

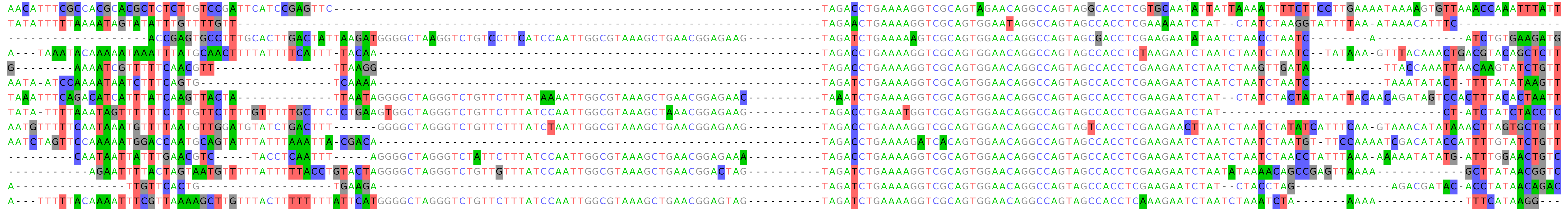
Start crop Point



End crop Point



MSA length = 4256



Start crop Point

End crop Point

0 -

10

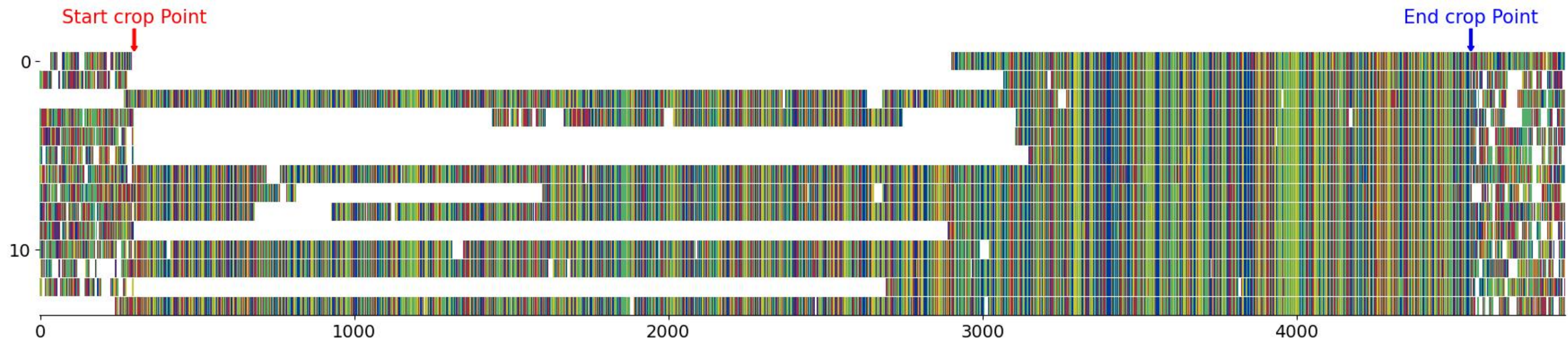
0

1000

2000

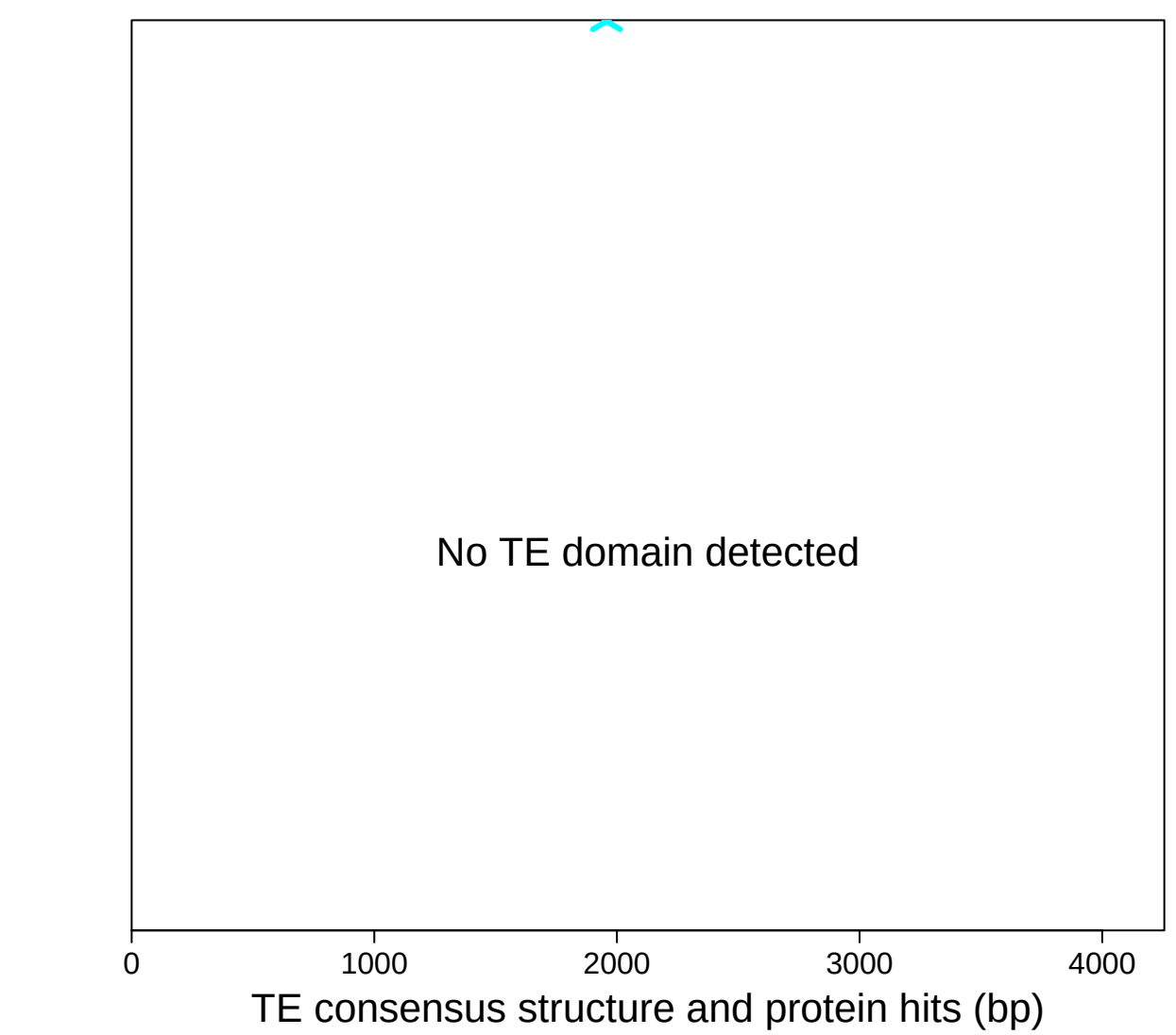
