

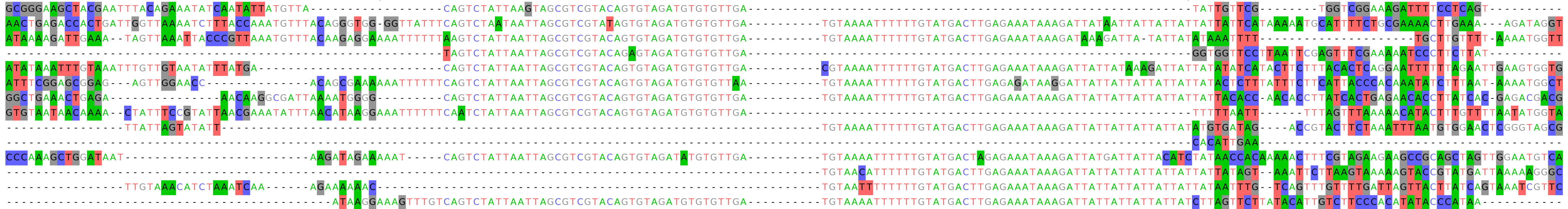
Start crop Point



End crop Point



MSA length = 2005



Start crop Point

End crop Point

0 -

10 -

0

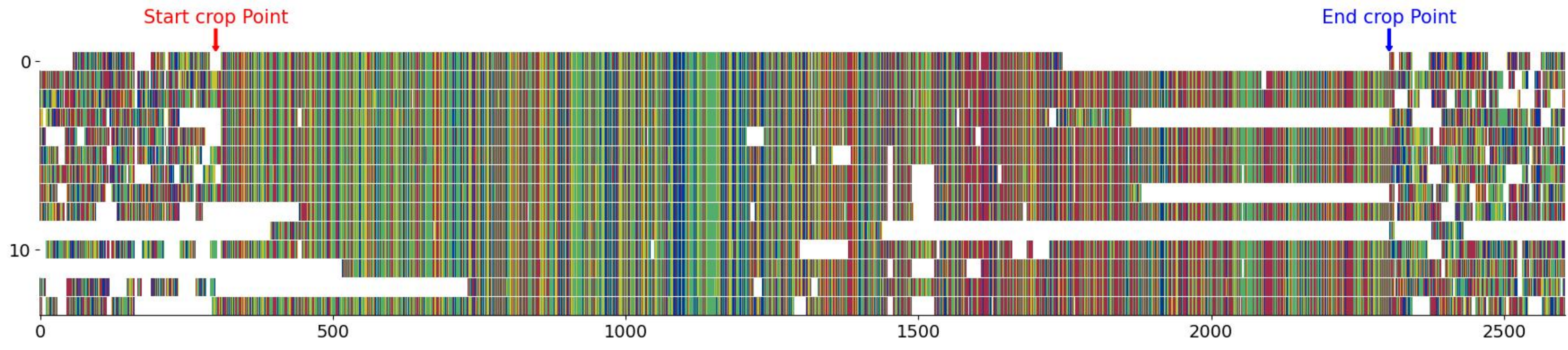
500

1000

1500

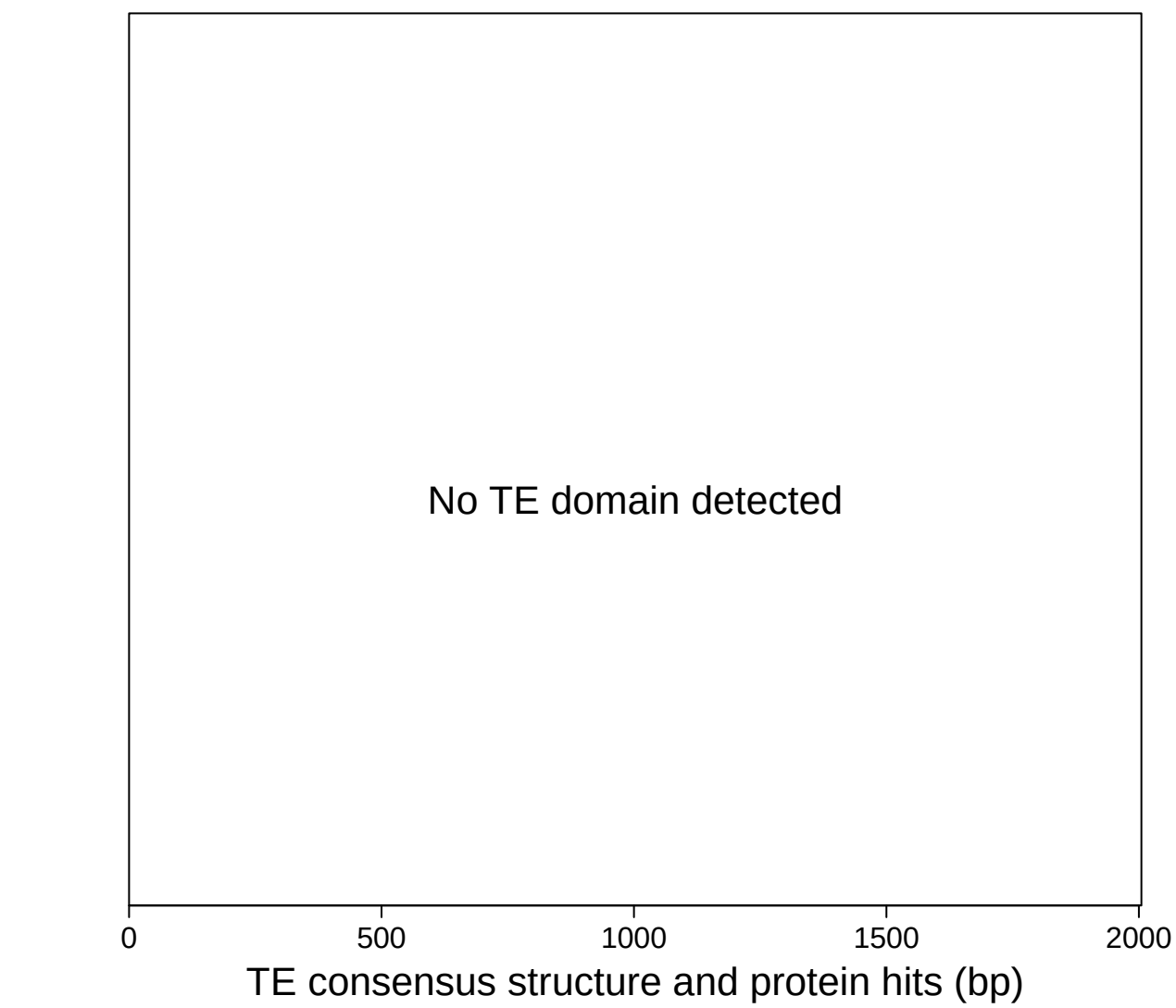
