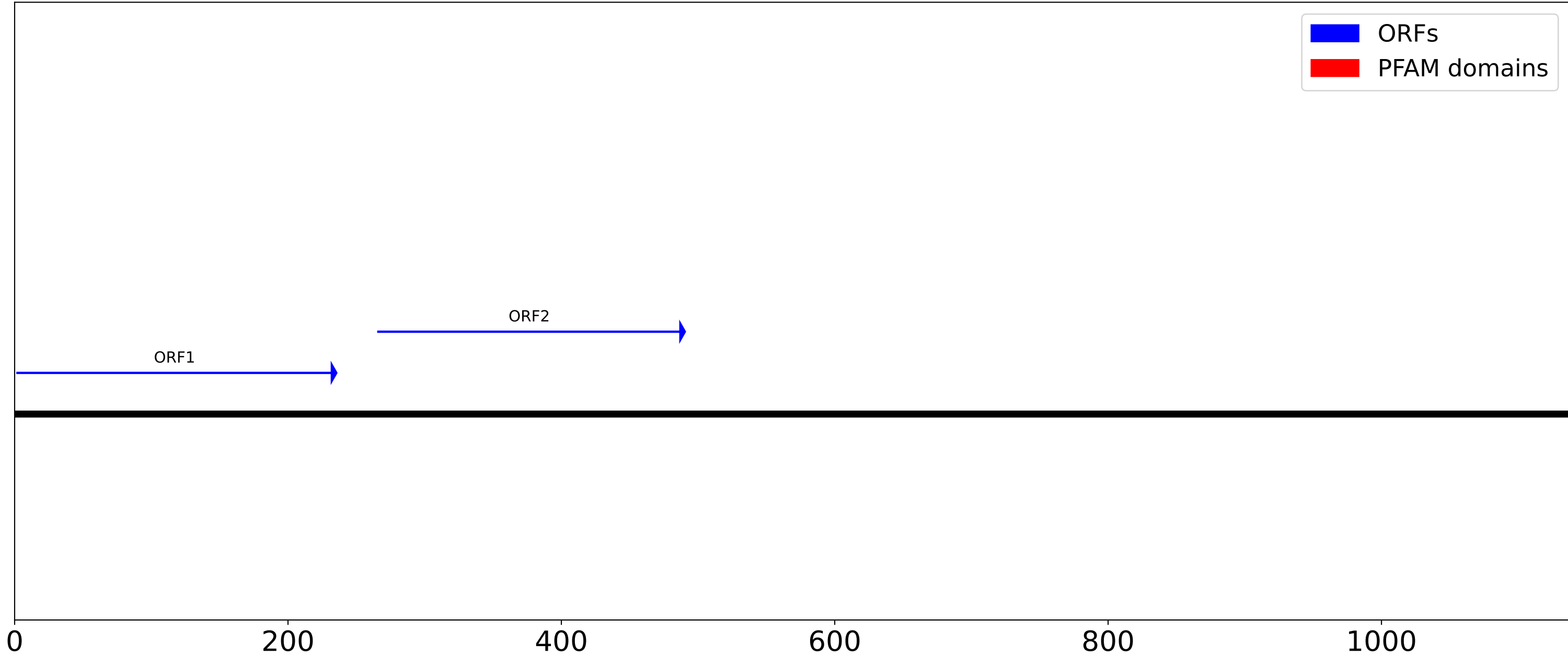
Before TEtrimmer 1138 bp



After TEtrimmer ORF and PFAM domain plot



Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