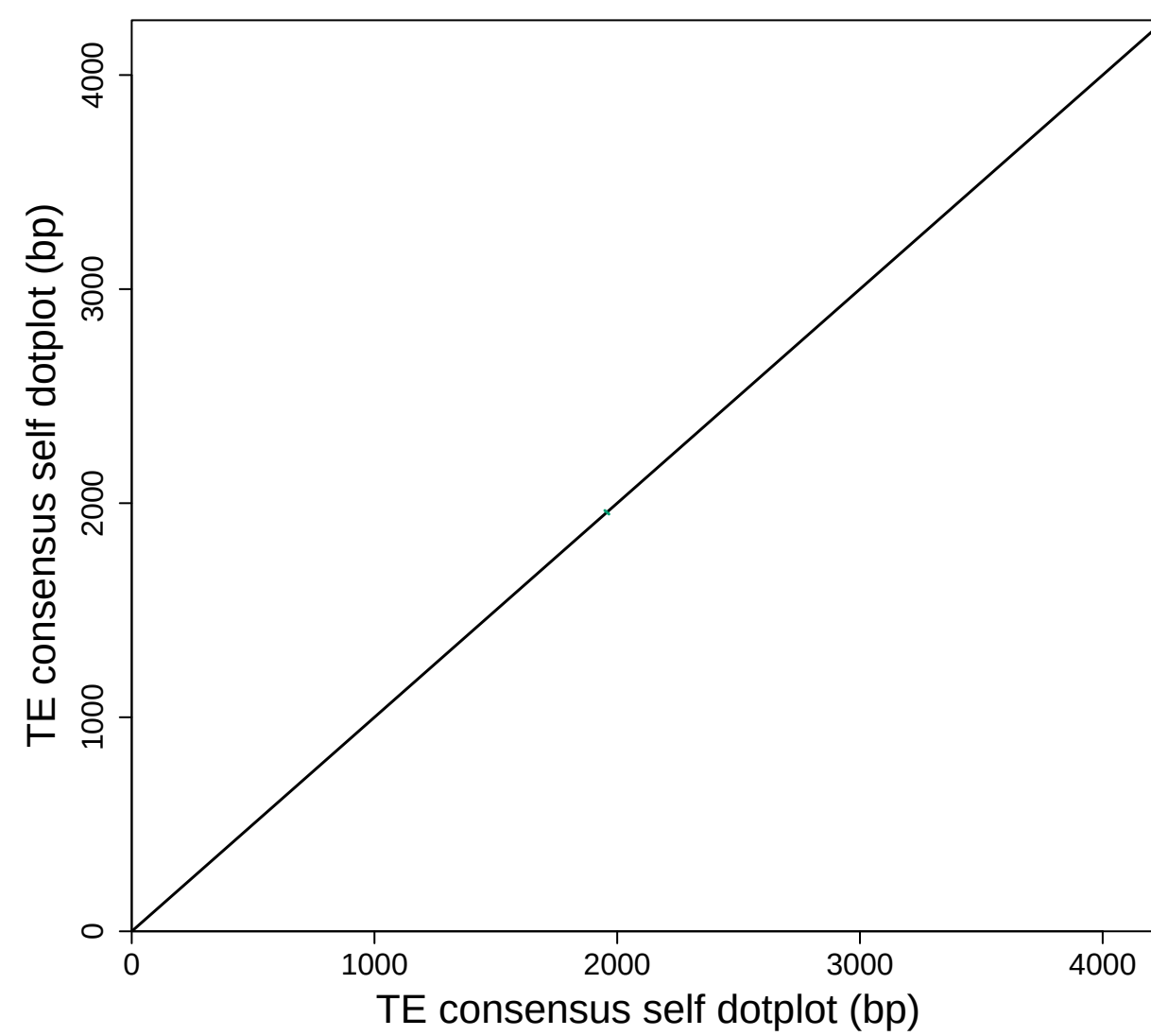
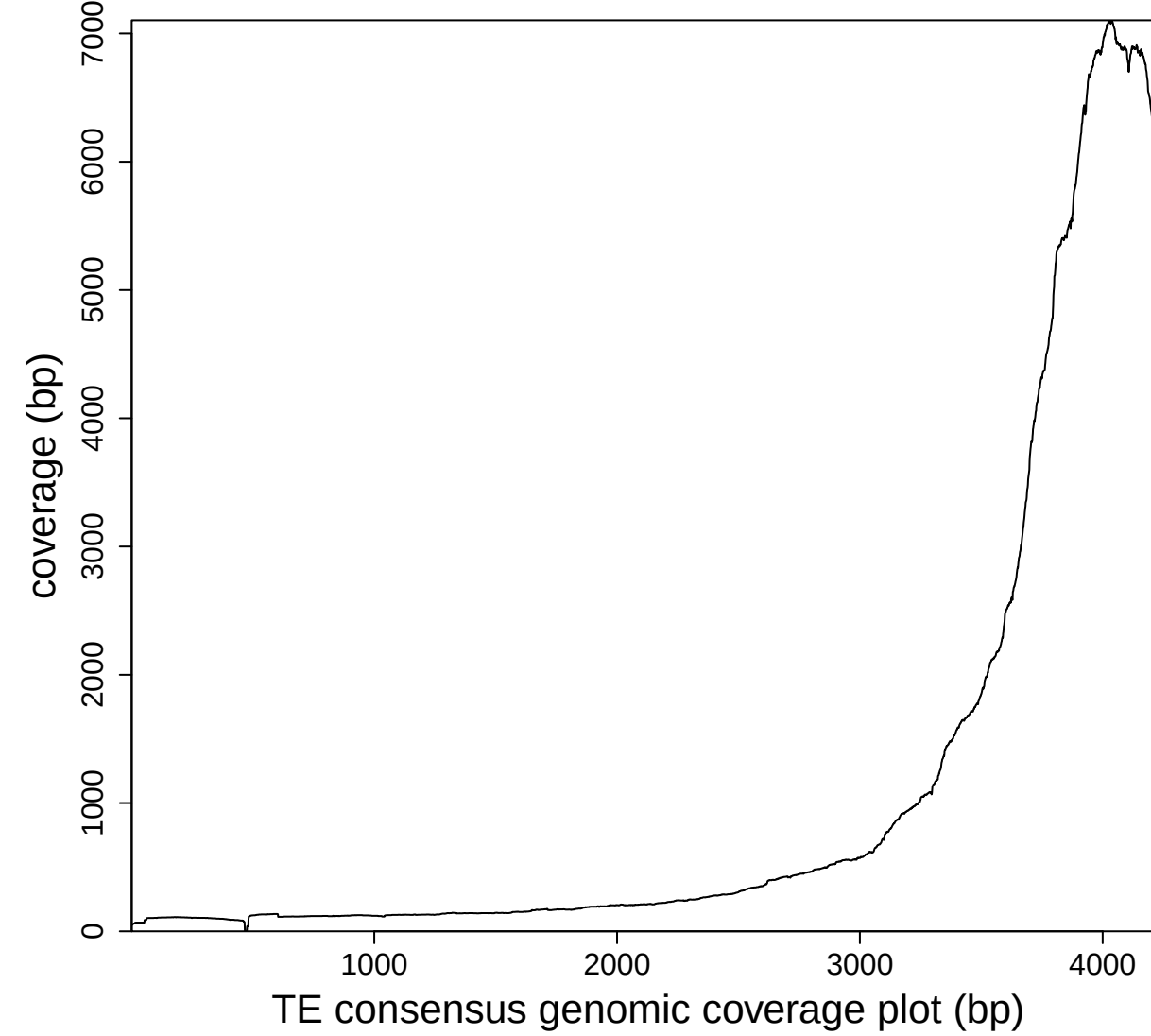
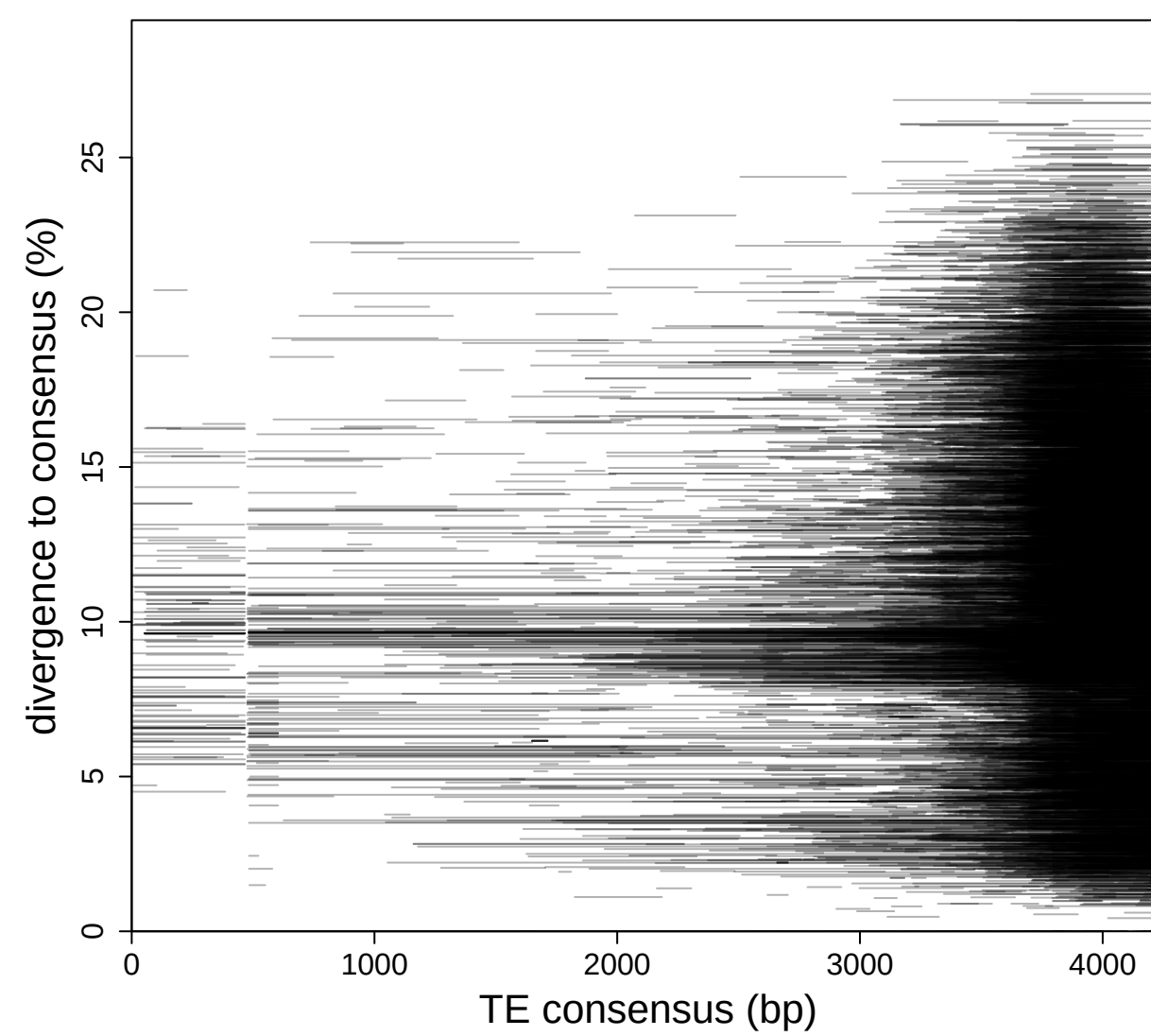
3000

