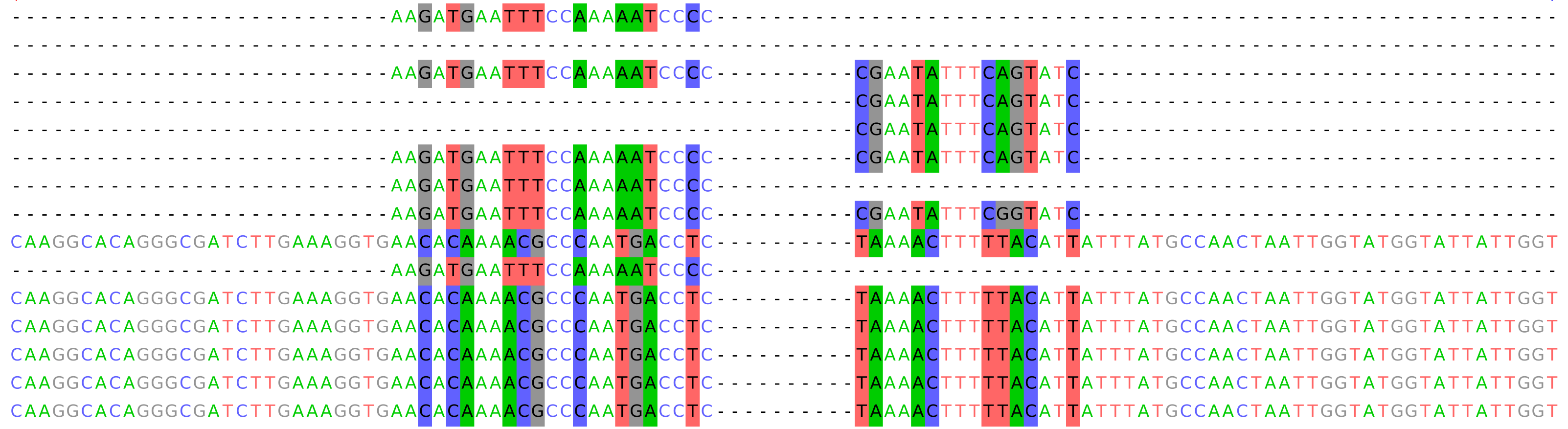