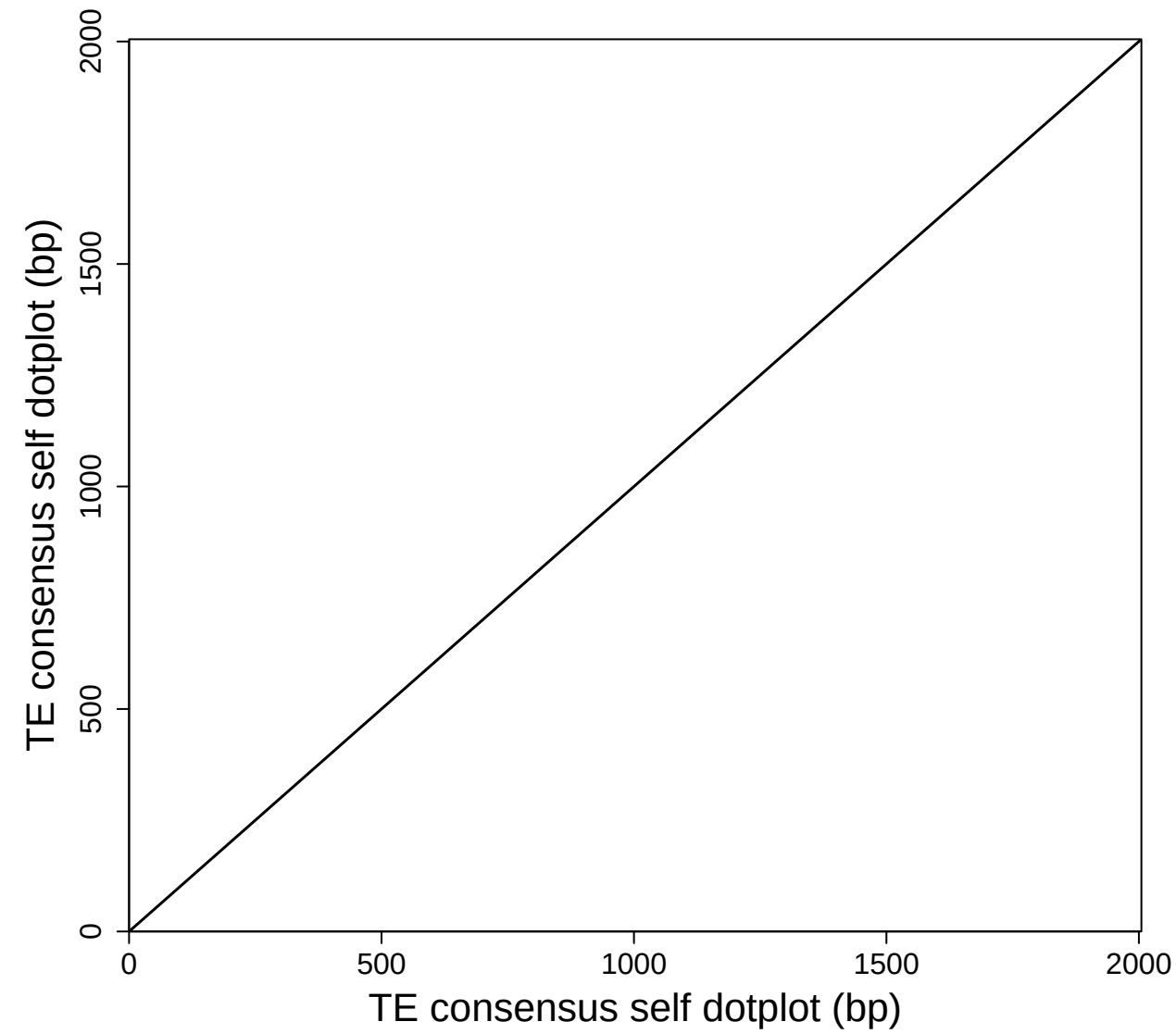
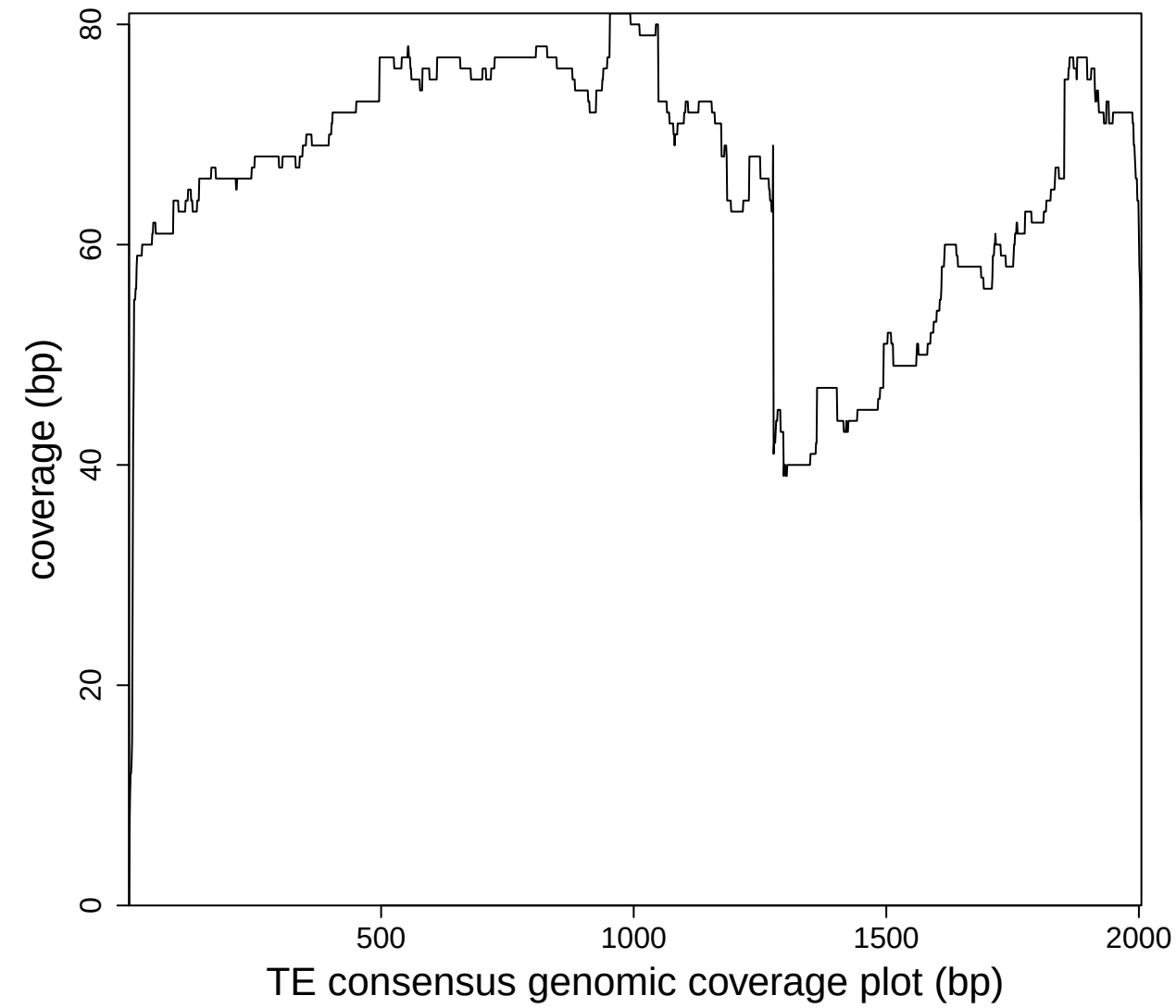
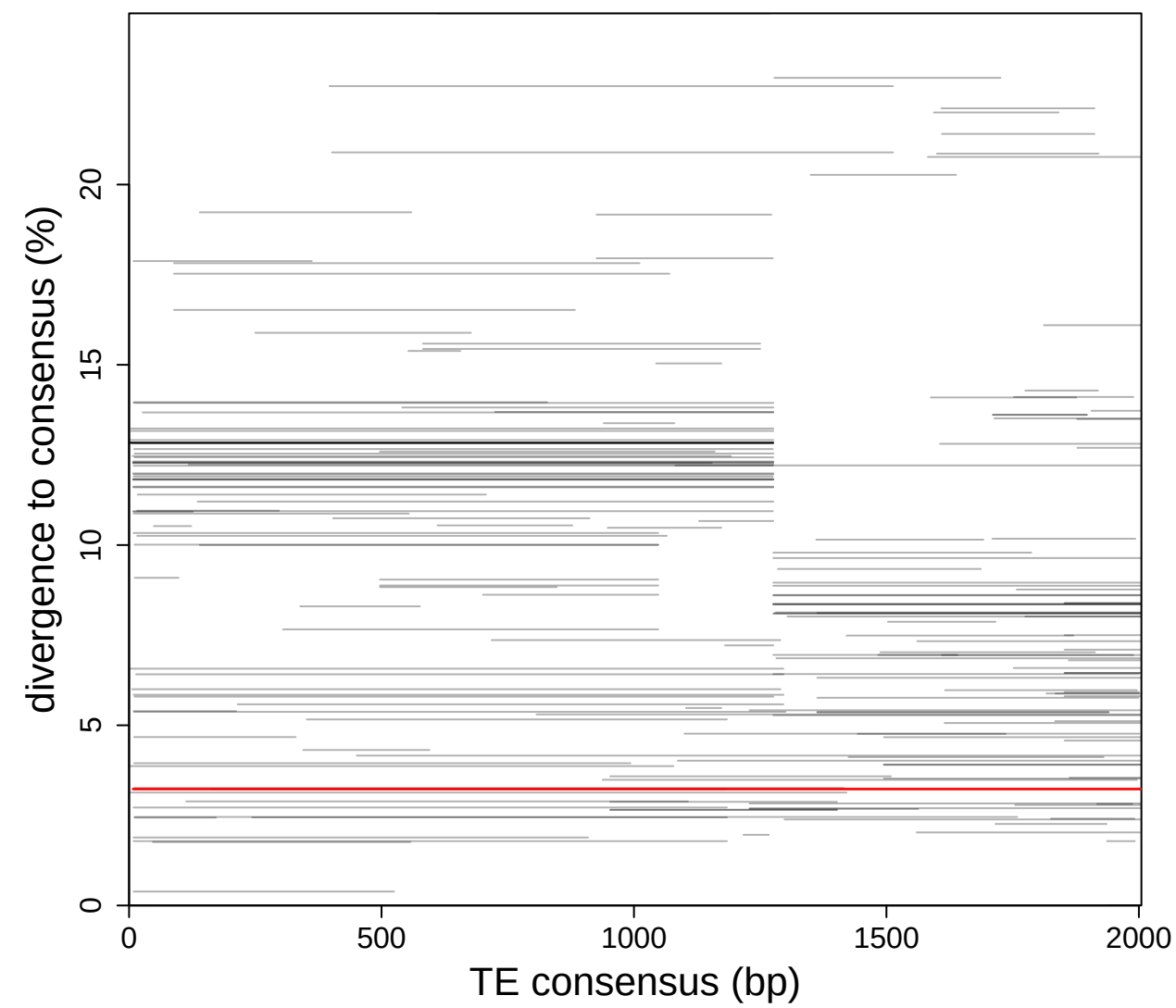
2000

