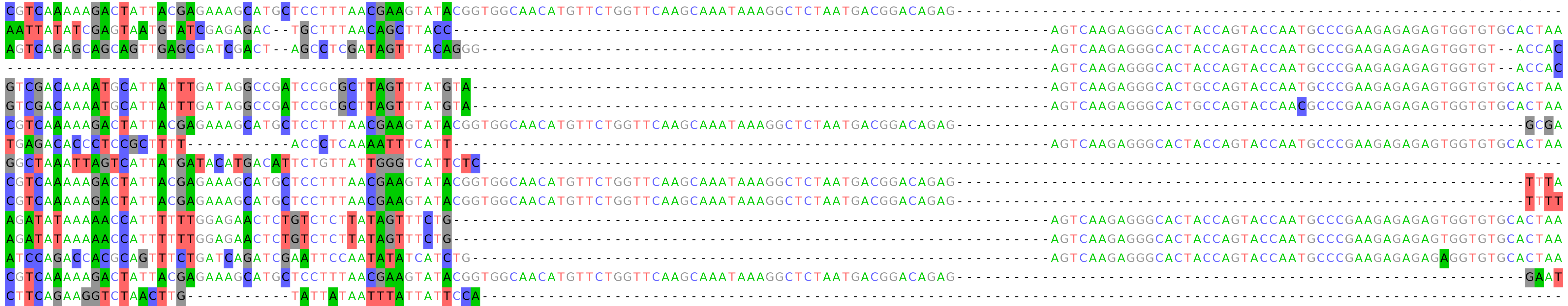


Start crop Point

End crop Point

MSA length = 1714



Start crop Point

End crop Point