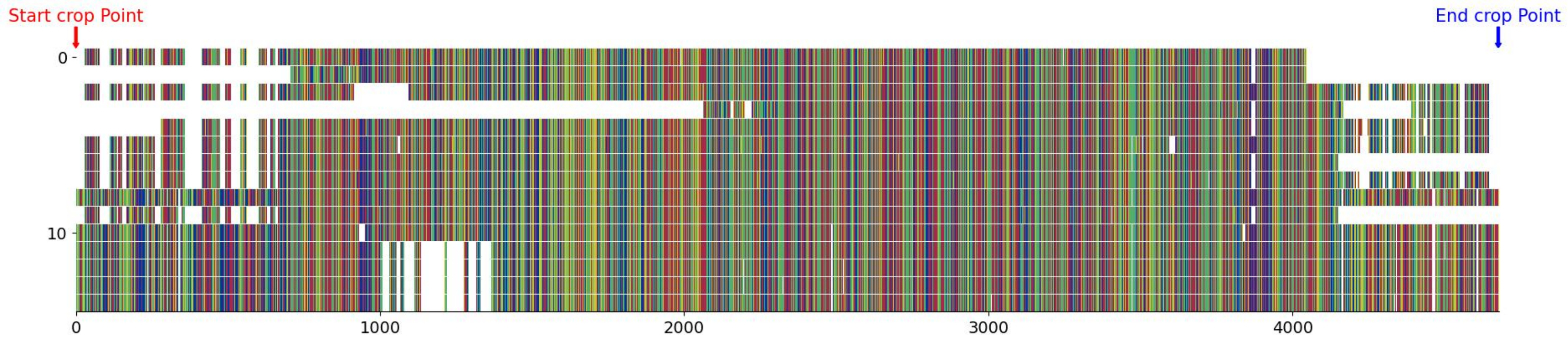


Start crop Point

End crop Point