2500



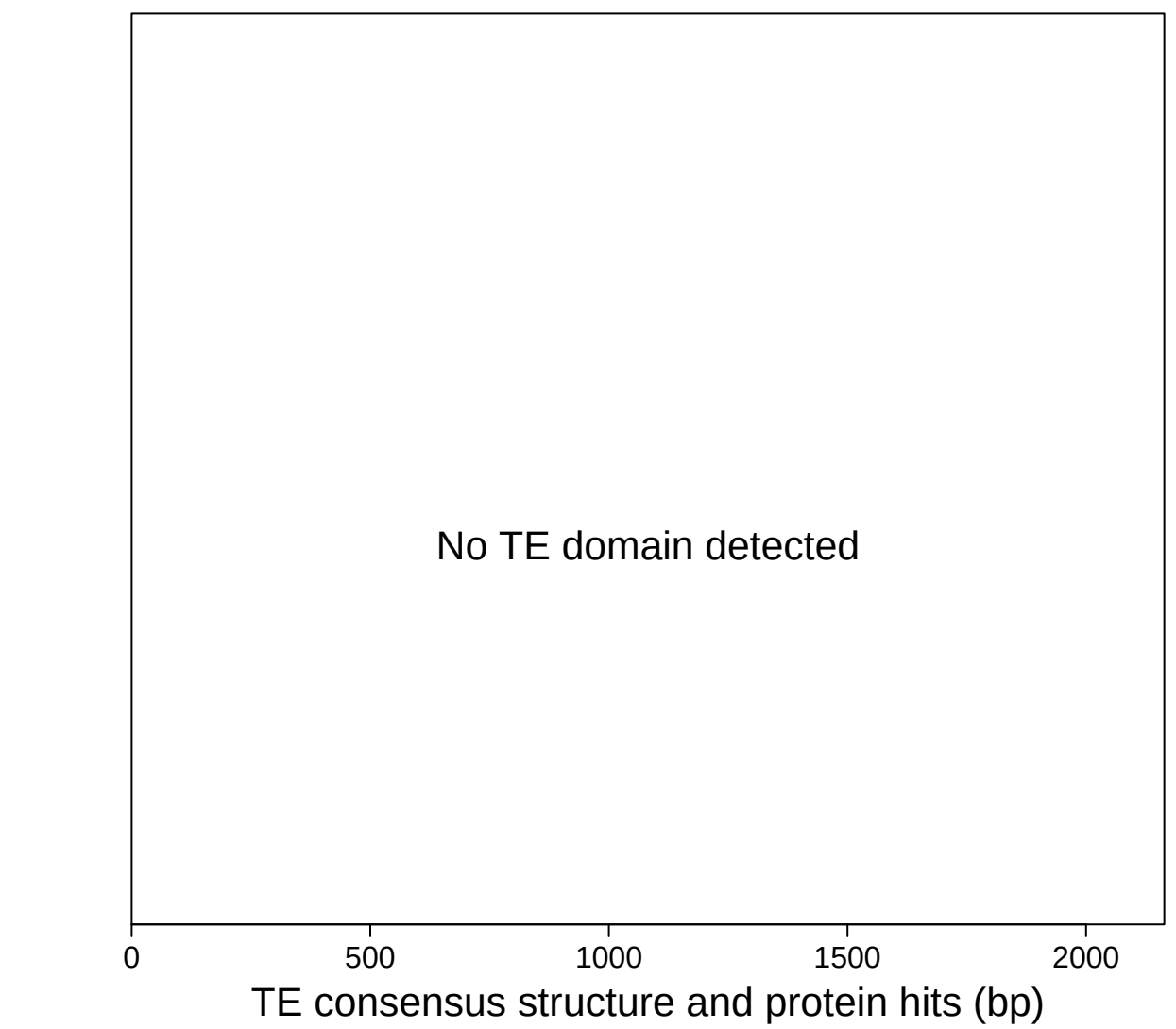
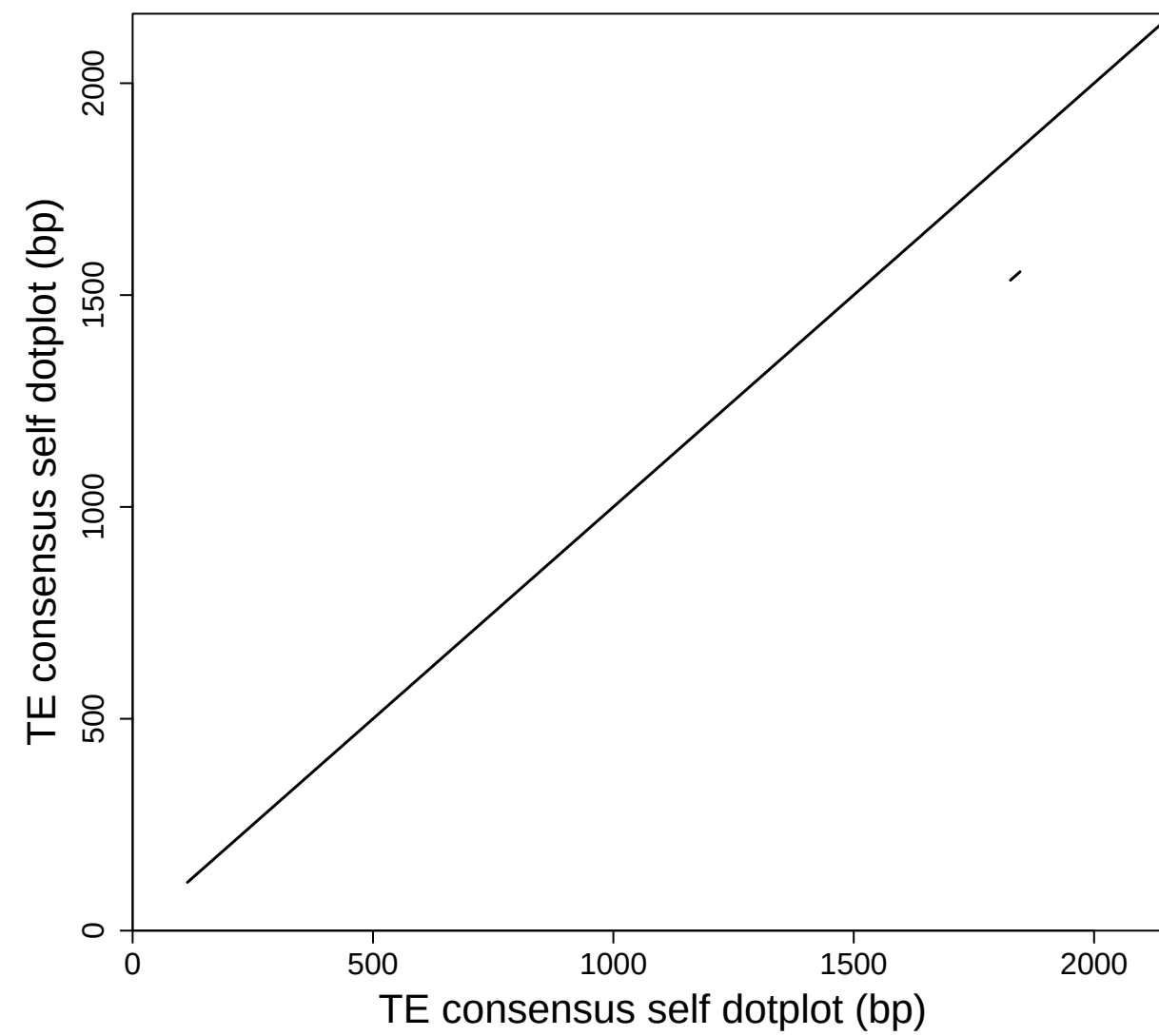
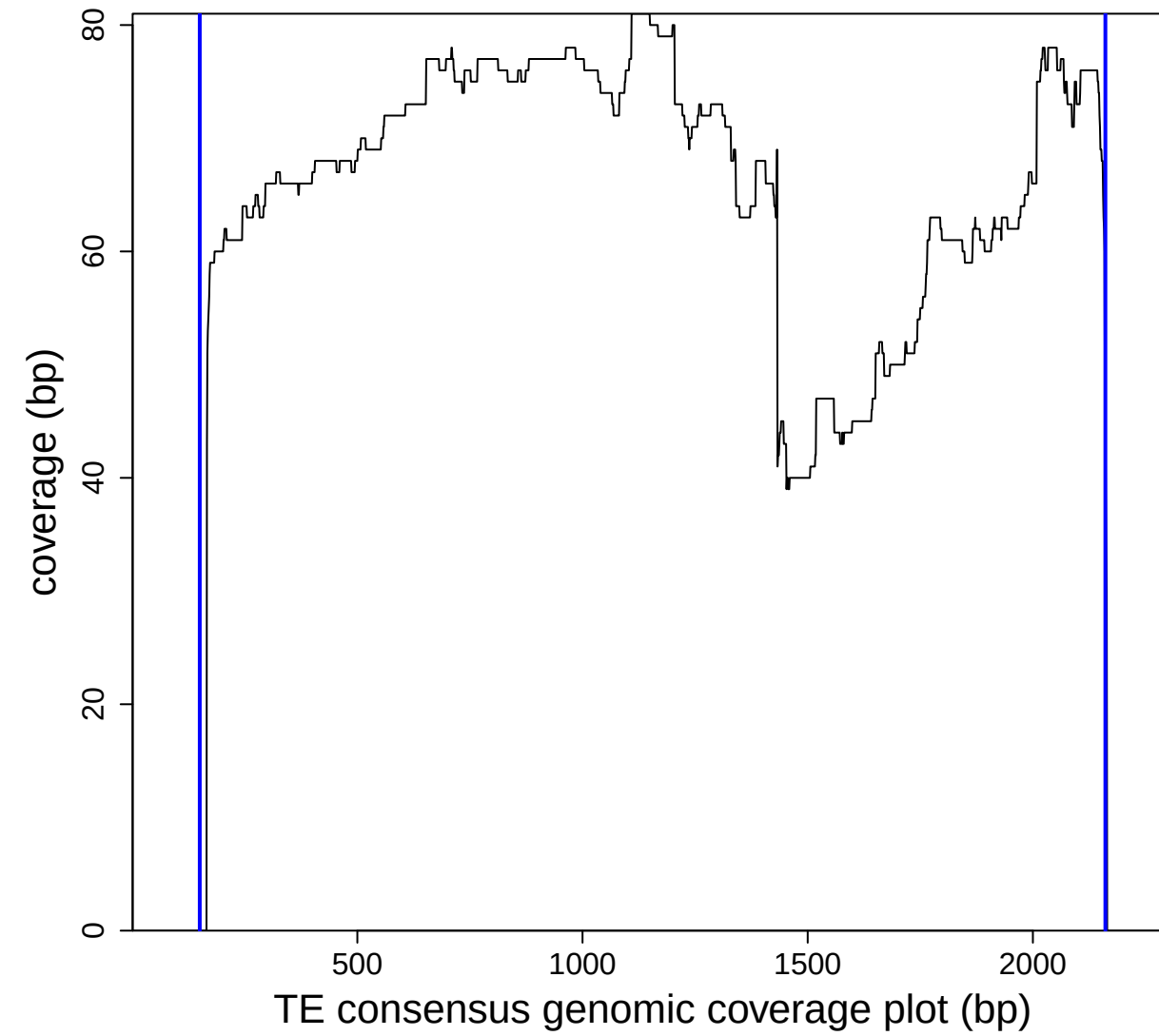
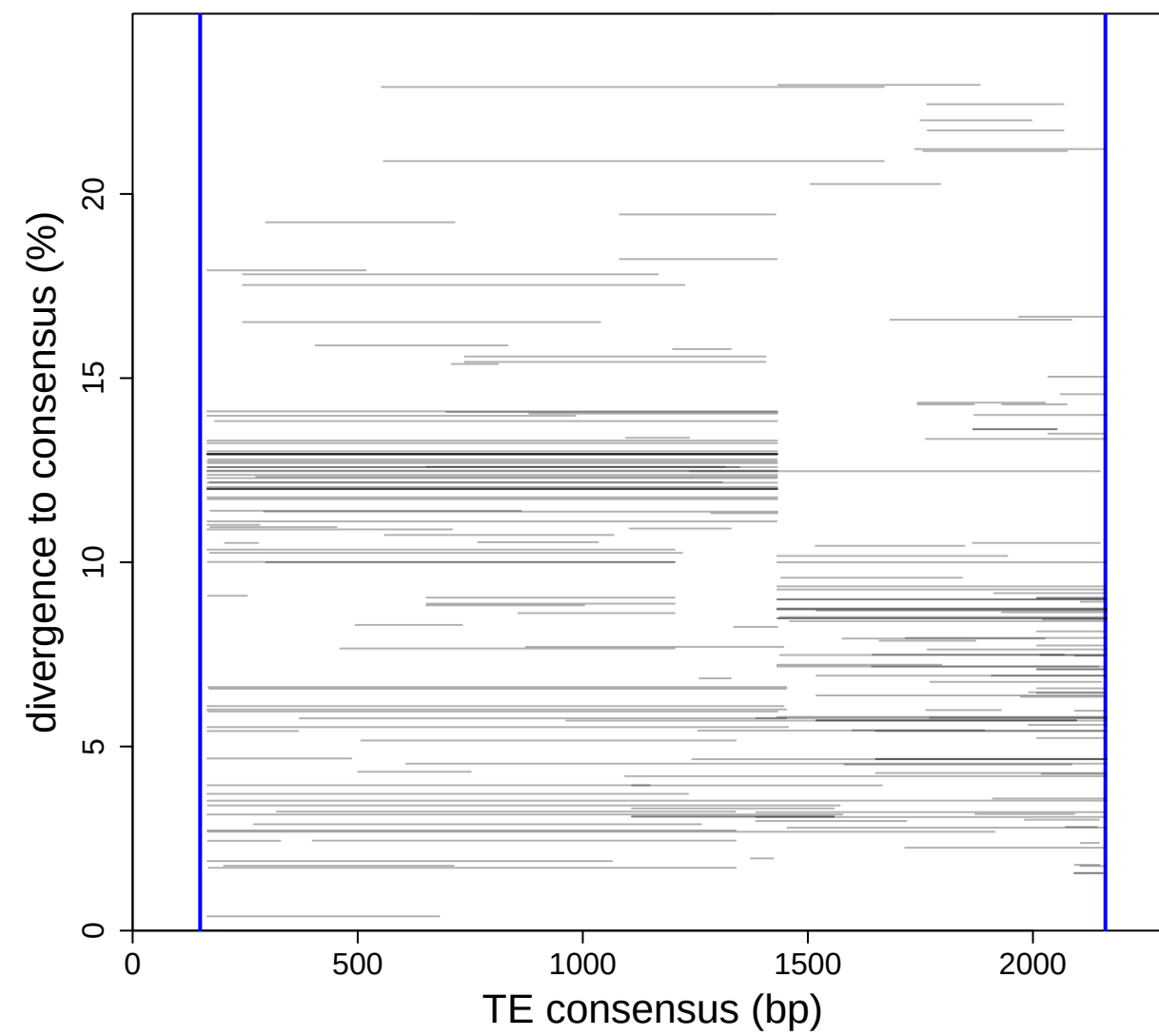
bed_uf.bed_g_2.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.fa_c
size: 2005bp; fragments: 201; full length: 1 (>=1804.5bp)

