

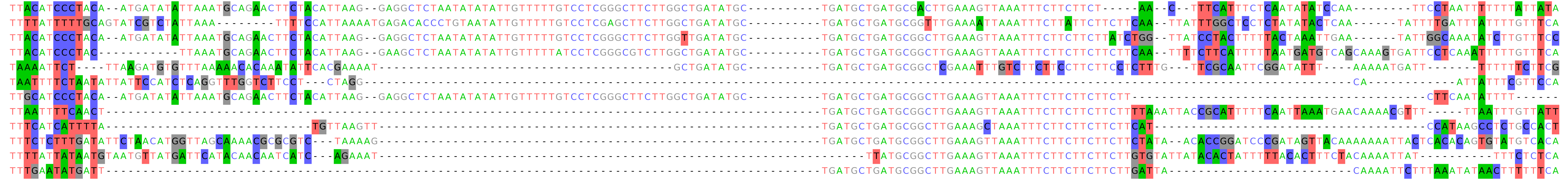
Start crop Point



End crop Point



MSA length = 951



Start crop Point

End crop Point

0 -

10 -

0

200

400

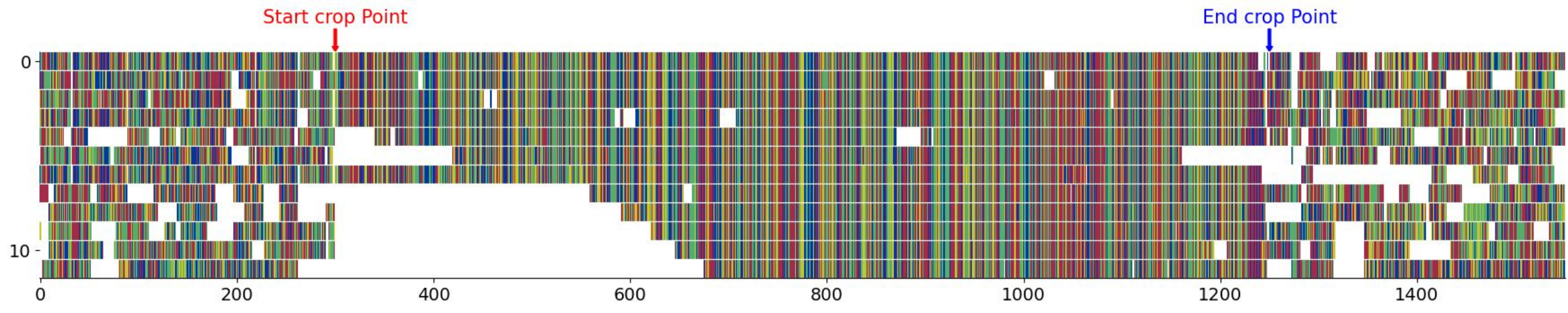
600

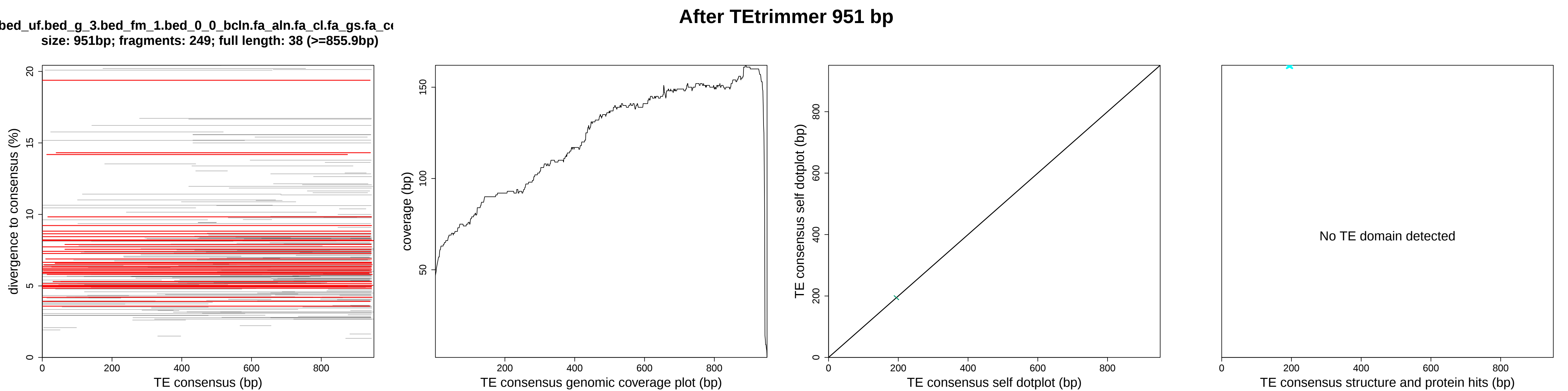
800

1000

1200

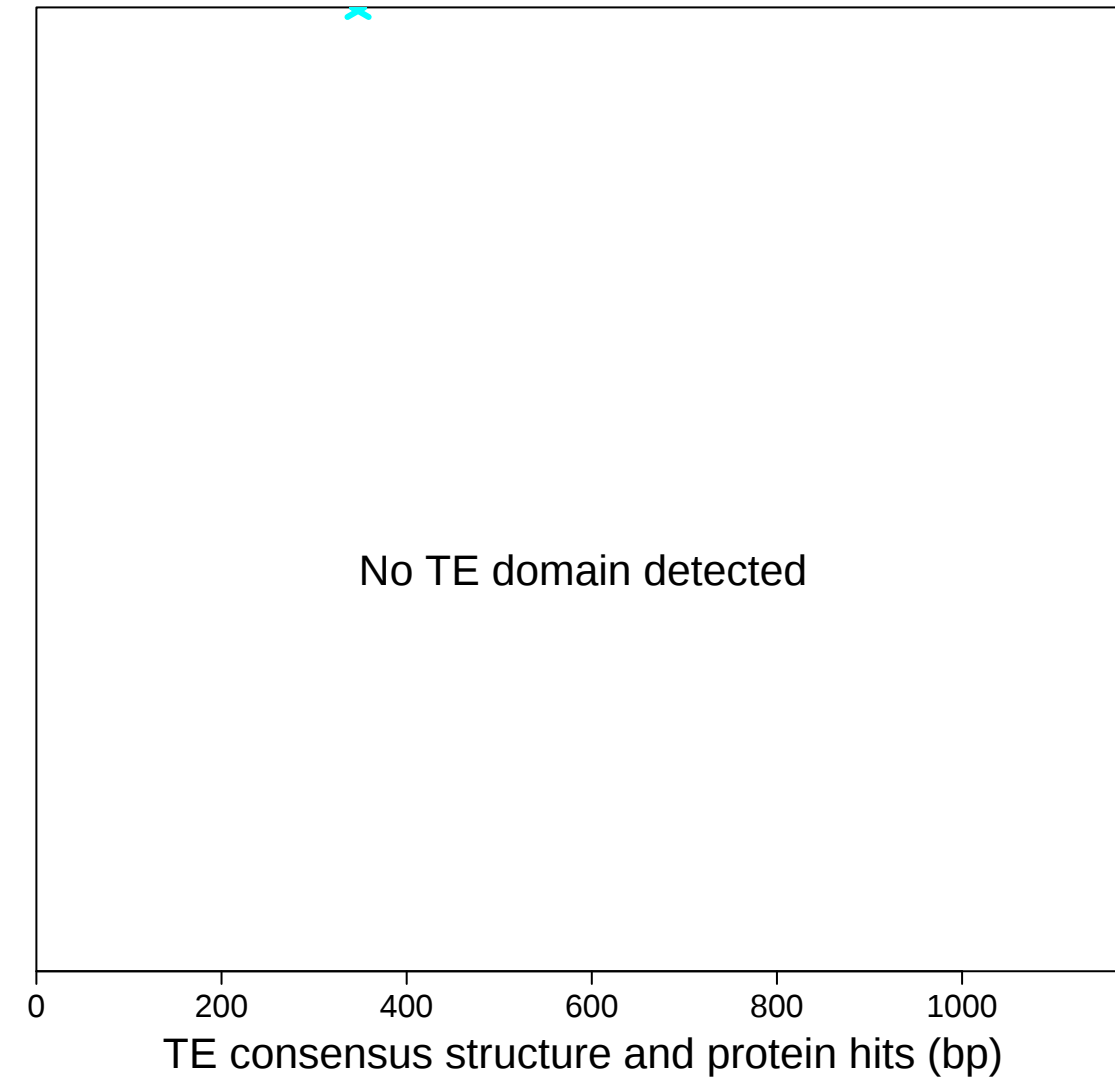