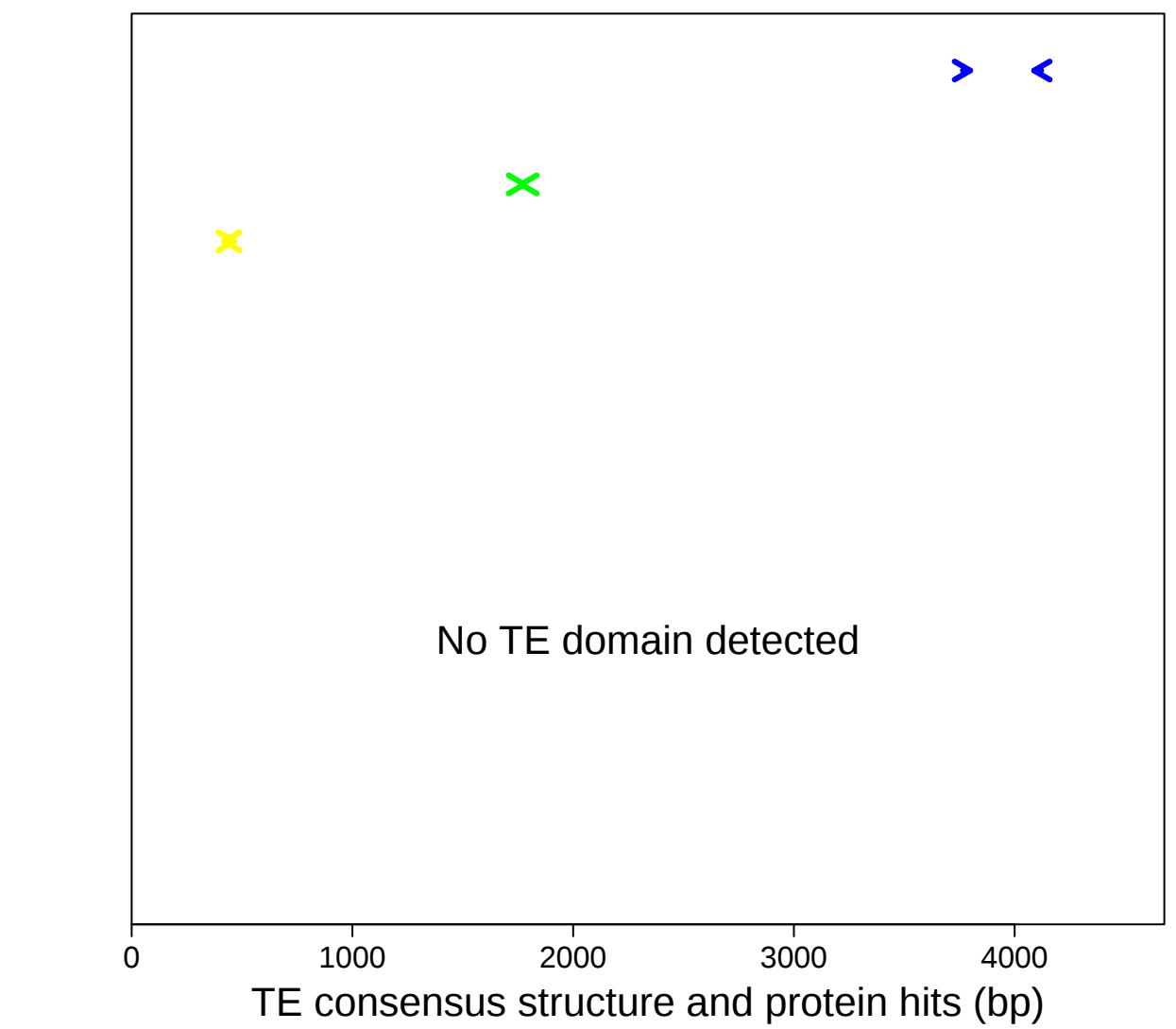
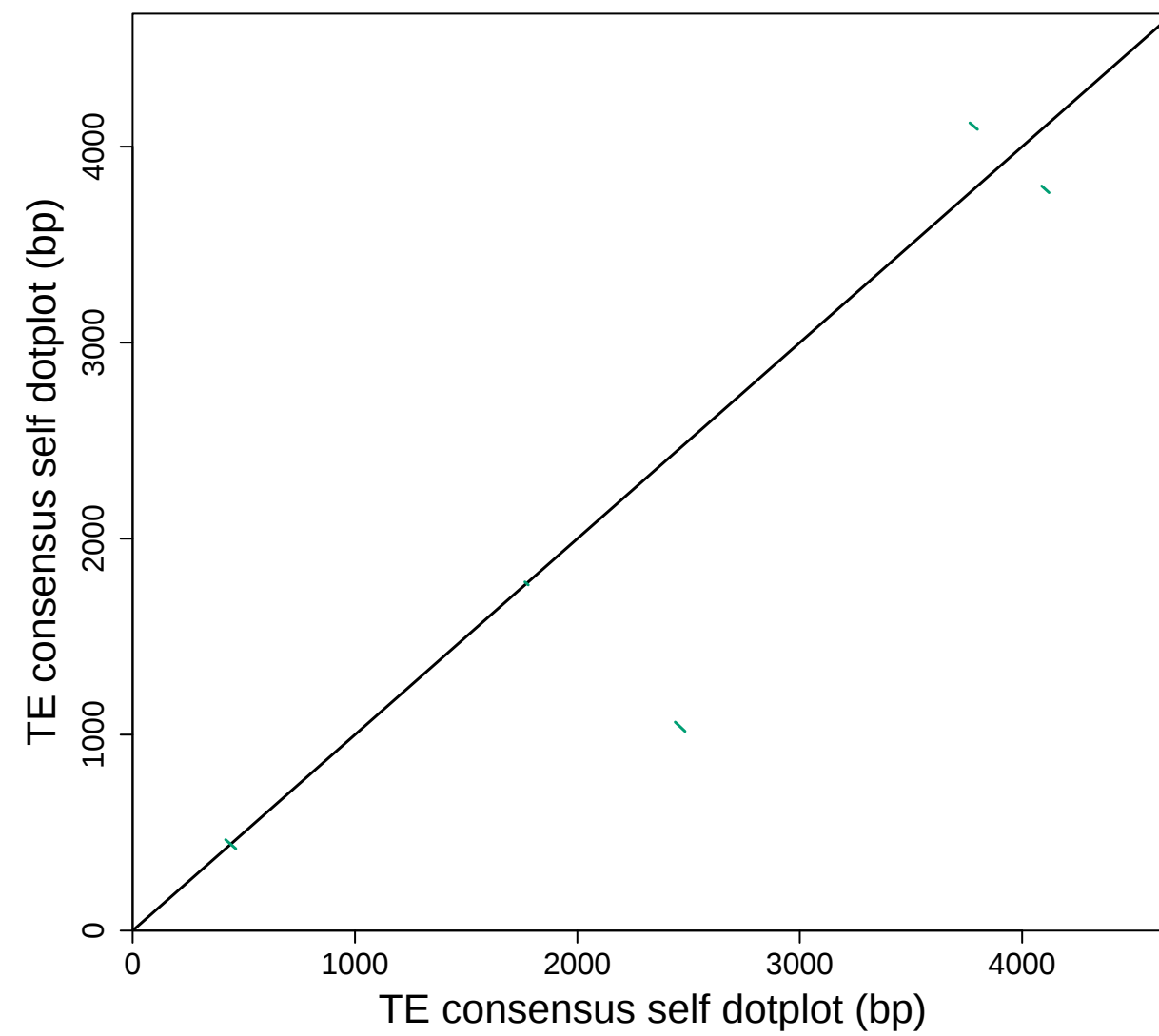