0

10

0

250

500

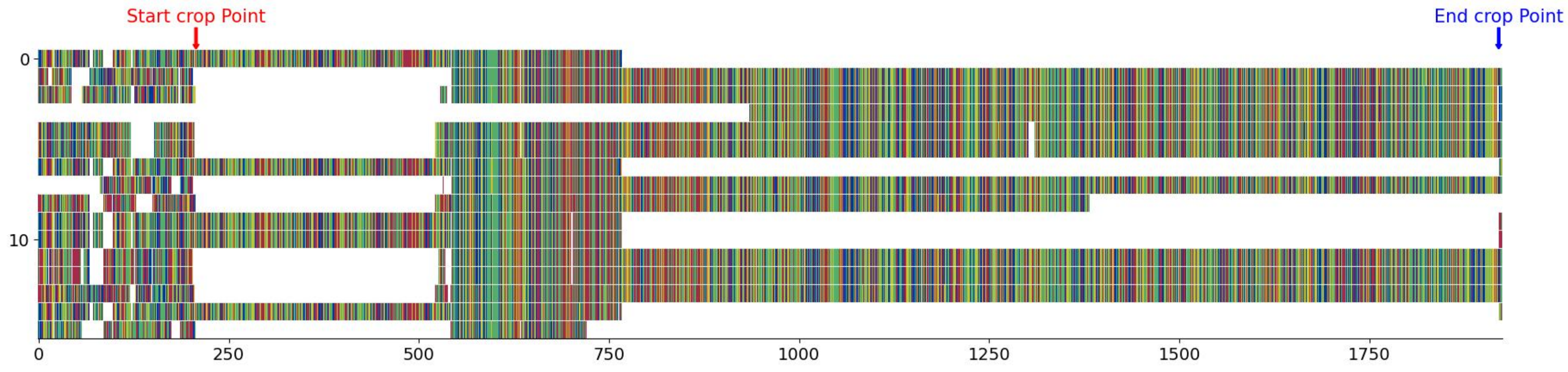
750

1000

1250

1500

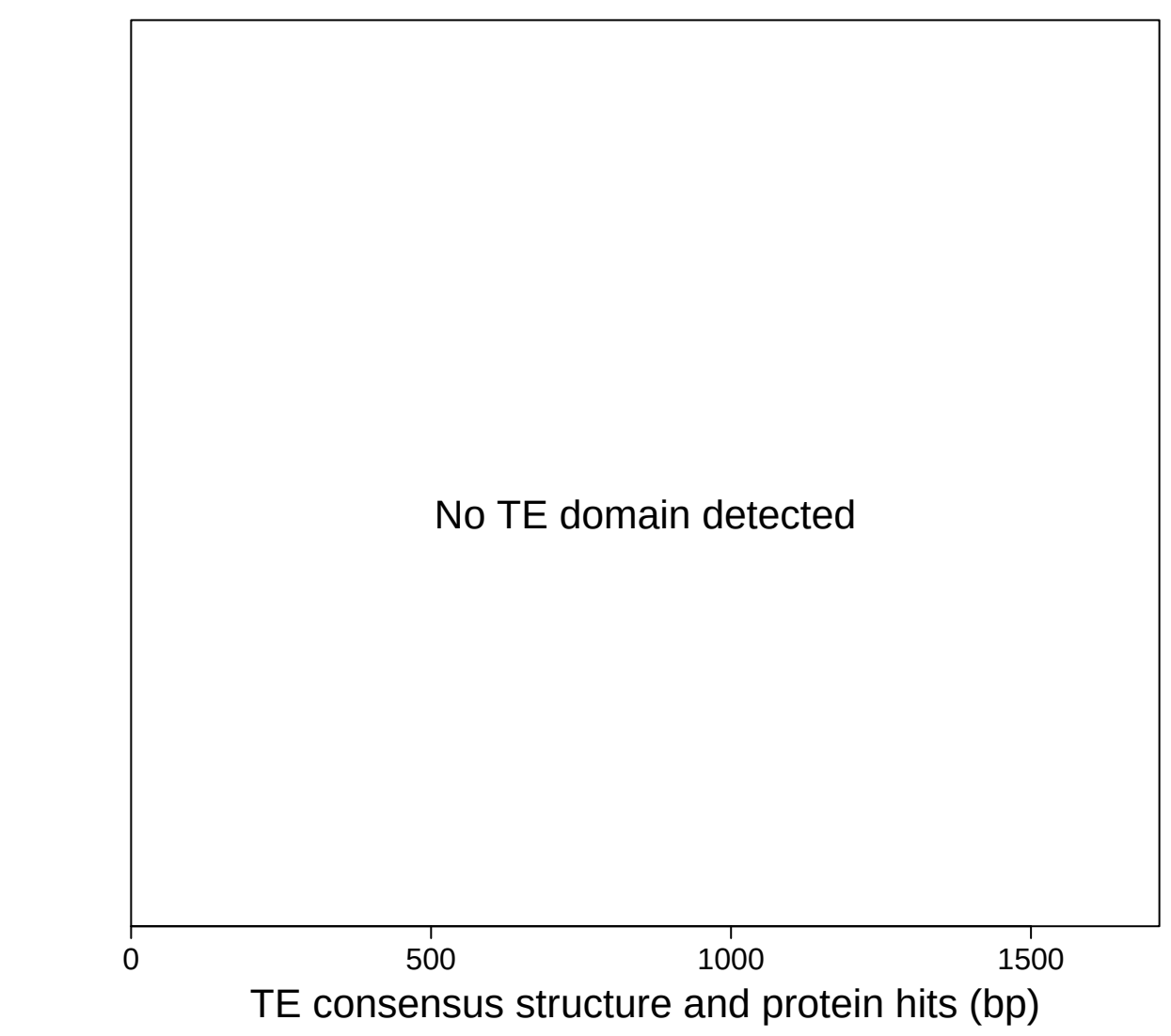