4000



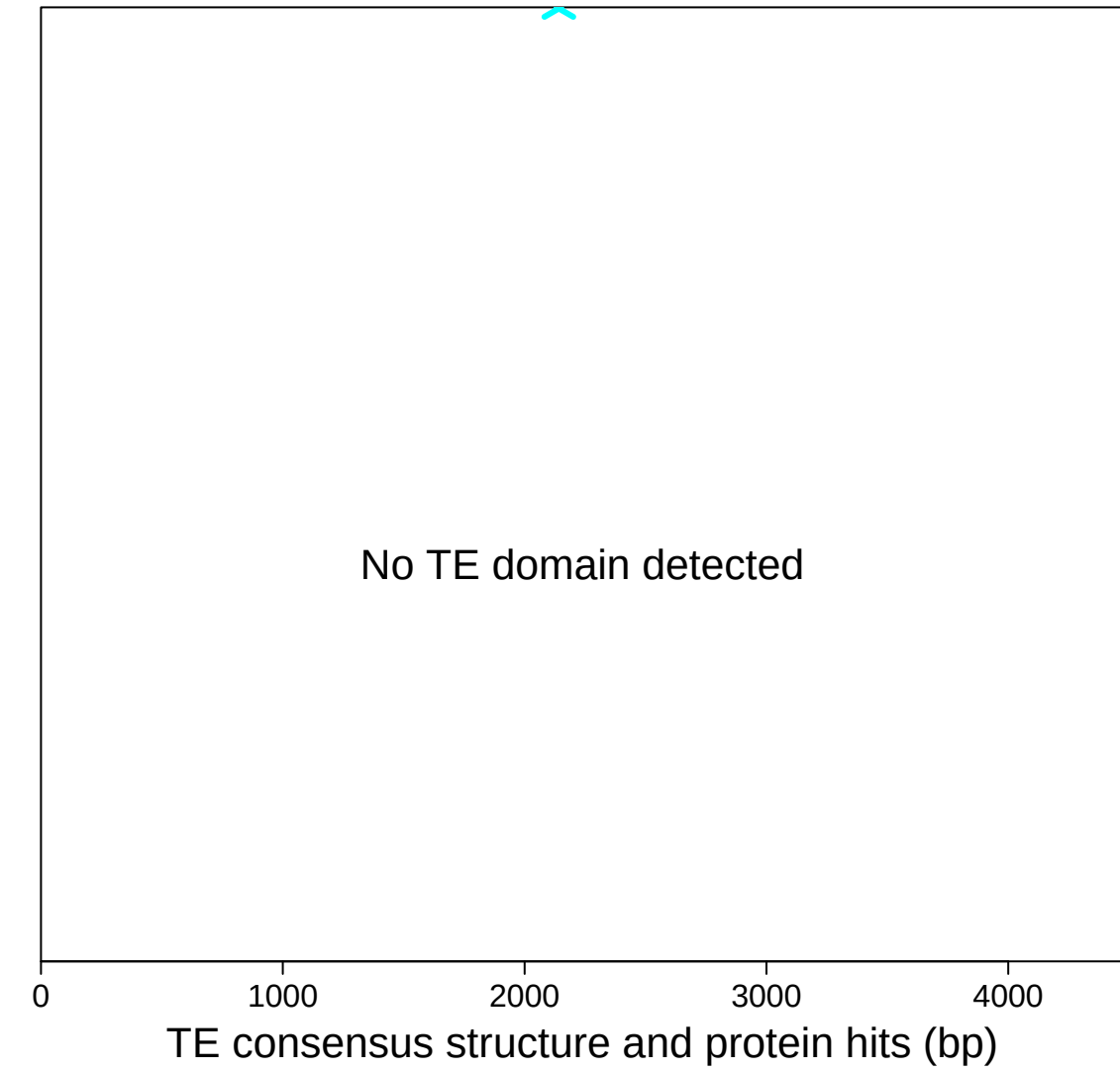
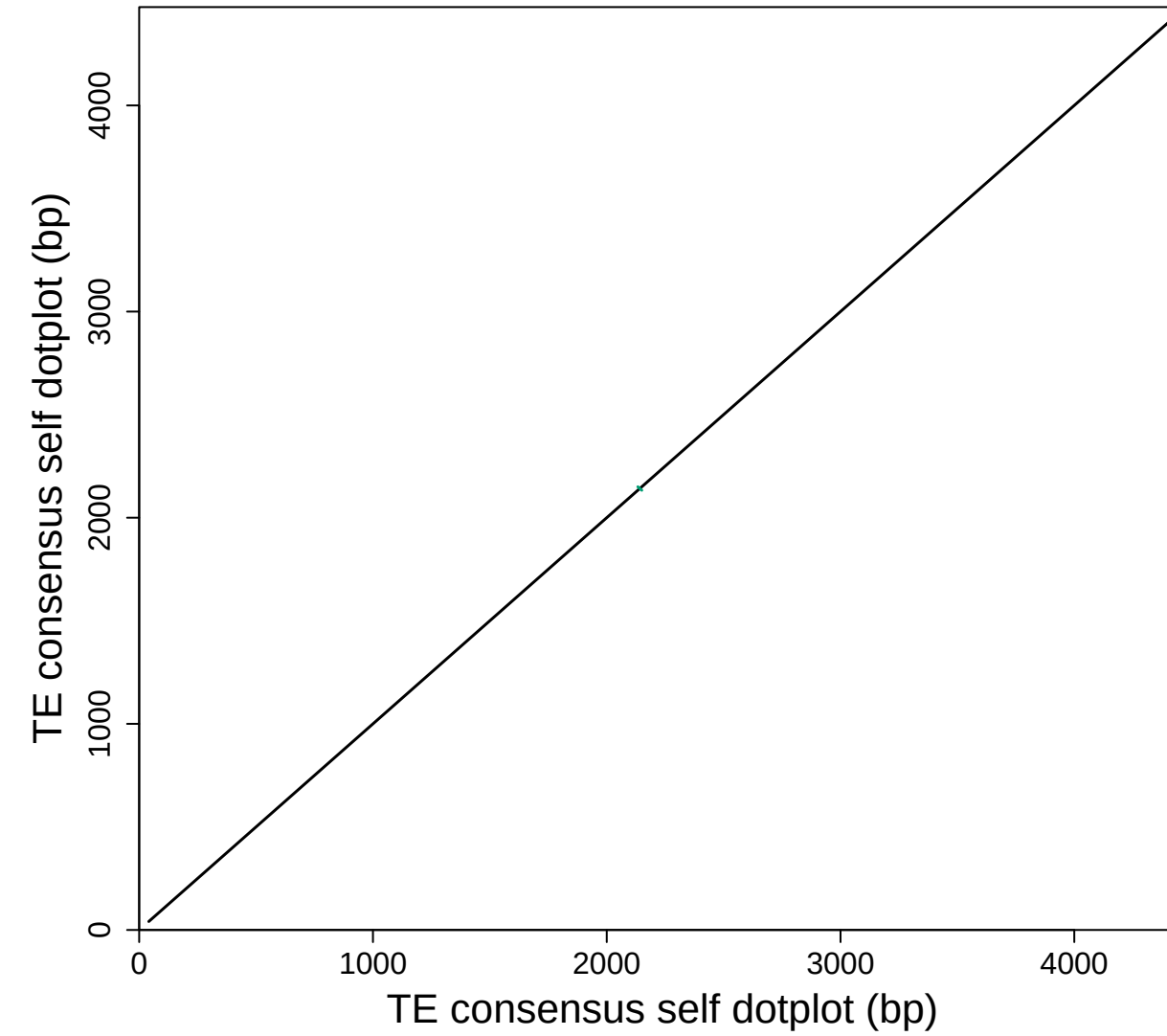
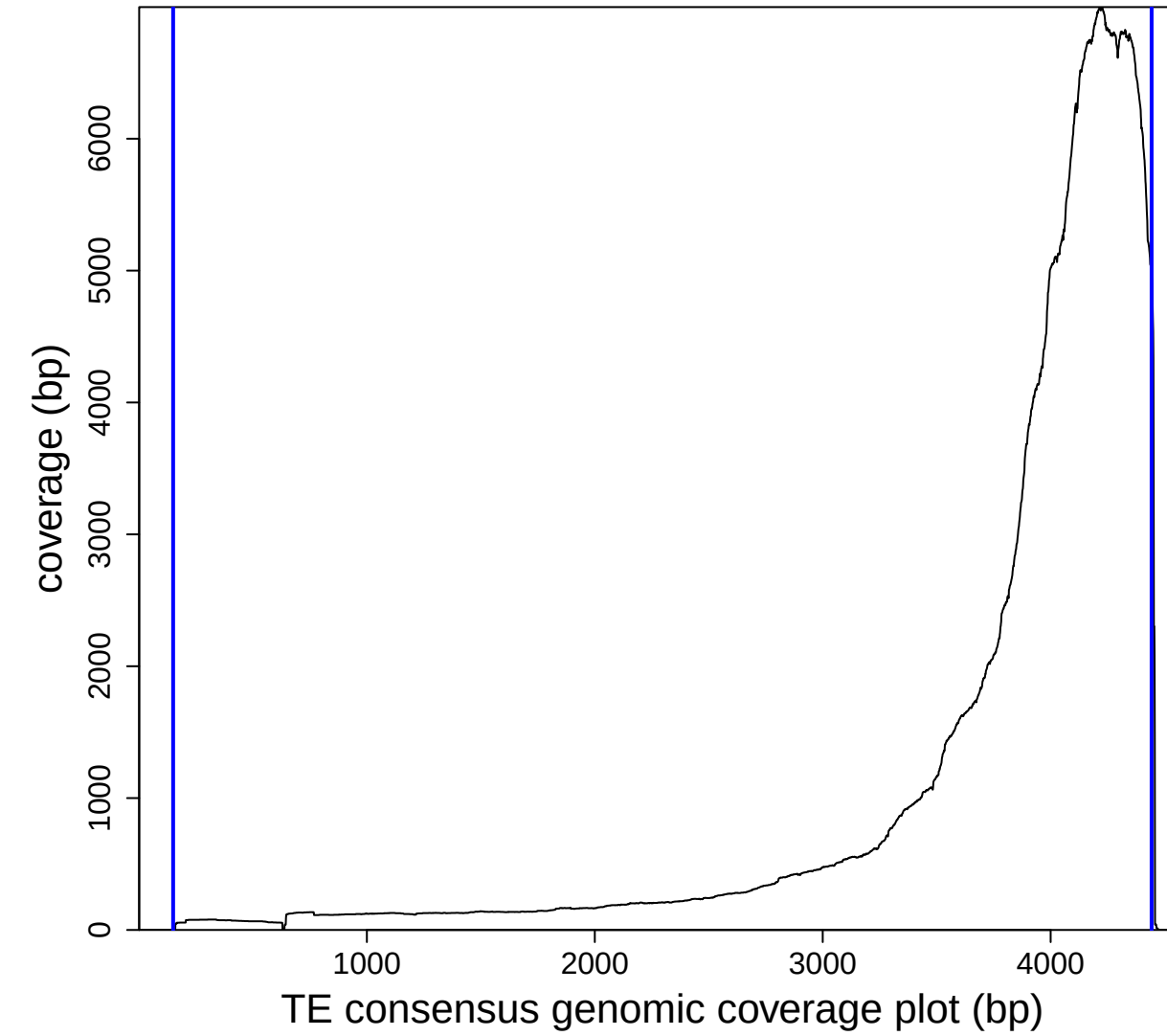