After TEtrimmer 2005 bp



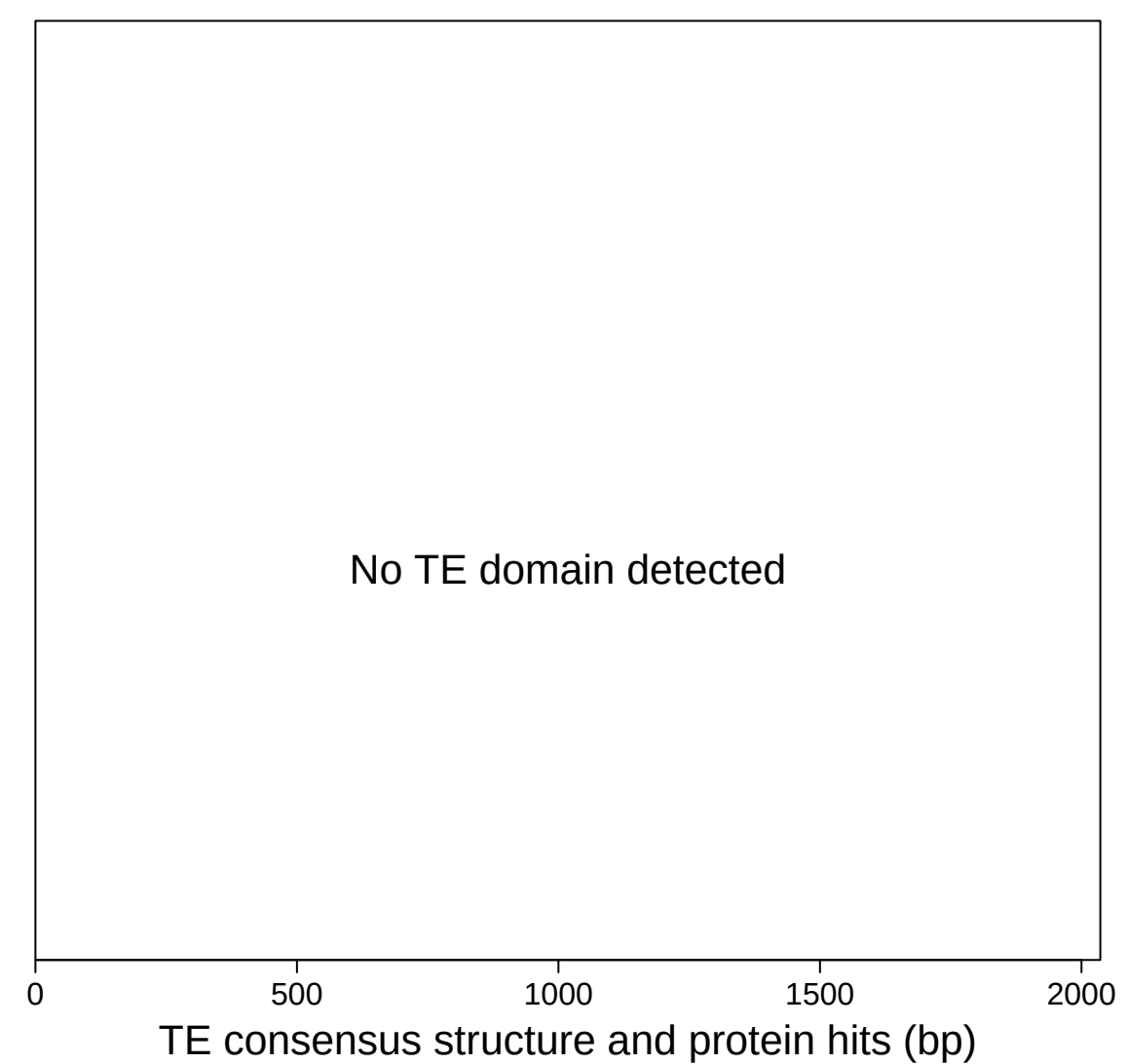
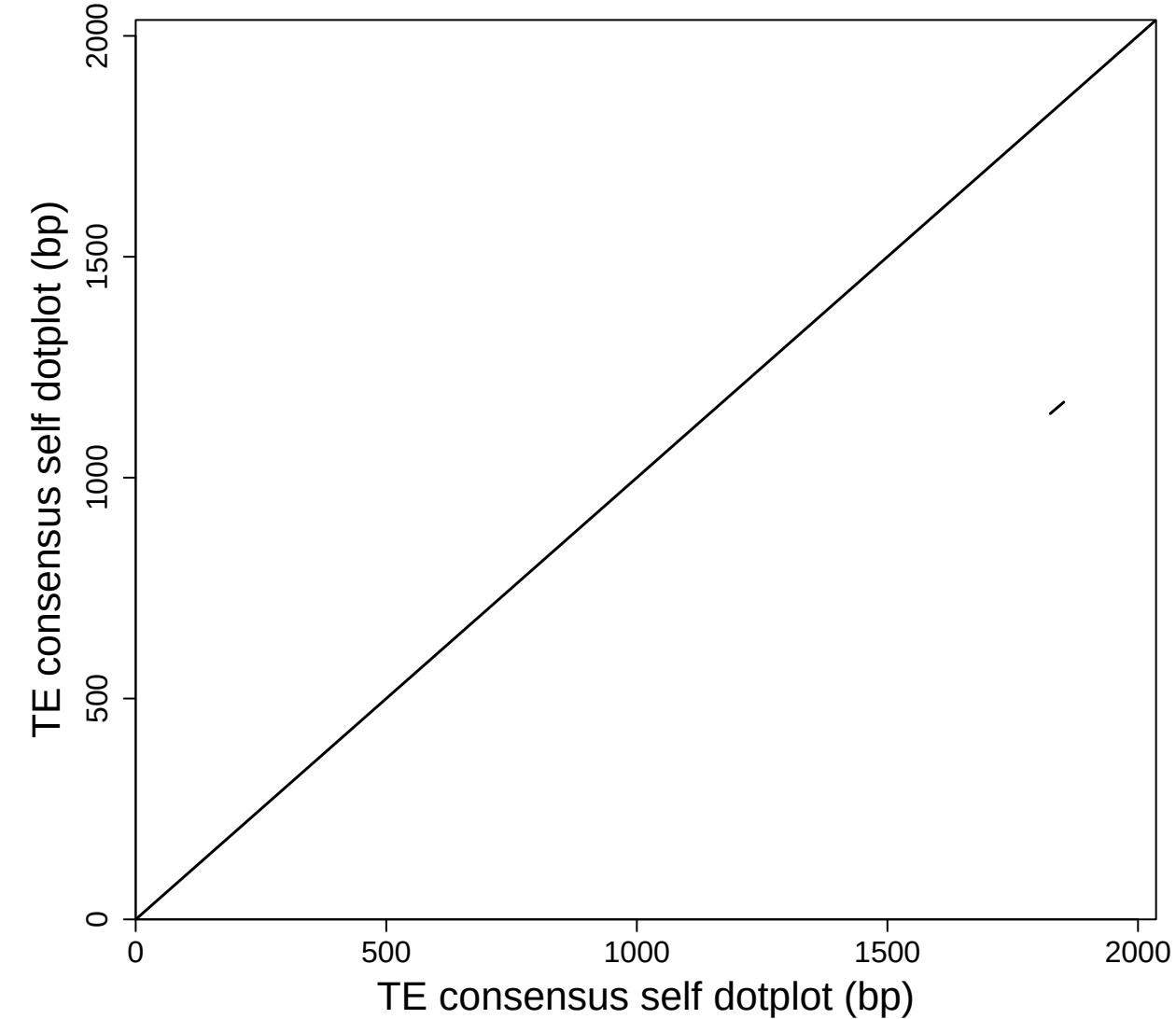
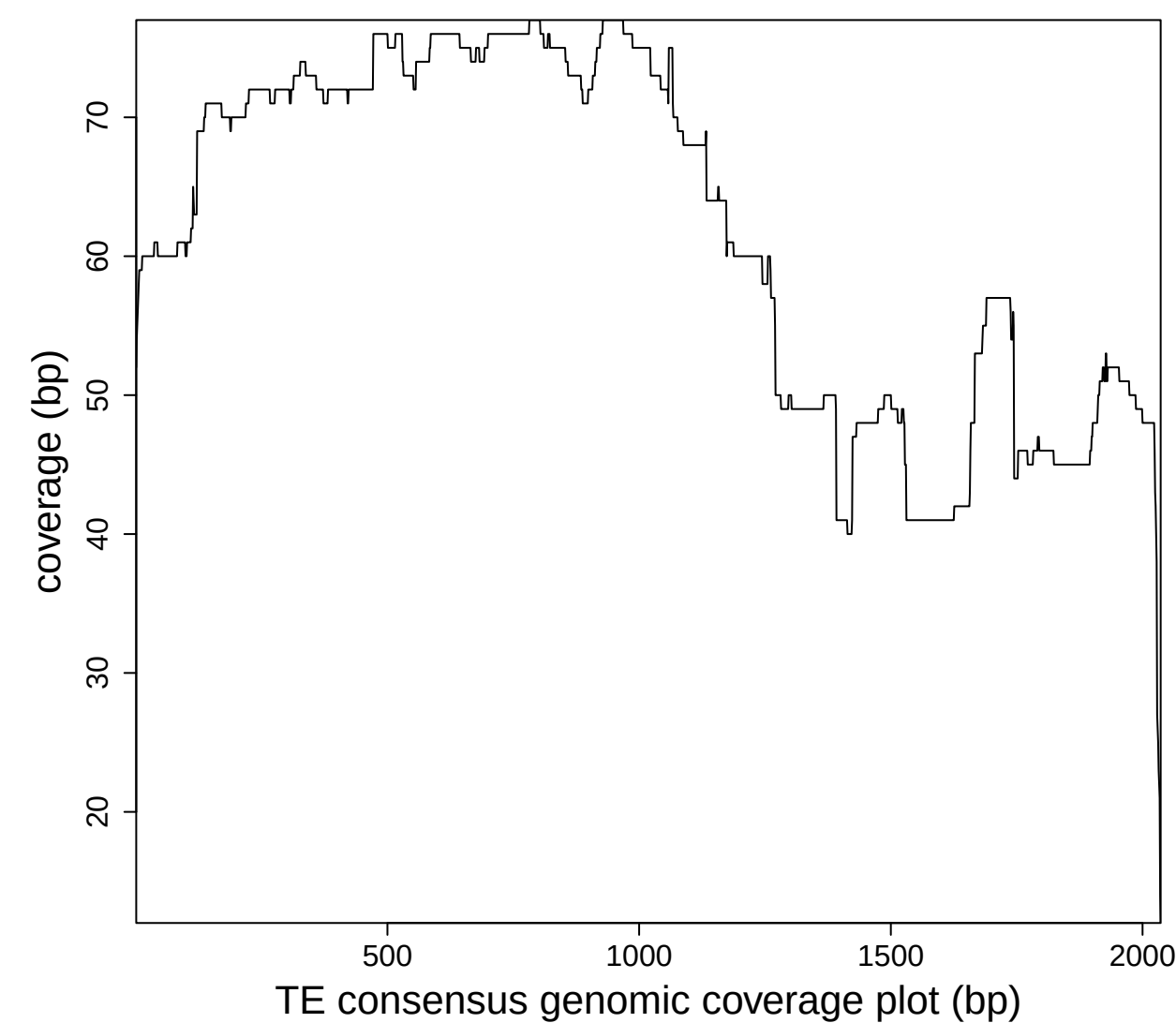