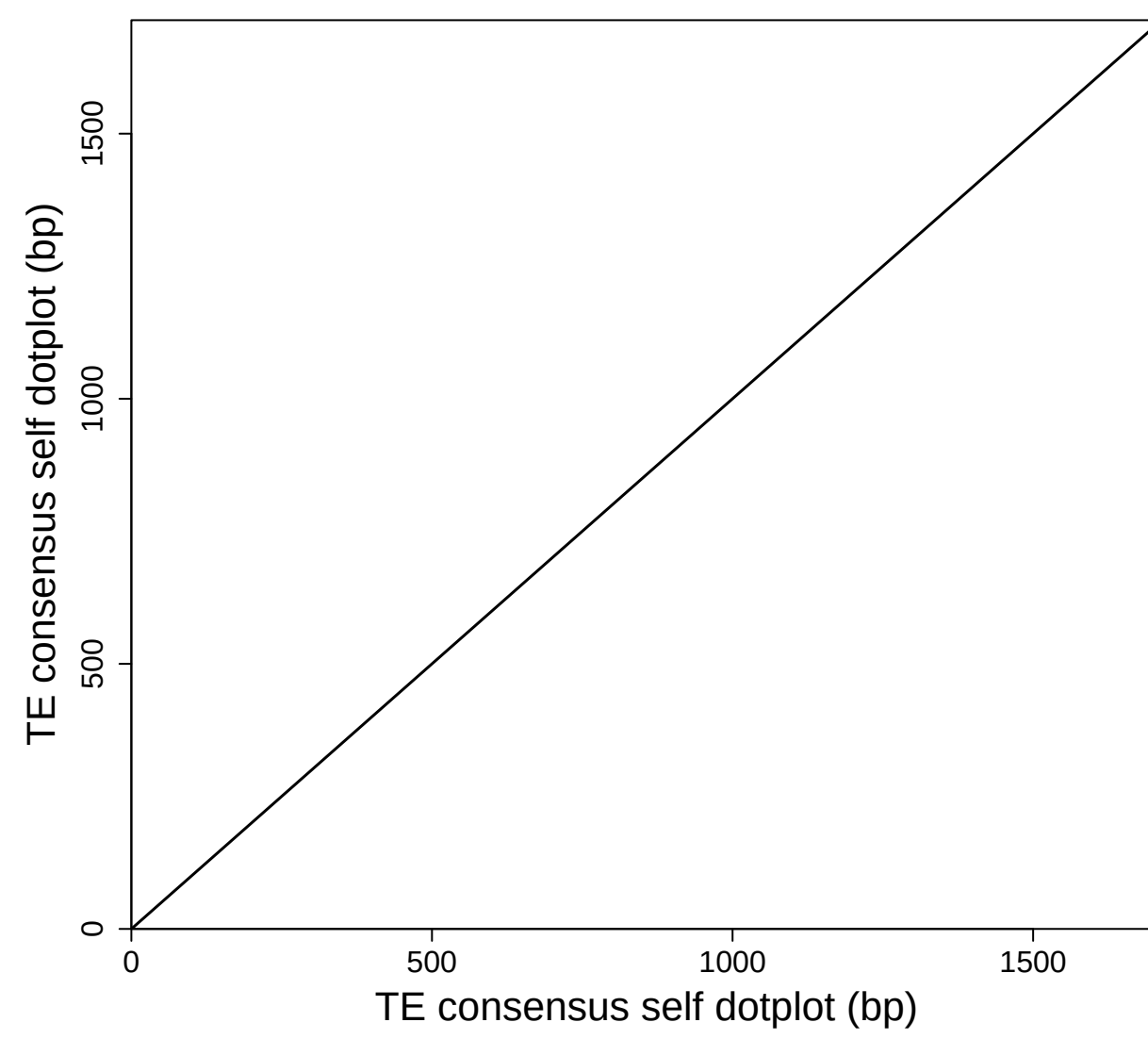
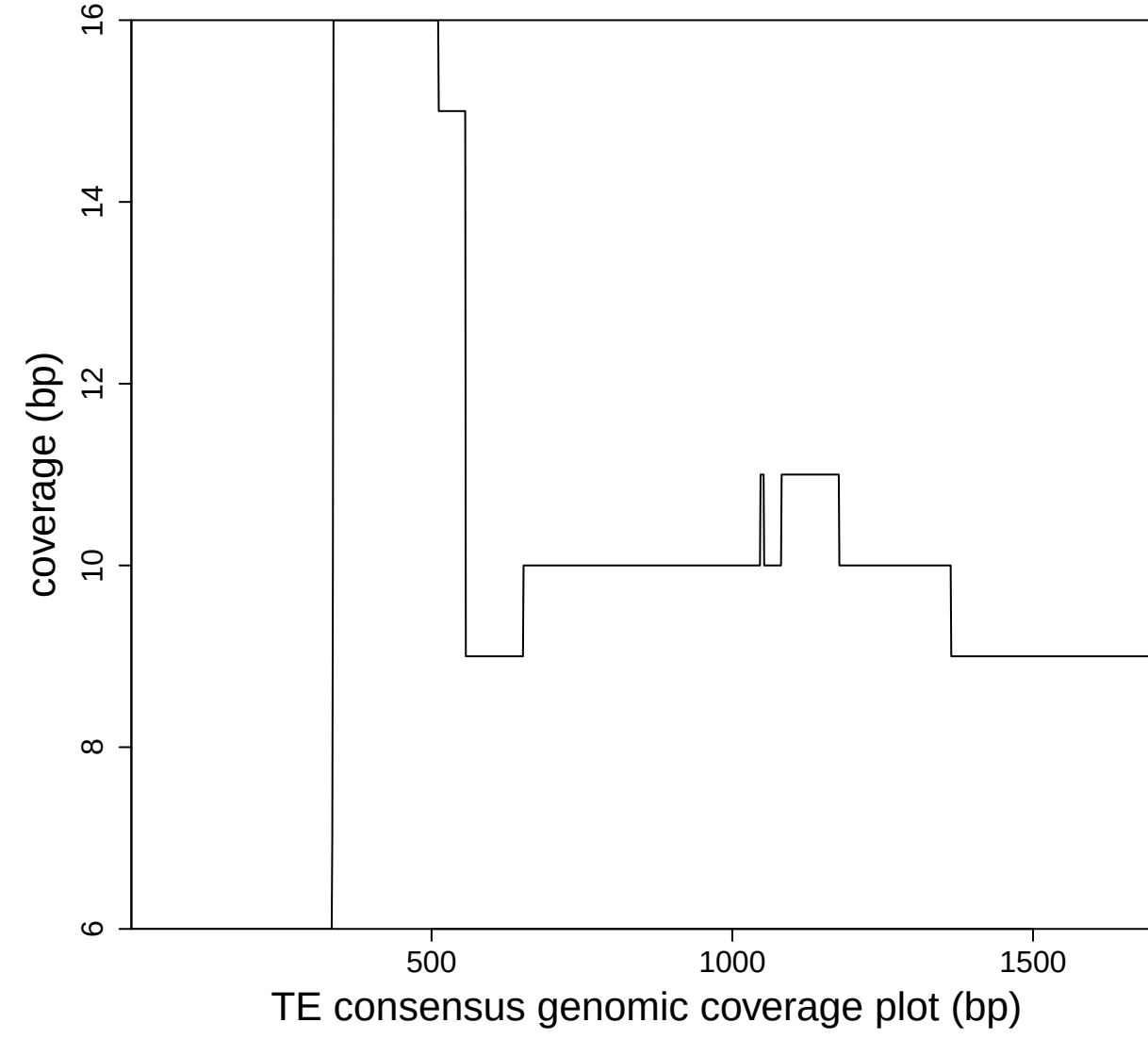
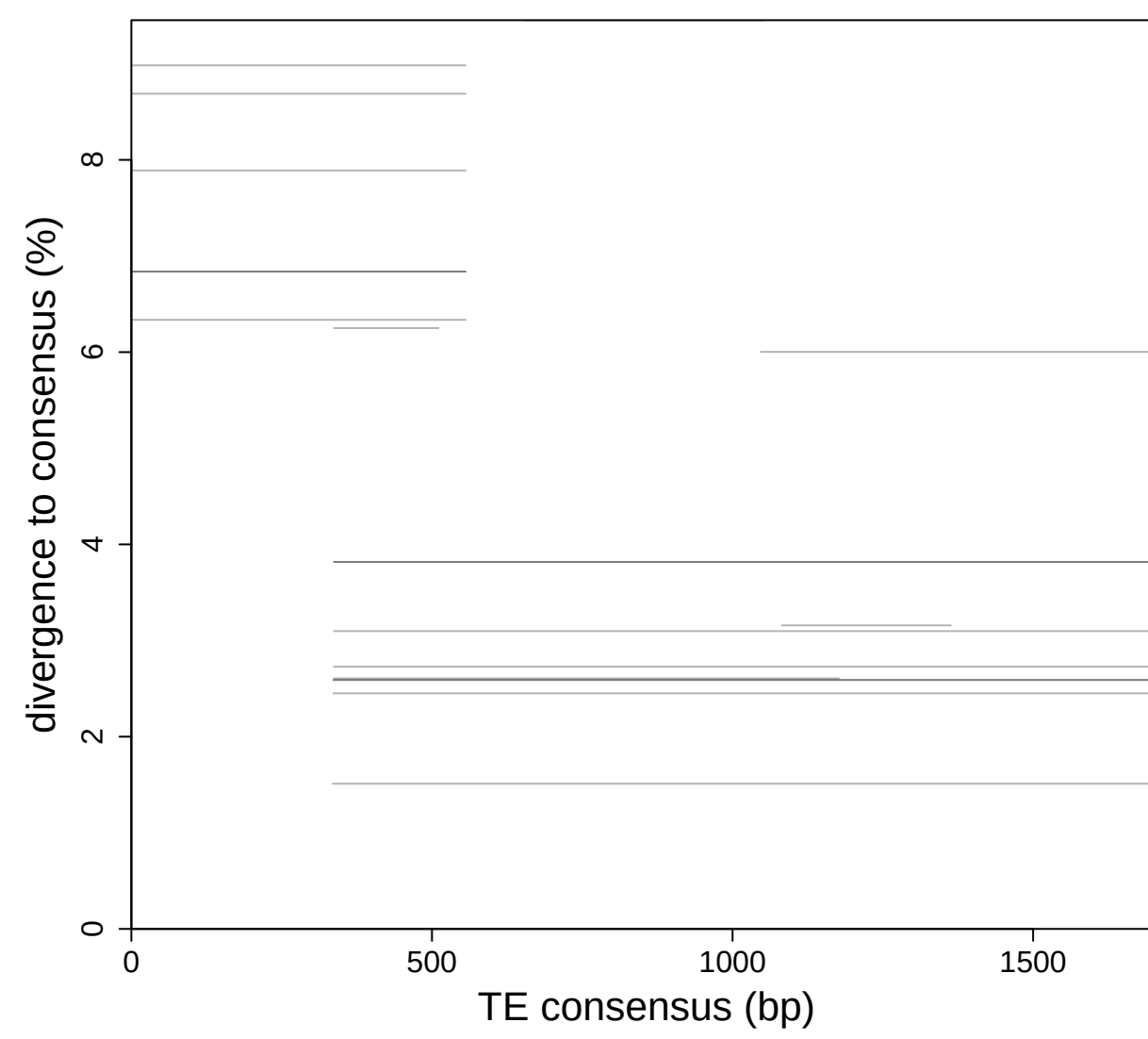
1750