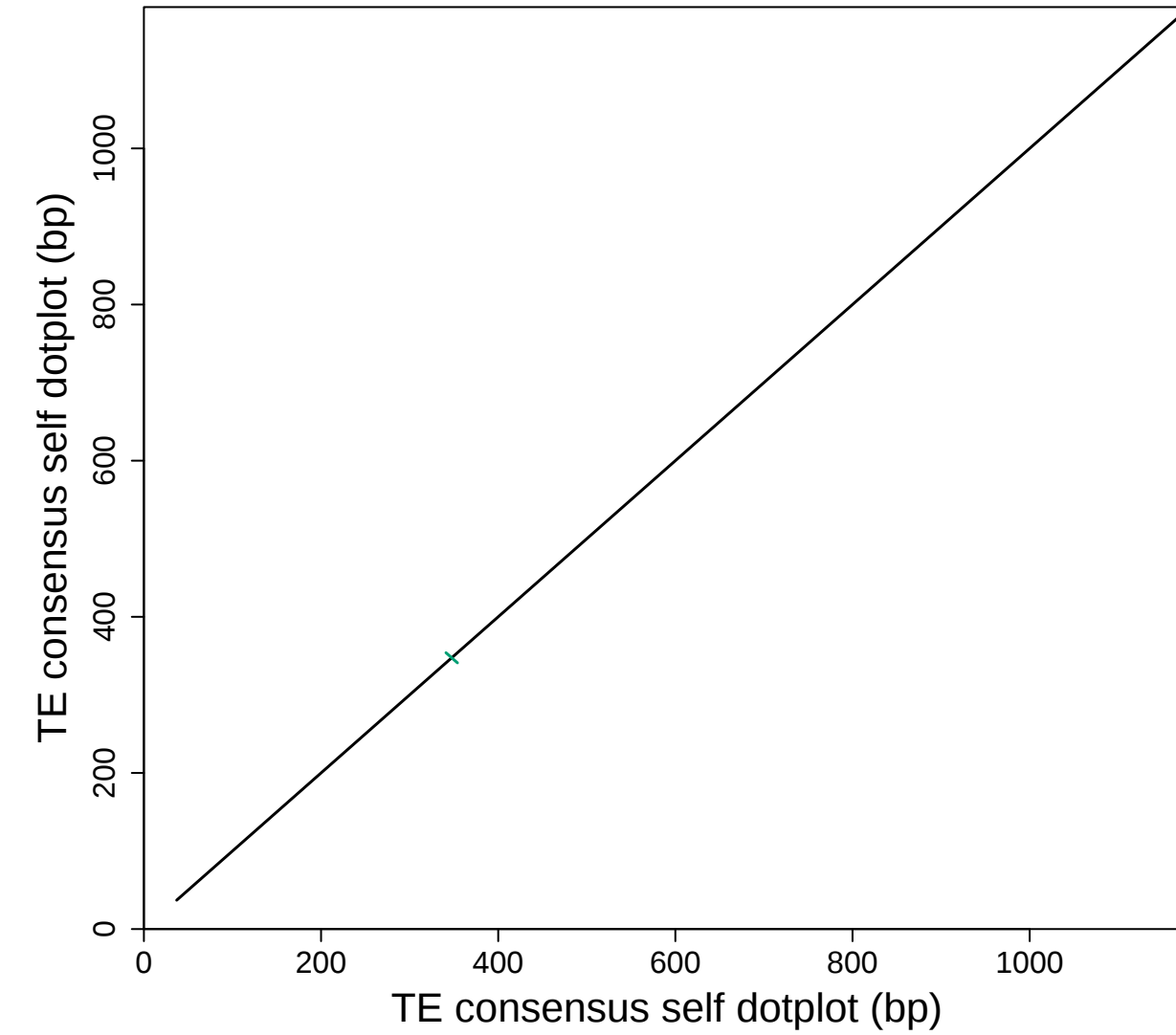
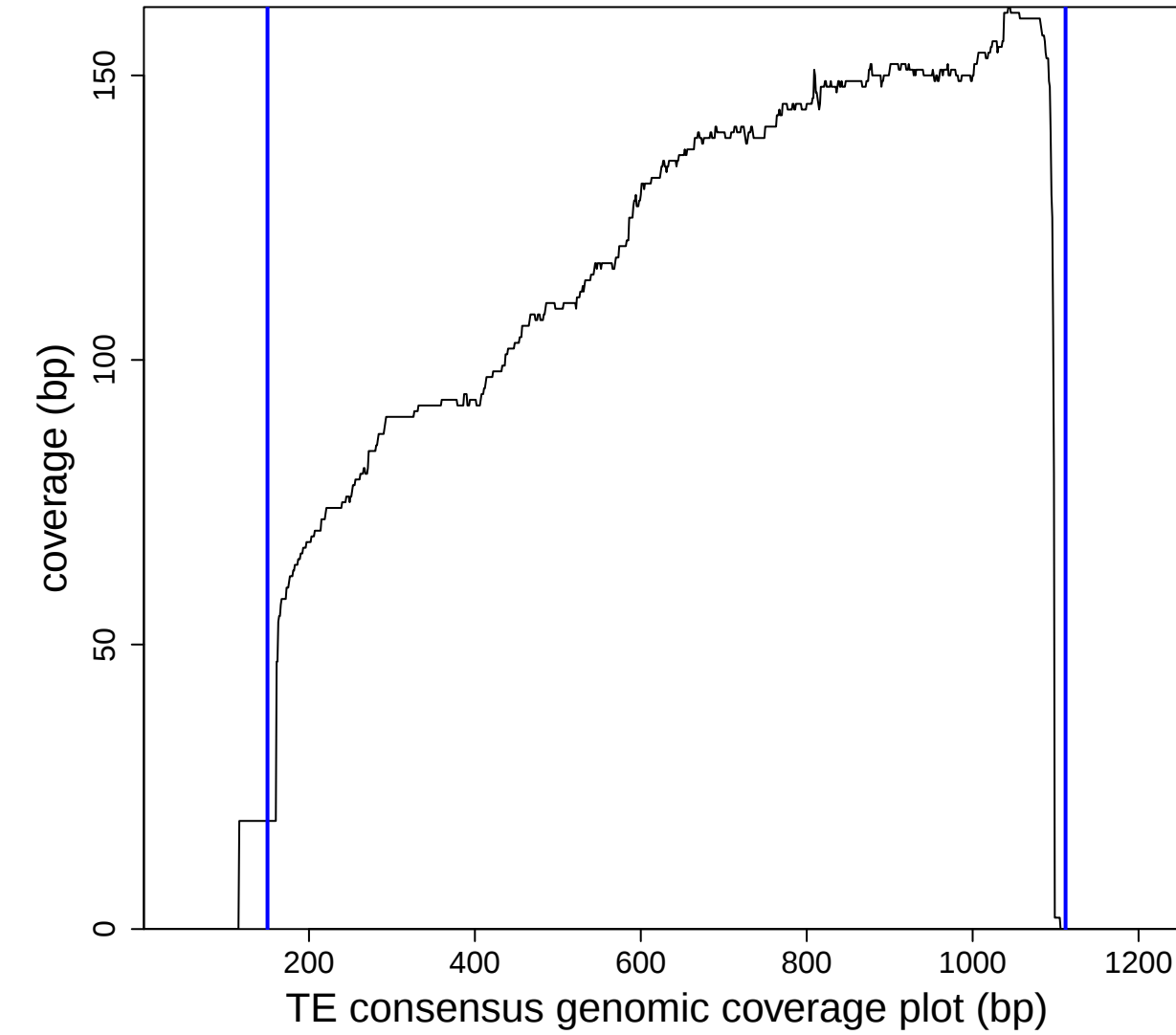
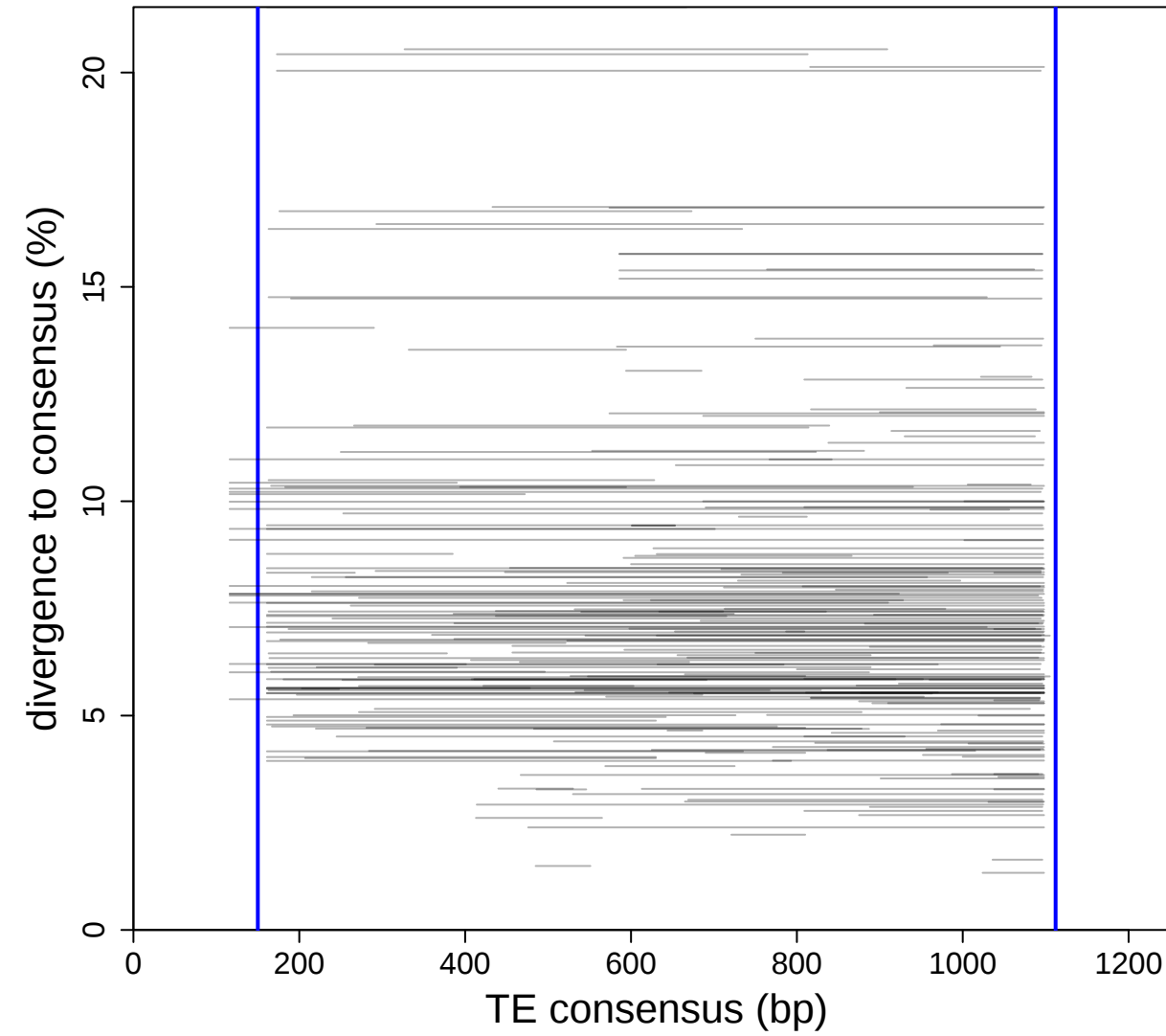
1400