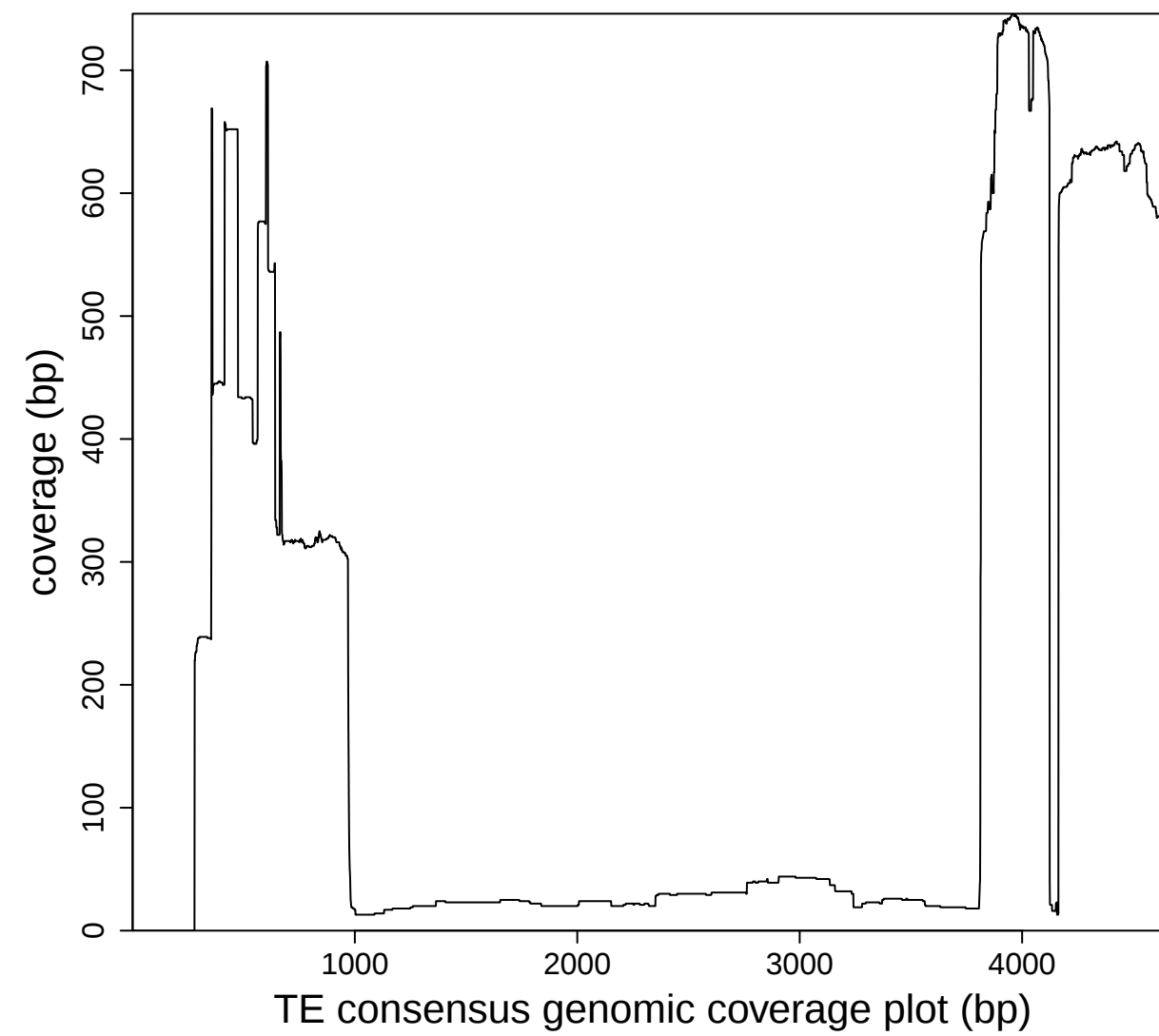