b.bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.fa_

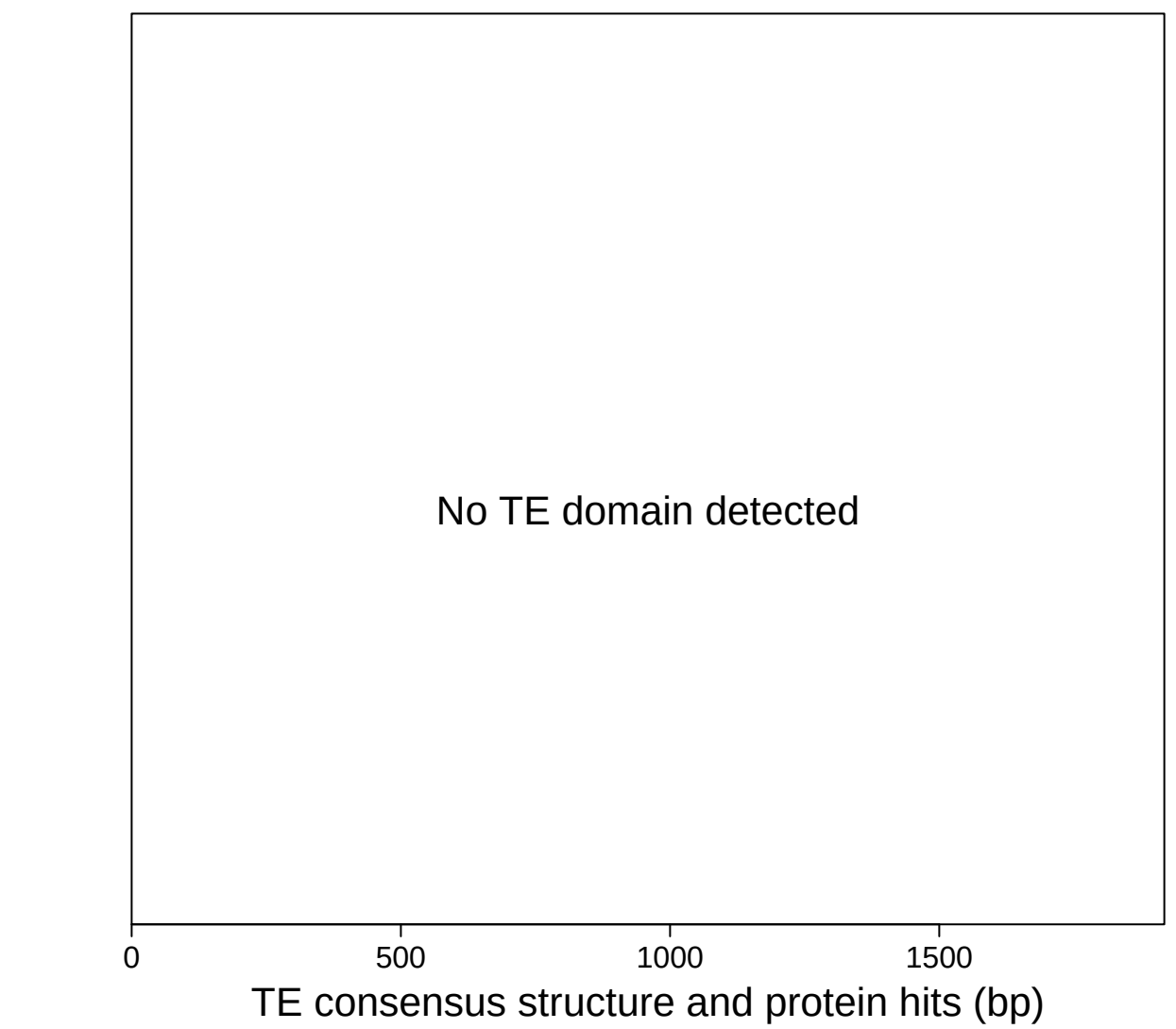
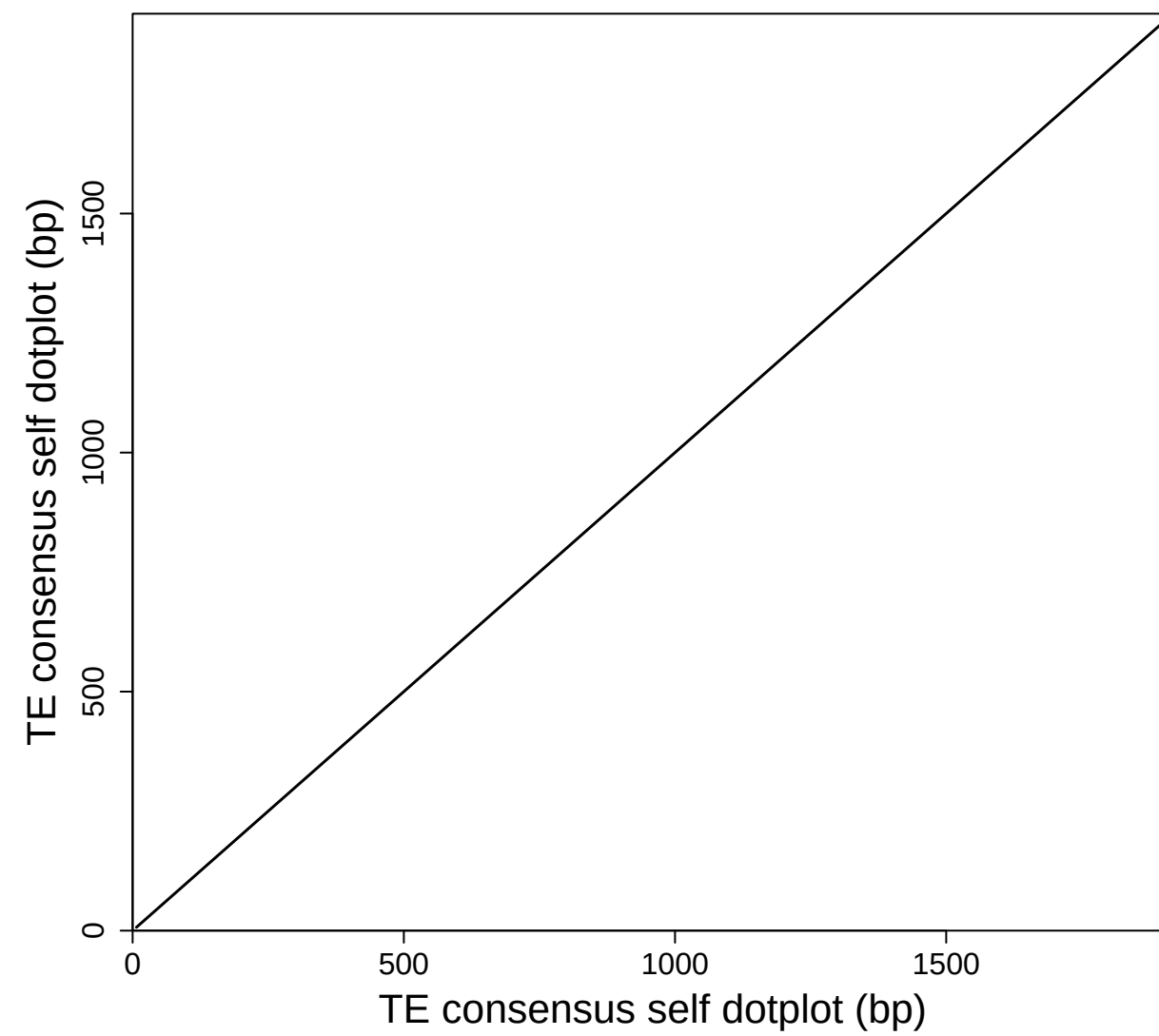