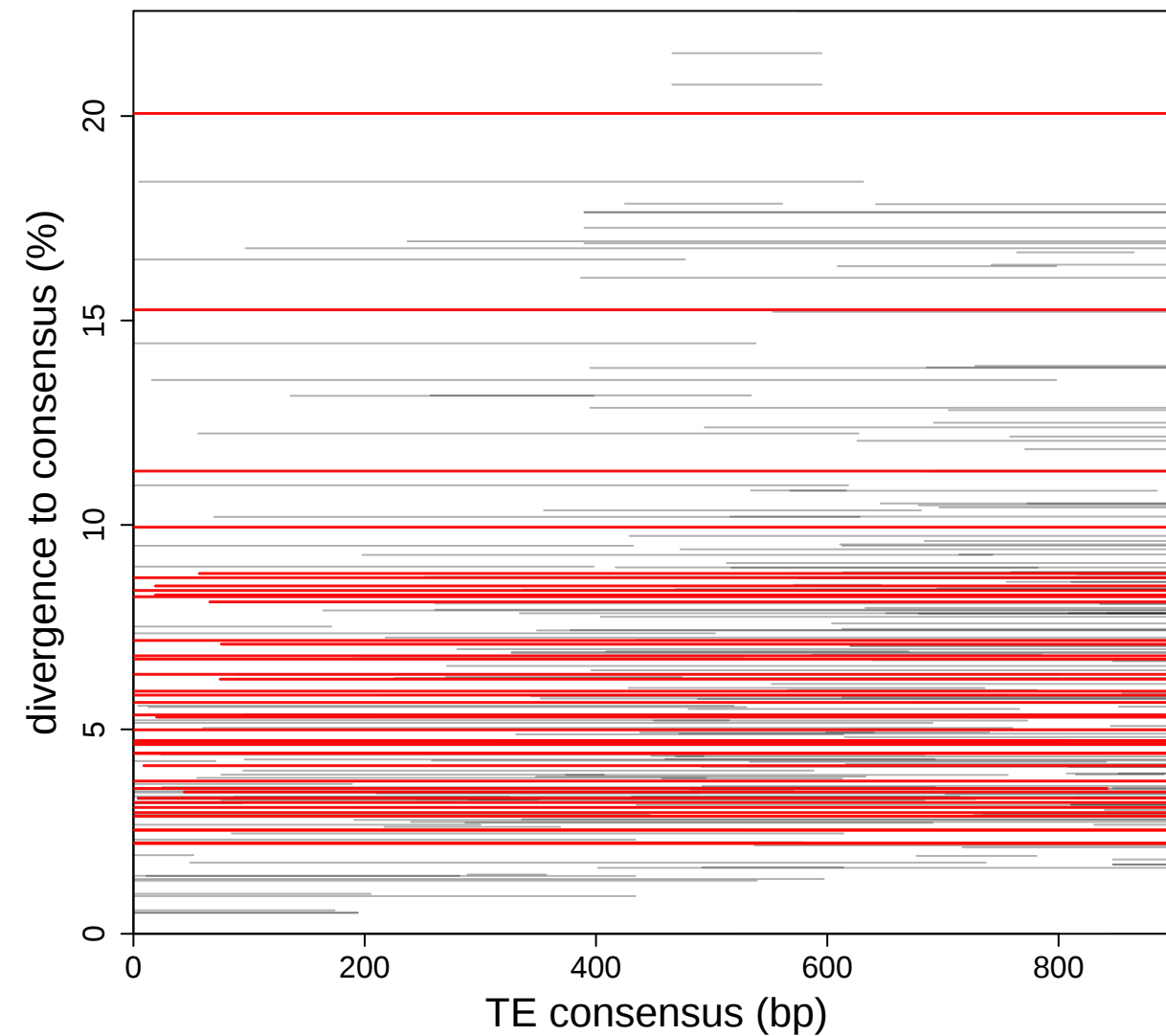


4.fasta.b.bed_uf.bed_g_3.bed_fm_1.bed_0_0_bcIn.fa_aln.fa_cl.f
size: 1262bp; fragments: 247; full length: 0 (>=1135.8bp)