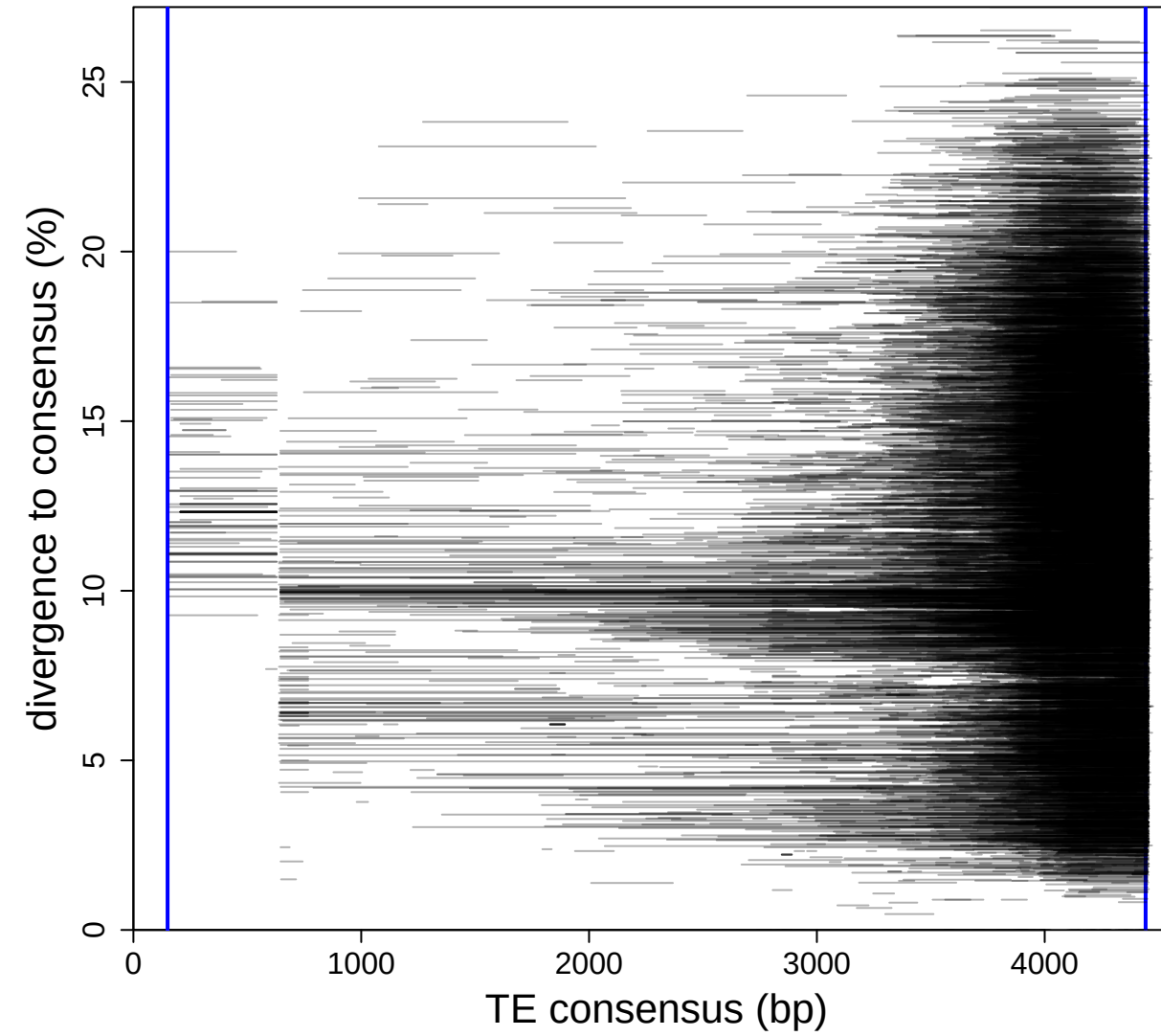
bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcln.fa_aln.fa
size: 4256bp; fragments: 12291; full length: 0 (>=3830.4bp)