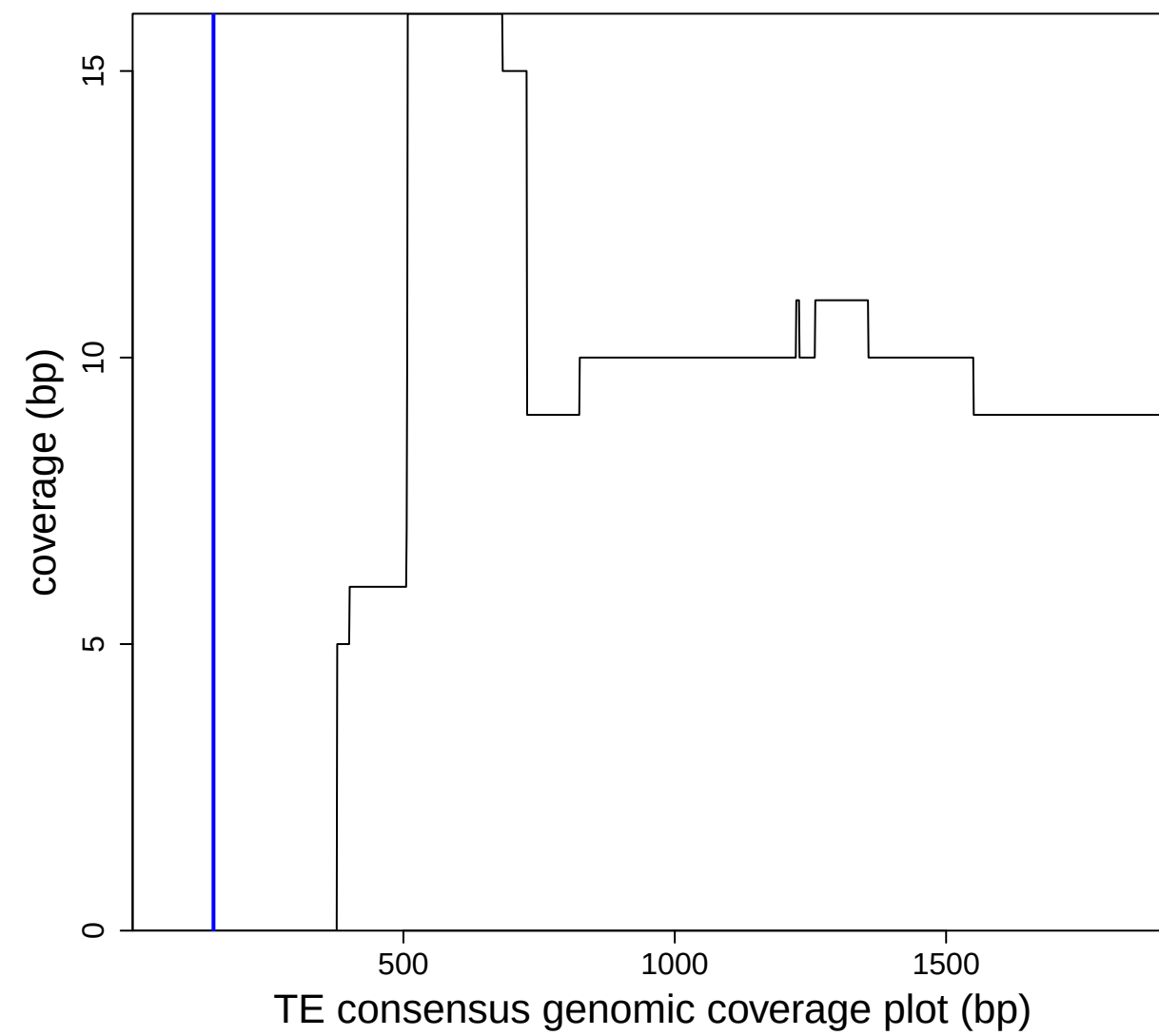
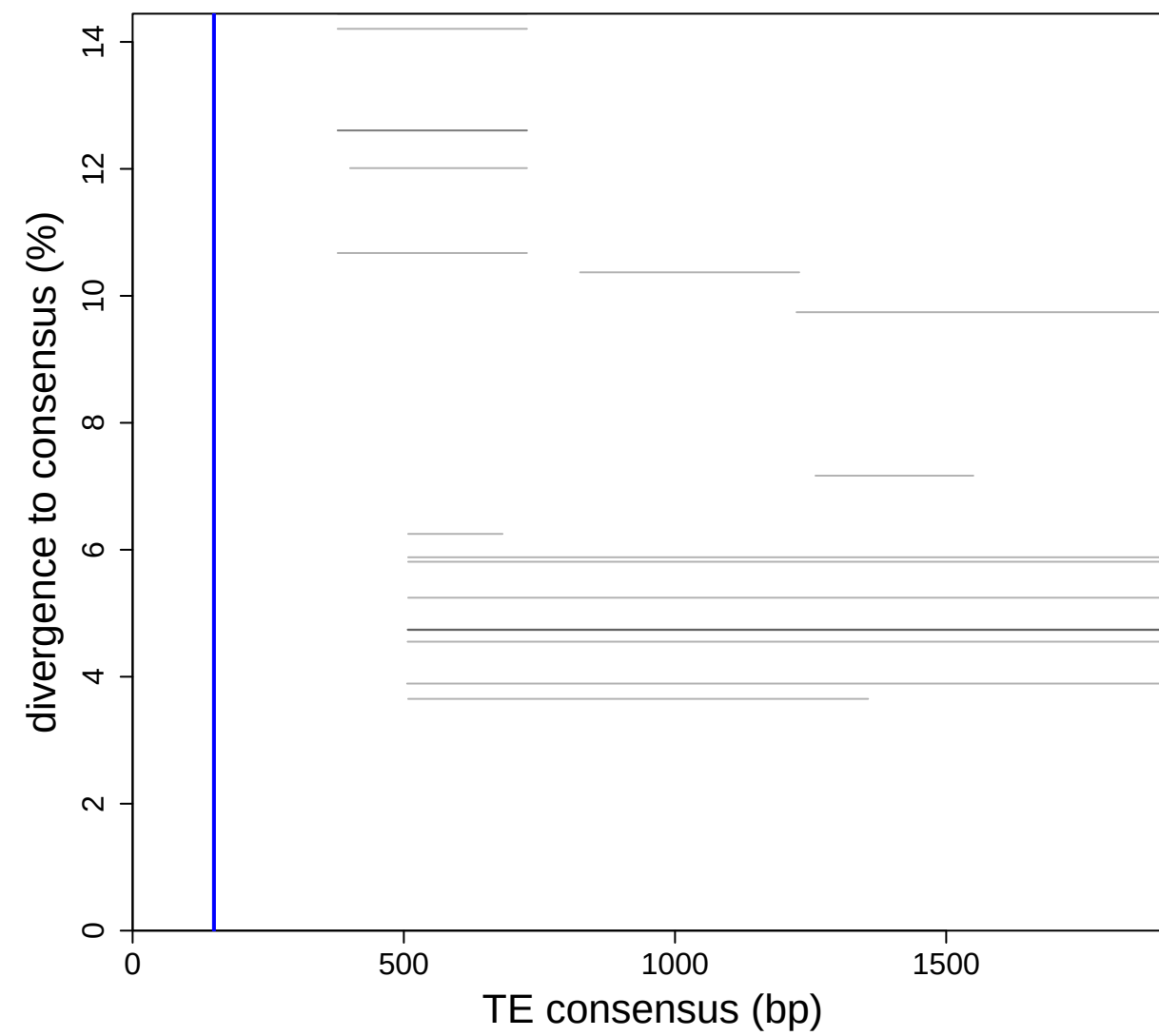
size: 1714bp; fragments: 19; full length: 0 (>=1542.6bp)