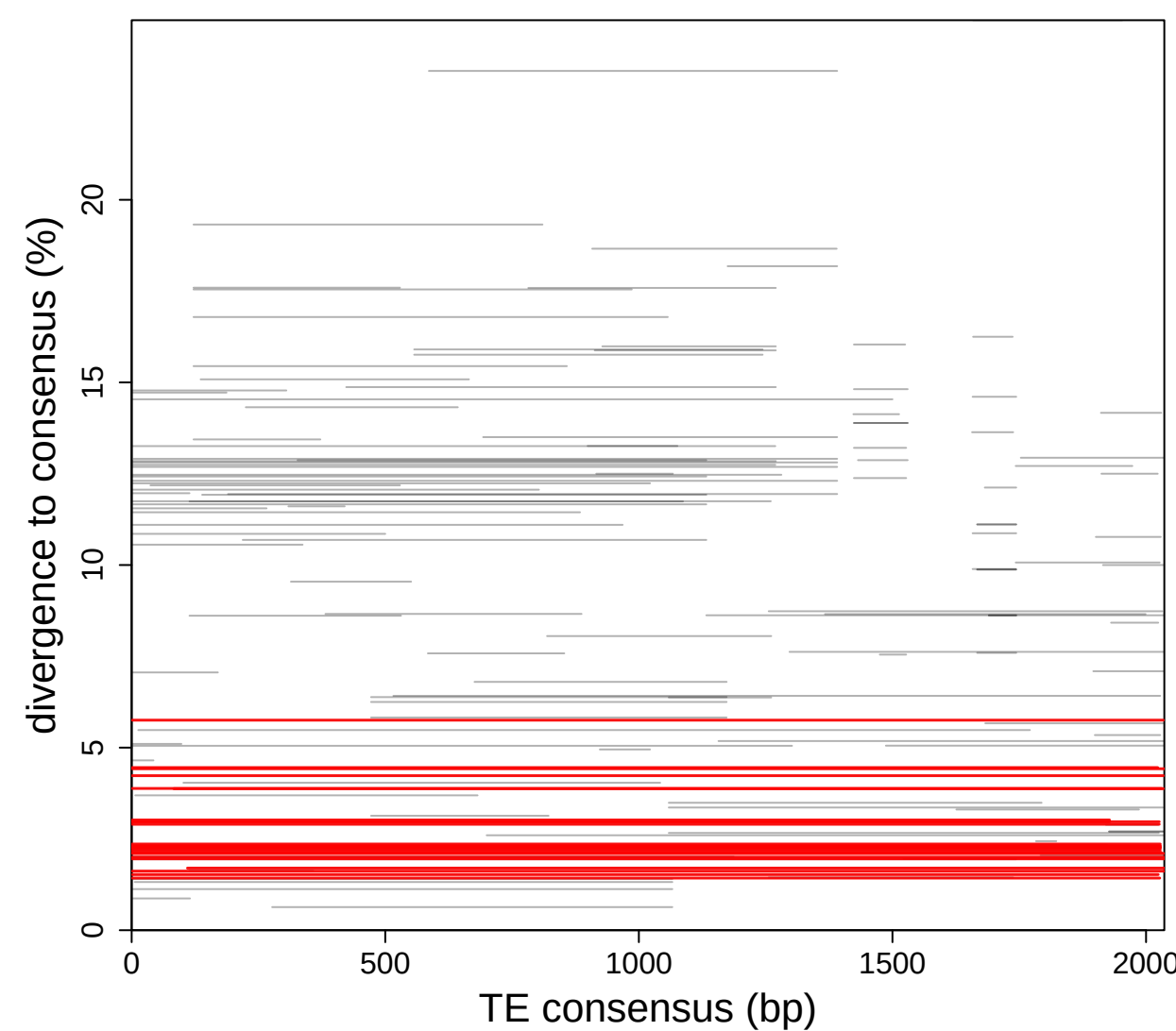
0.fasta.b.bed uf.bed g_2.bed fm_1.bed 0_0_bcIn.fa aln.fa cl.f
size: 2311bp; fragments: 208; full length: 0 (>=2079.9bp)