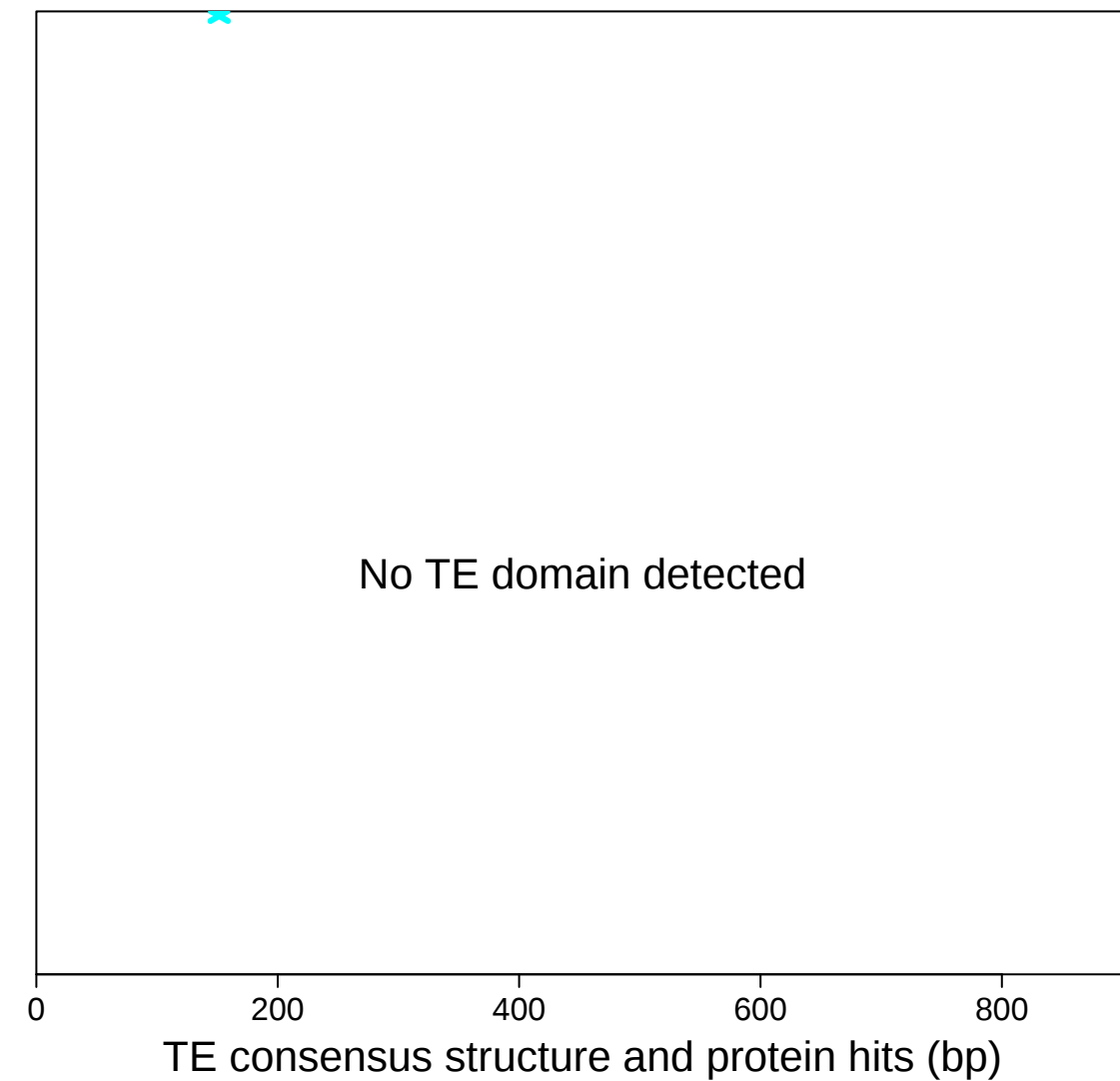
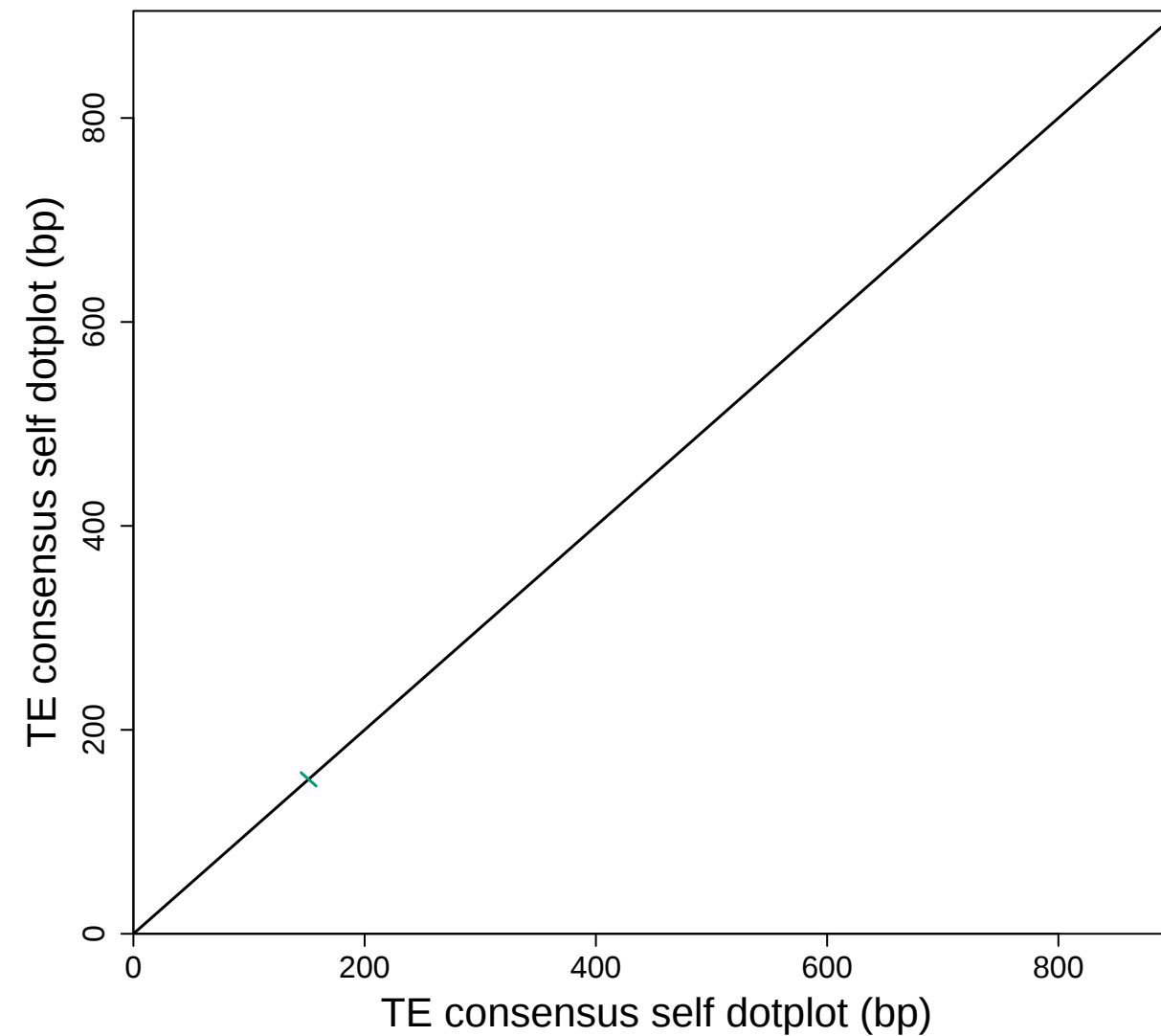