After TEtrimmer 1714 bp



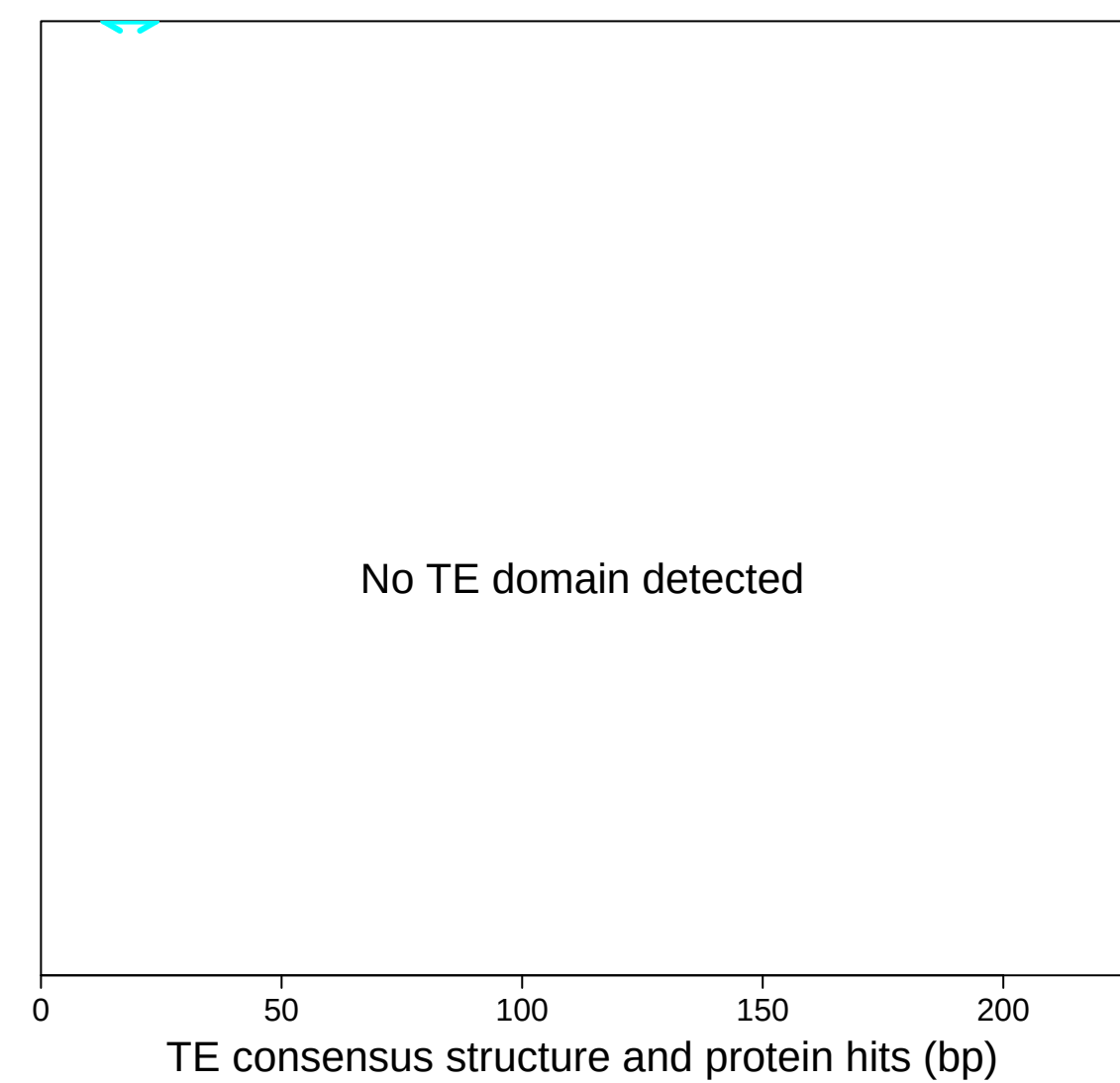
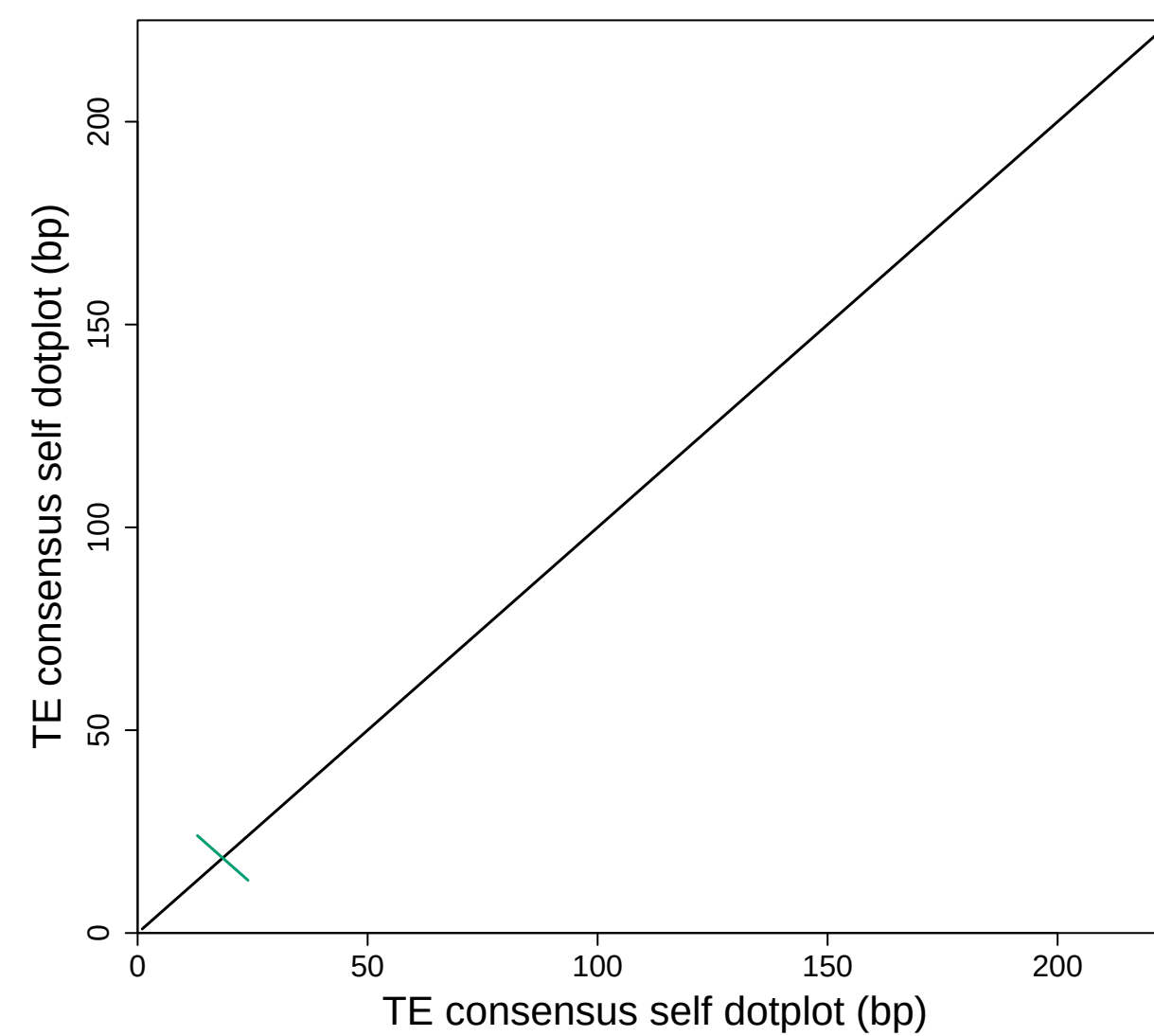
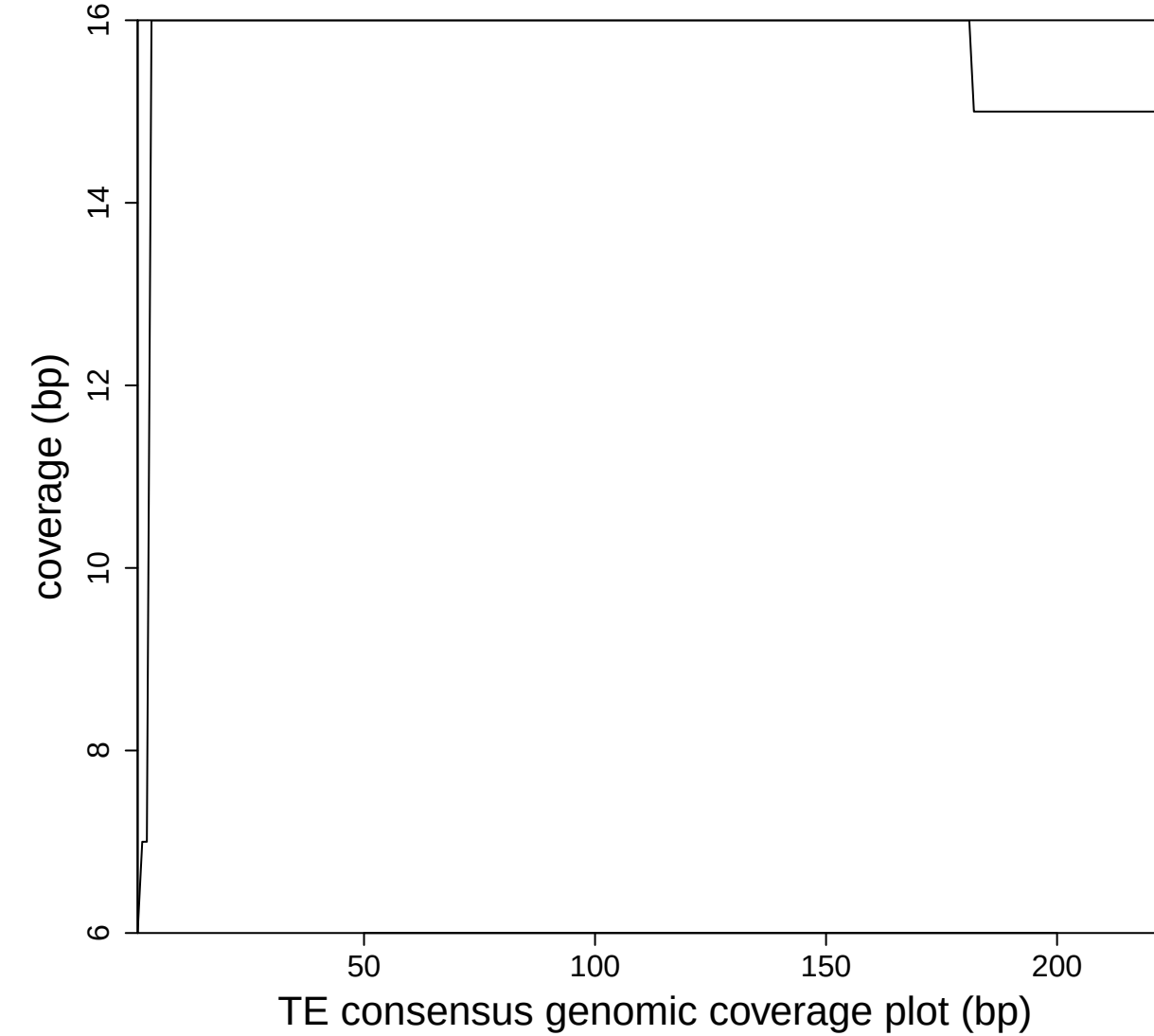