After TEtrimmer 4256 bp



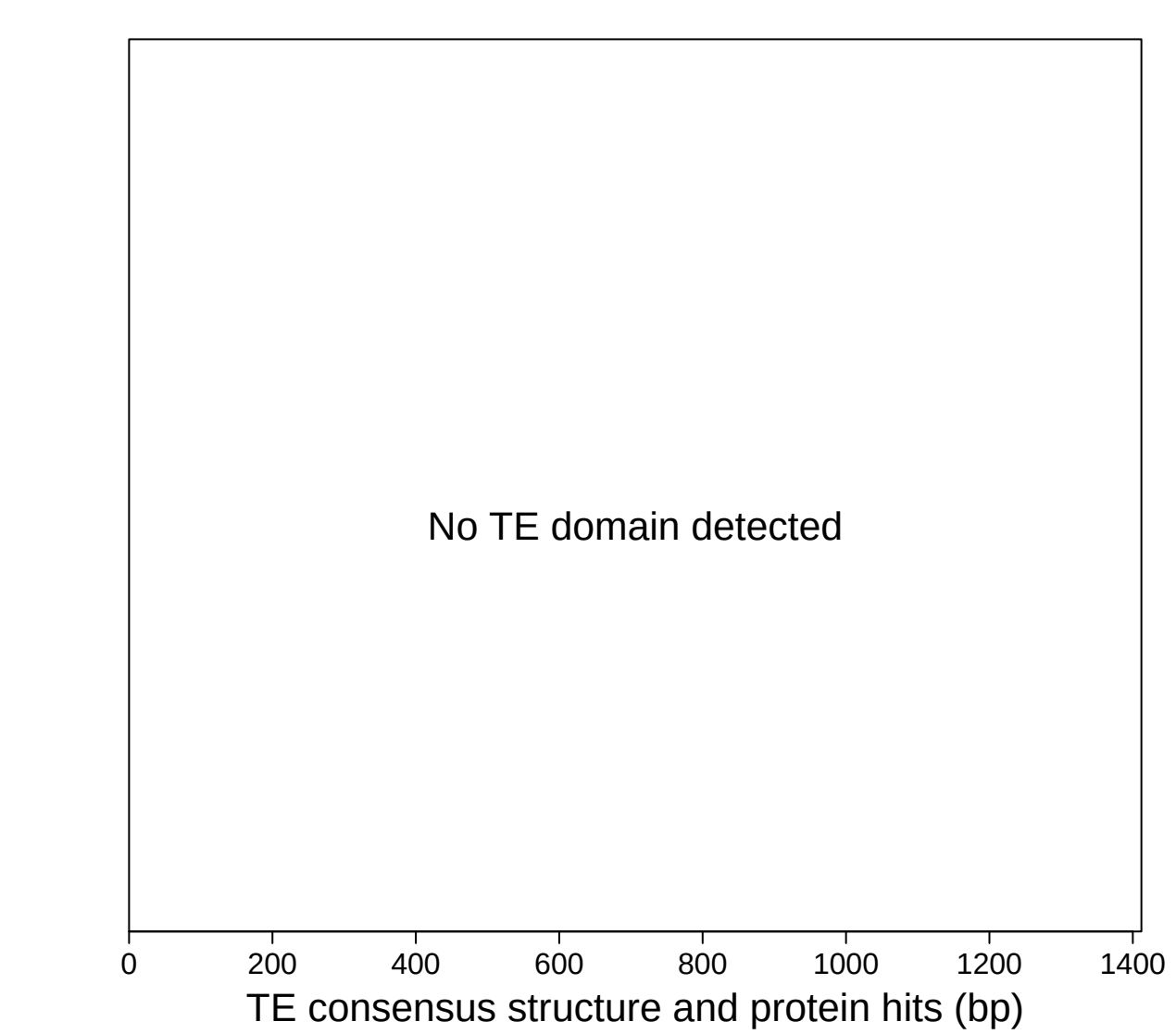
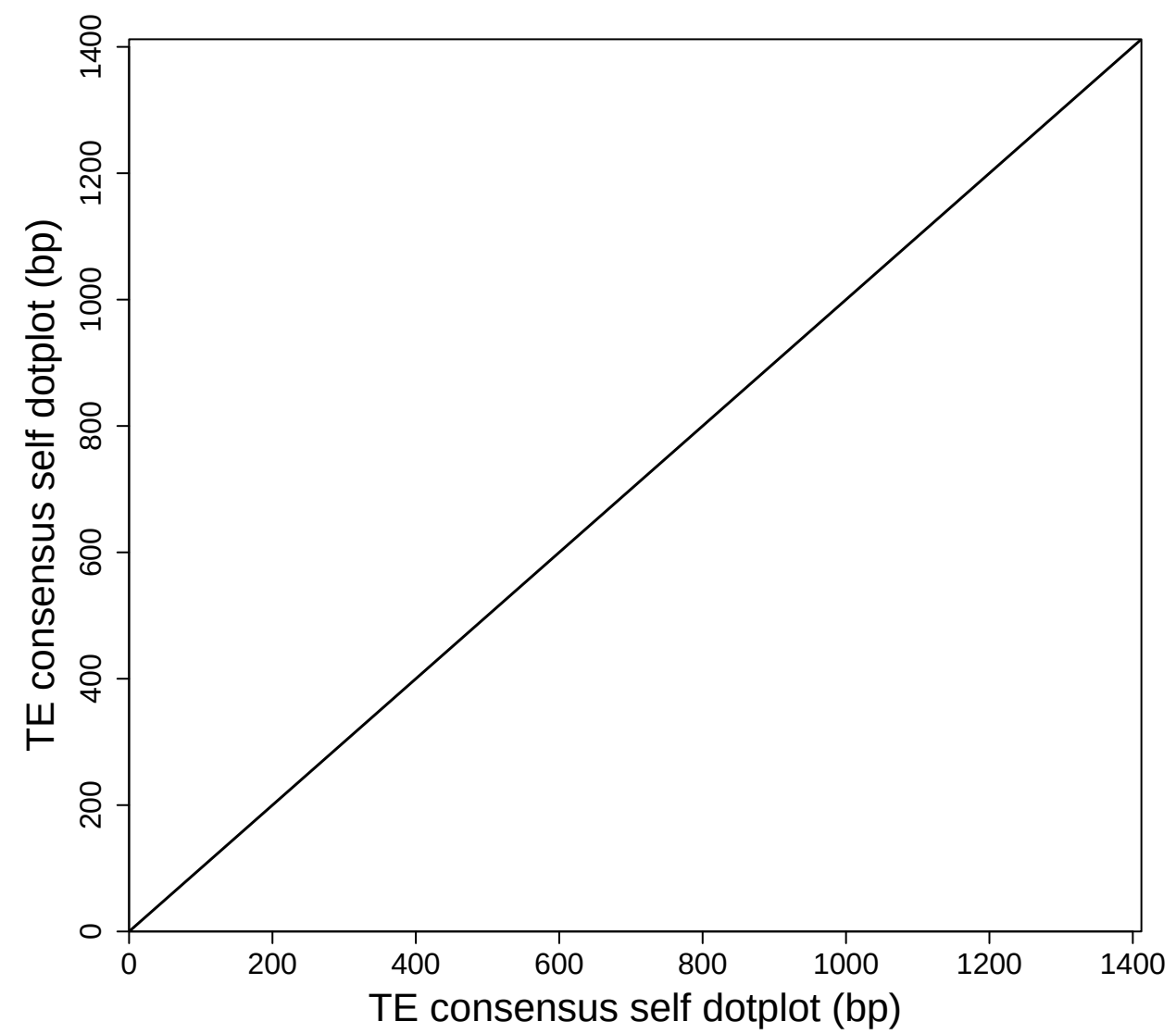
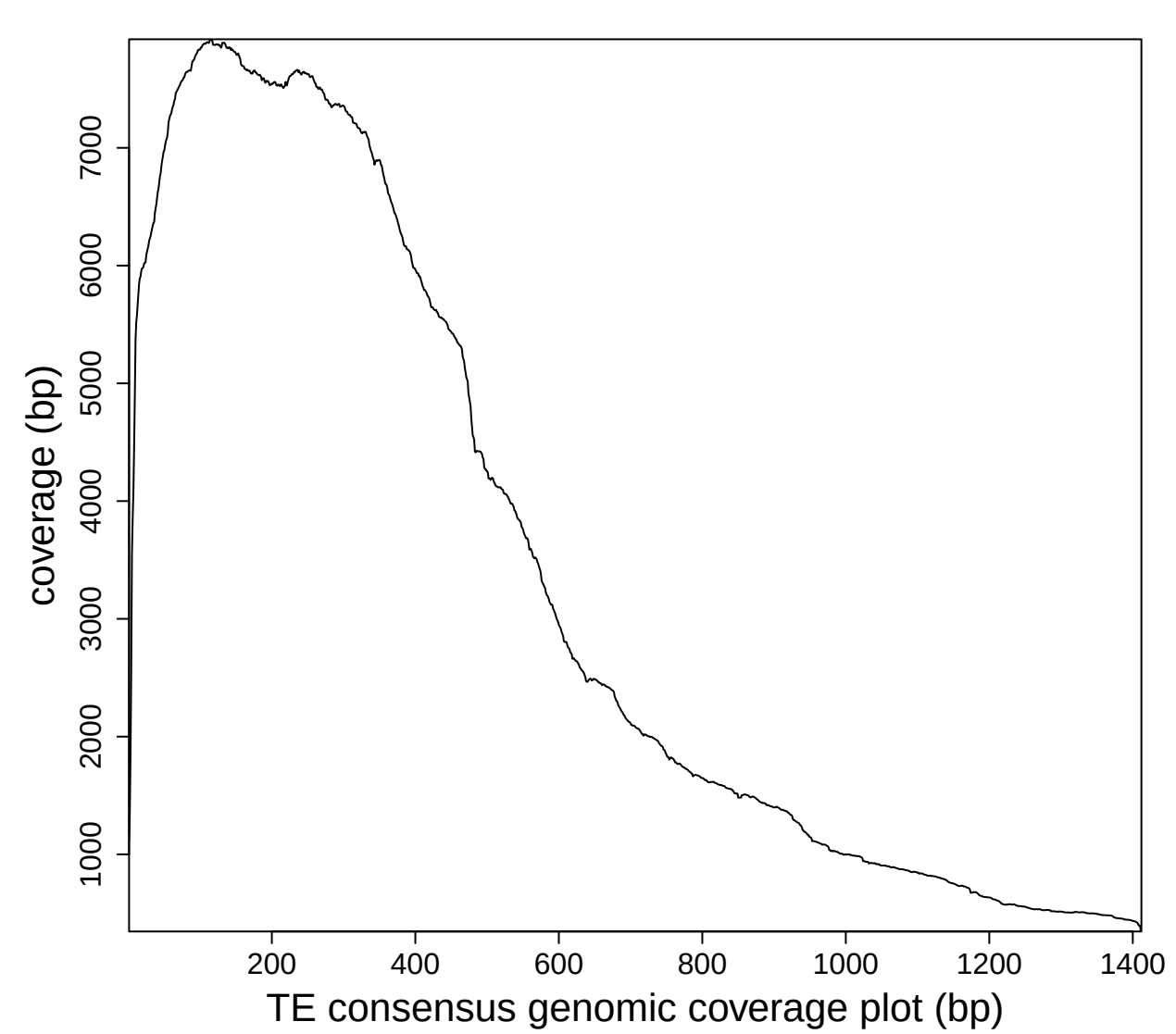