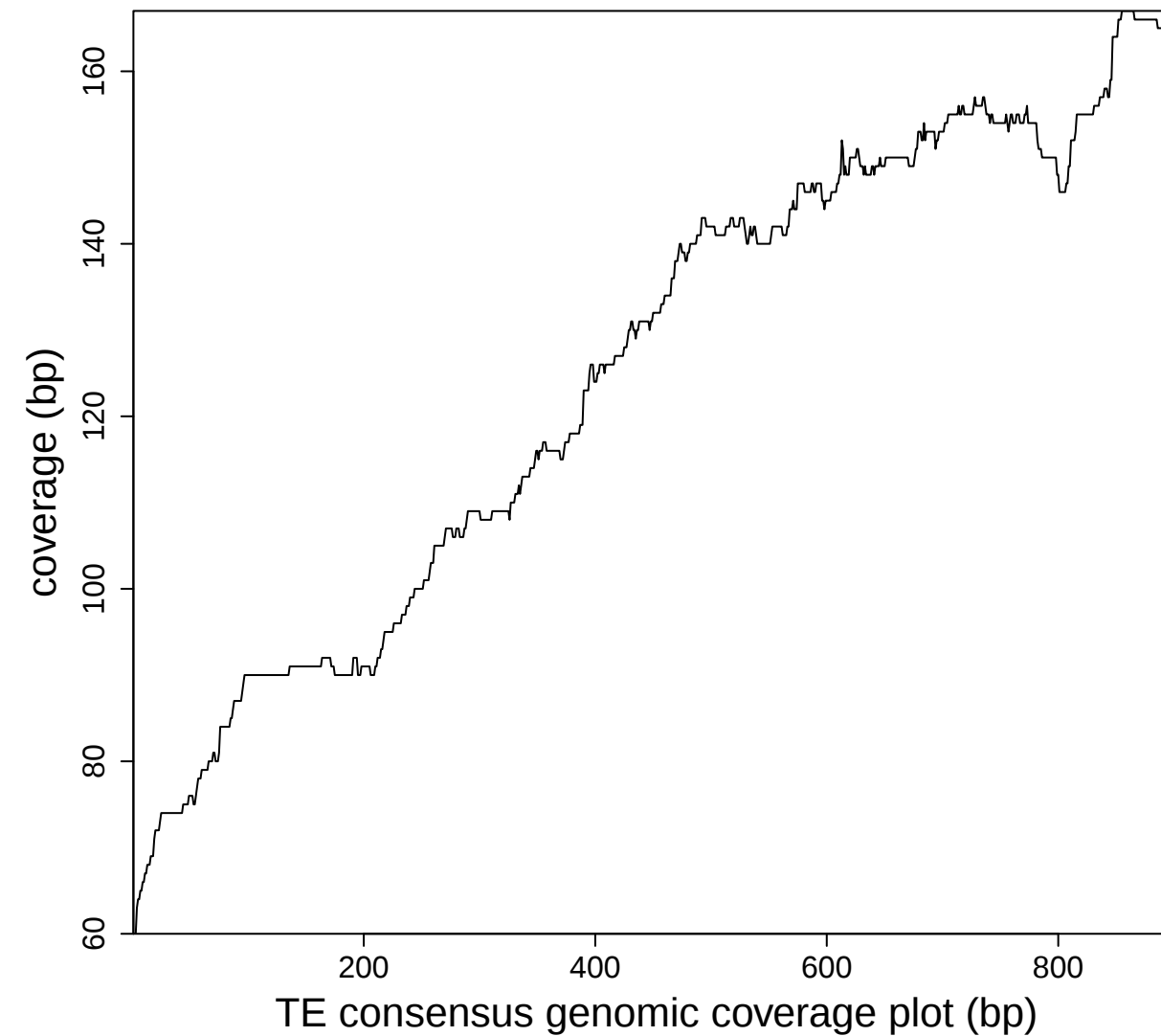
After TEtrimmer Extended plot Blue lines are boundaries