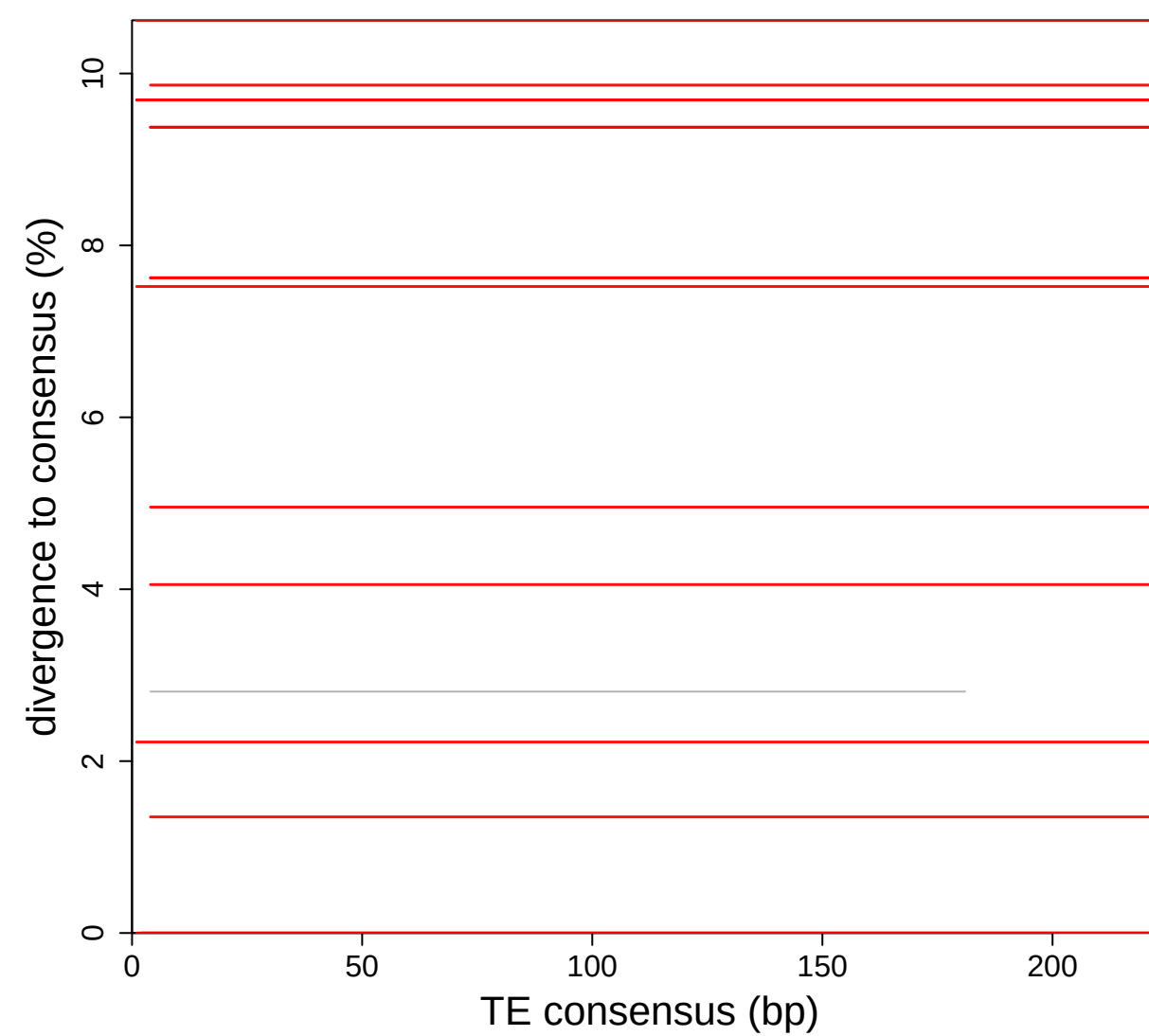
fasta.b.bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_
size: 1918bp; fragments: 19; full length: 0 (>=1726.2bp)