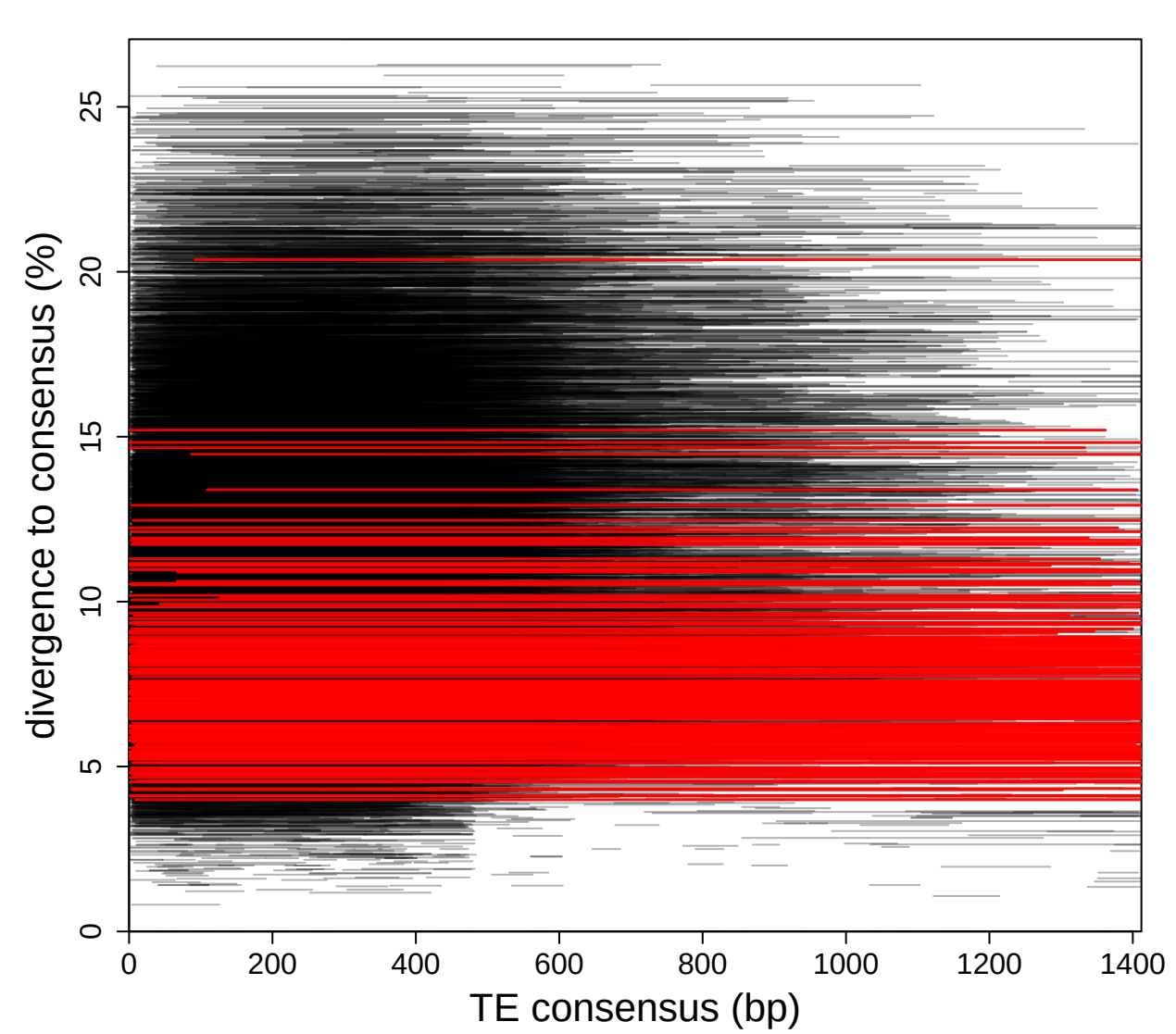
d_g_3.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcln.fa
size: 4593bp; fragments: 11840; full length: 0 (>=4133.7bp)