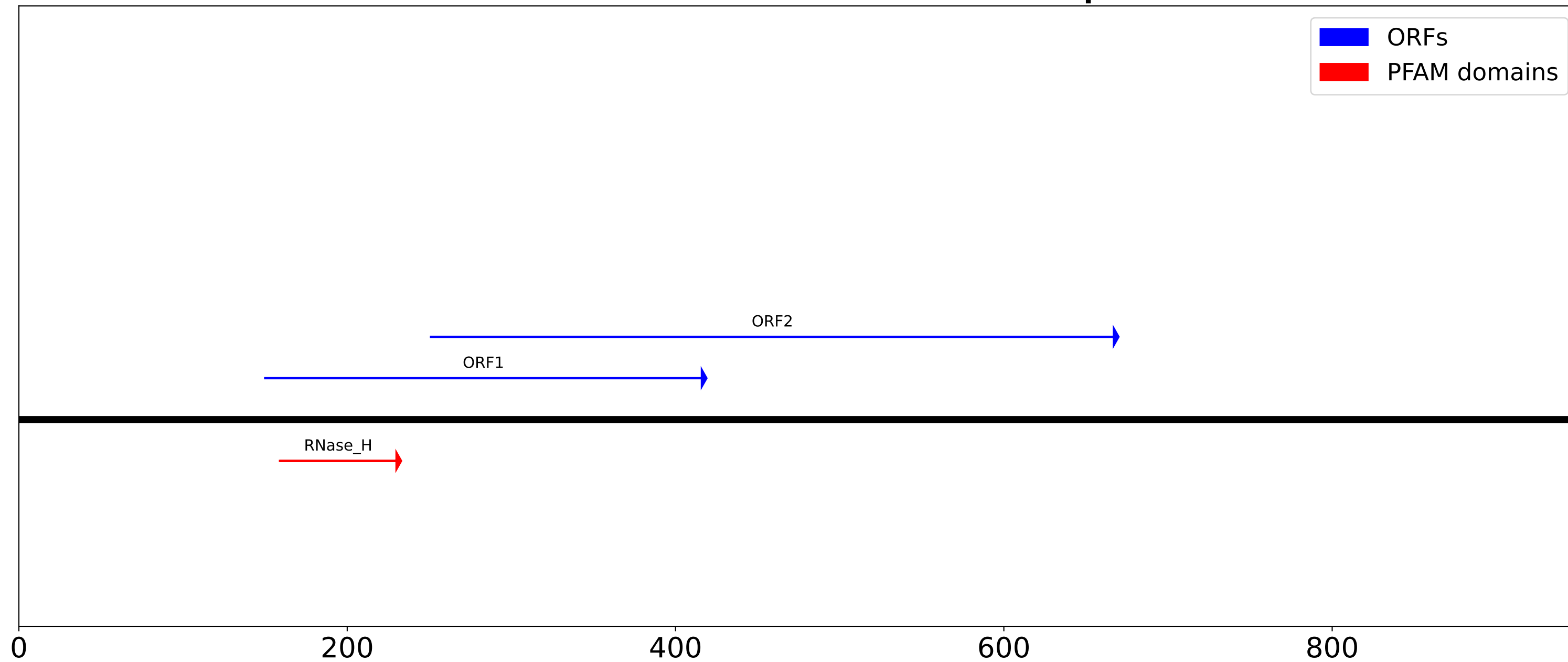
TE: rnd_5_family_3414
size: 905bp; fragments: 257; full length: 42 (≥ 814.5 bp)