After TEtrimmer Extended plot Blue lines are boundaries

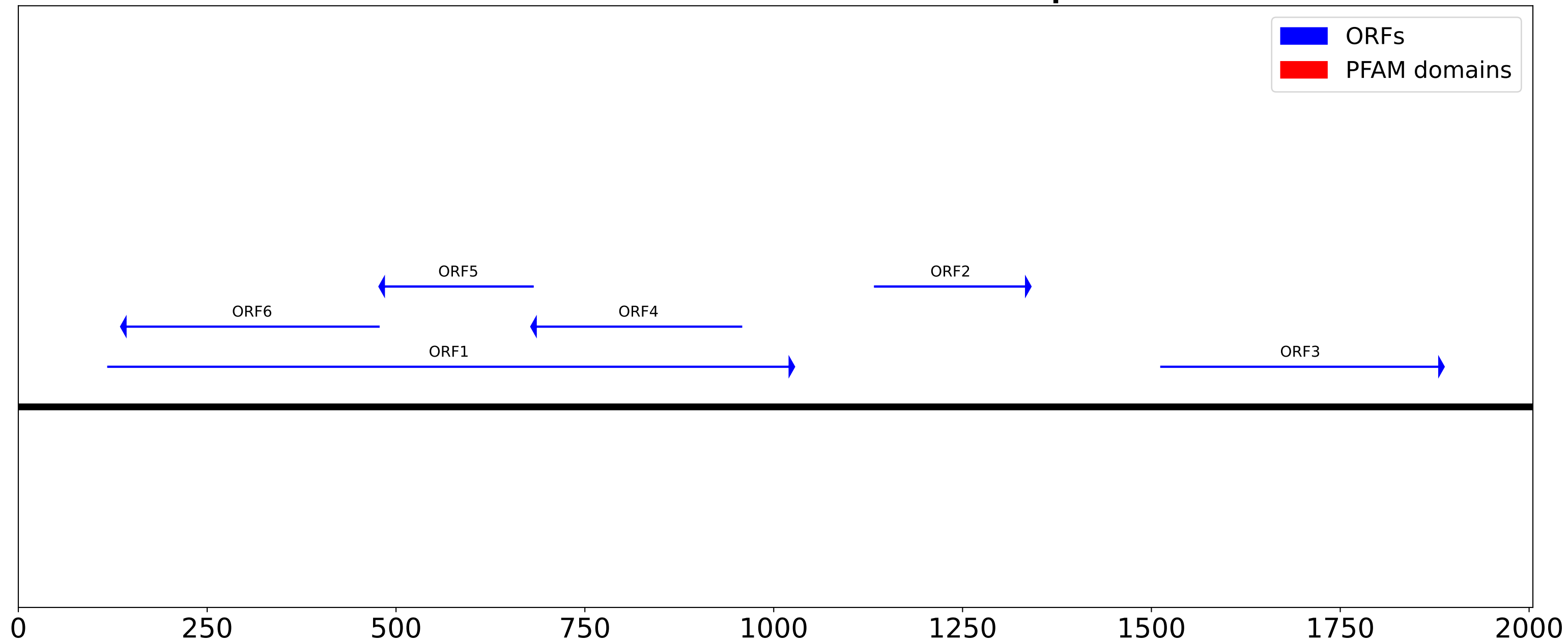


TE: rnd_5_family_4690
size: 2036bp; fragments: 158; full length: 26 (≥ 1832.4 bp)

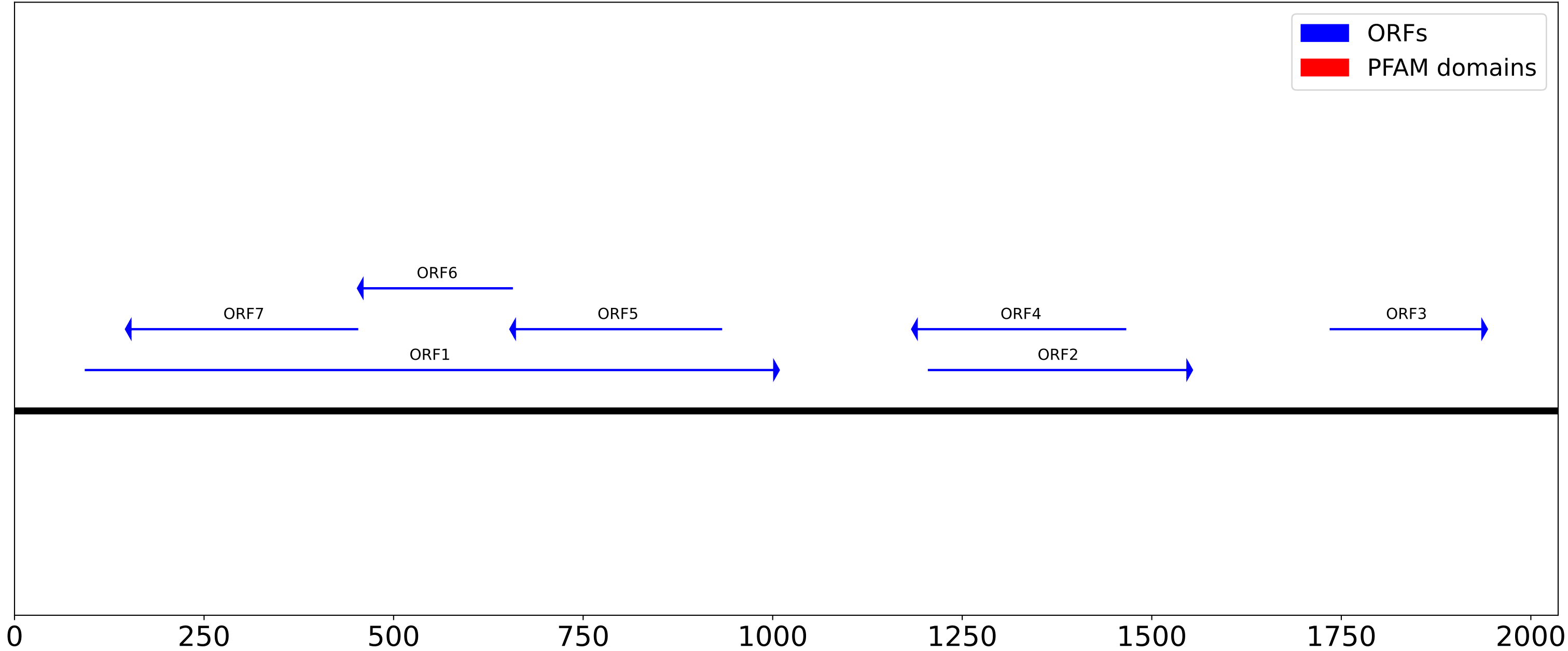
Before TEtrimmer 2036 bp



After TEtrimmer ORF and PFAM domain plot



Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