After TEtrimmer Extended plot Blue lines are boundaries

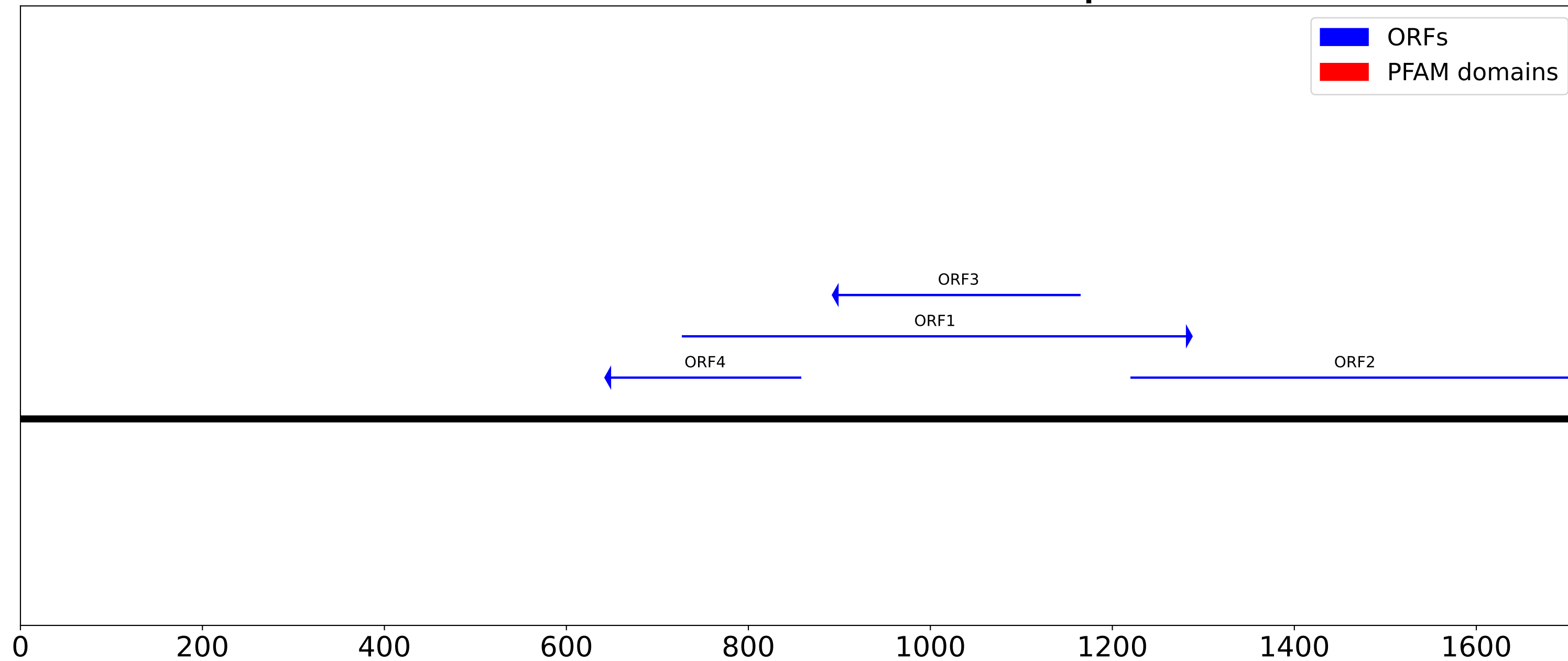


Before TEtrimmer 225 bp

TE: ltr_1_family_58
size: 225bp; fragments: 16; full length: 15 (≥ 202.5 bp)



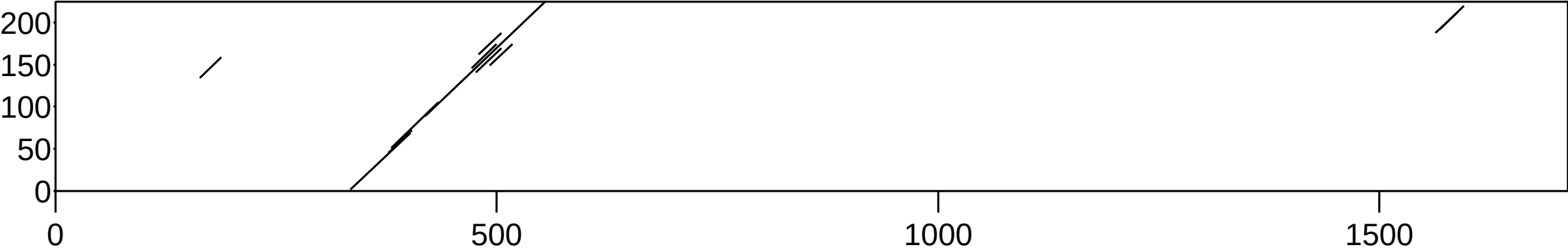
After TEtrimmer ORF and PFAM domain plot



Dotplot

(windowsize = 25, threshold = 50.00 09/10/25)

sensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)