After TEtrimmer Extended plot Blue lines are boundaries

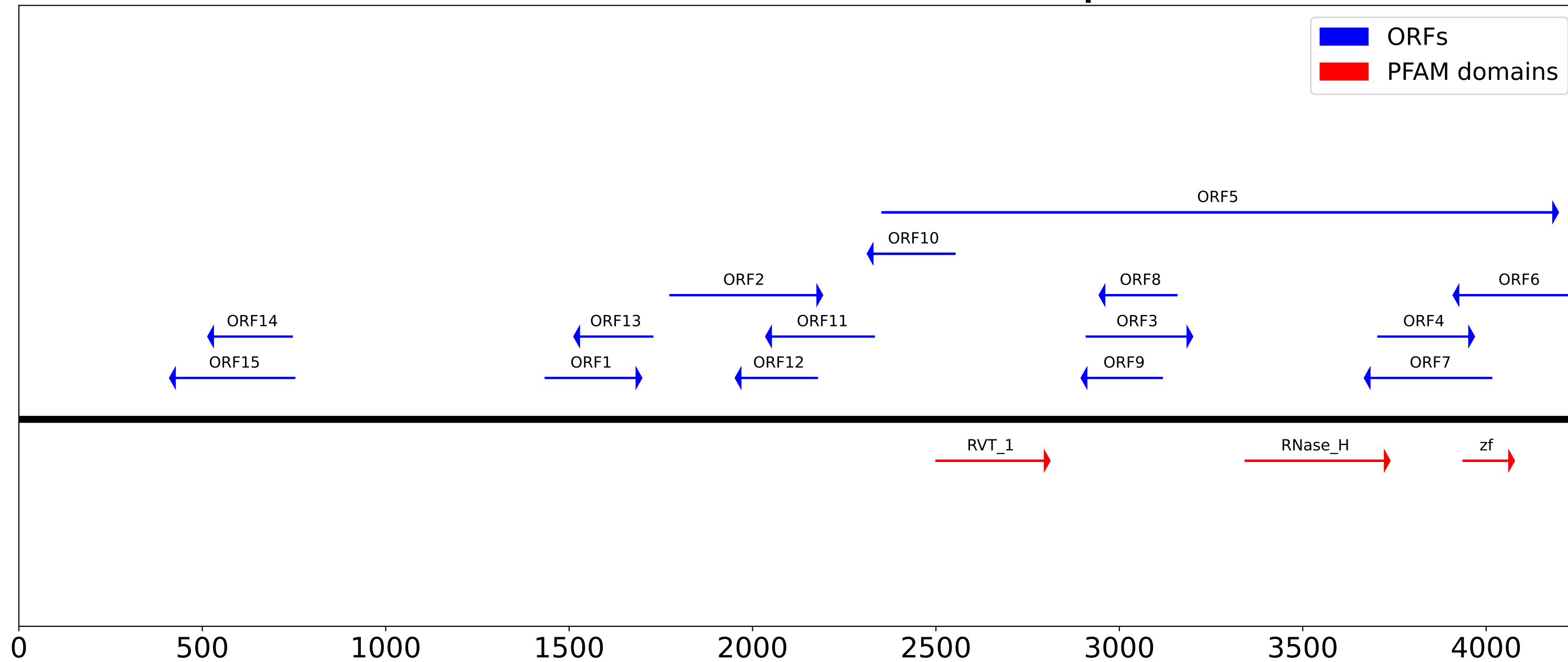


TE: rnd_1_family_93
size: 1412bp; fragments: 12810; full length: 263 (≥ 1270.8 bp)

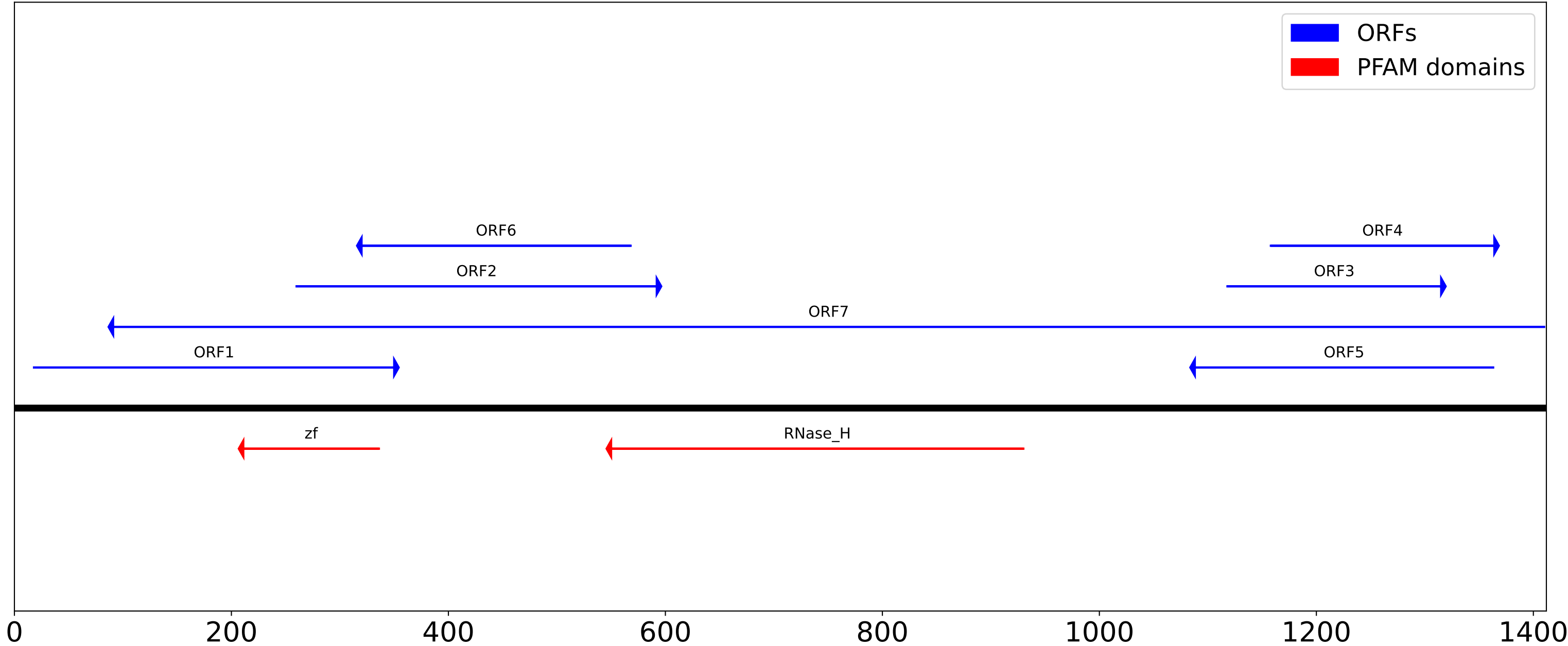
Before TEtrimmer 1412 bp



After TEtrimmer ORF and PFAM domain plot



Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)