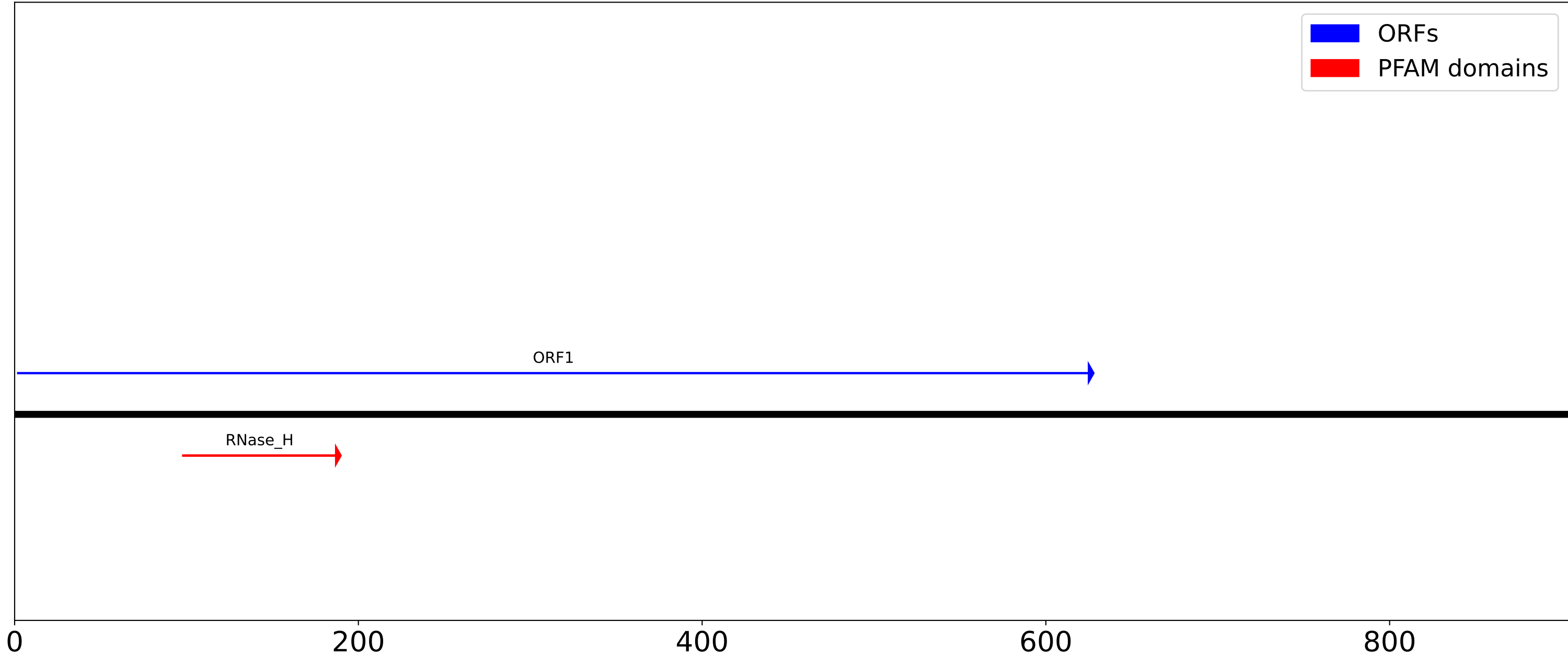
Before TEtrimmer 905 bp



After TEtrimmer ORF and PFAM domain plot



Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