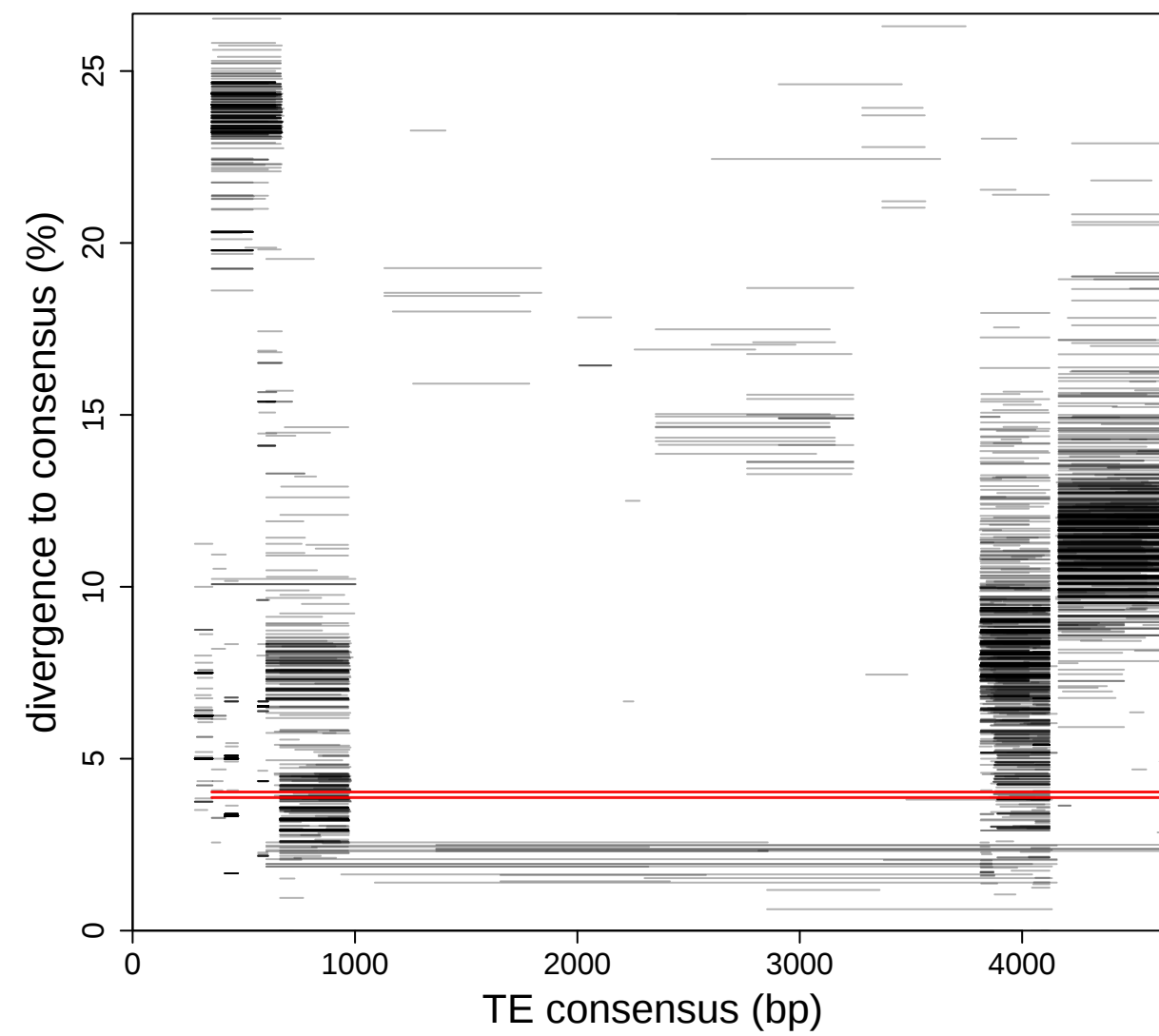
MSA length = 4678





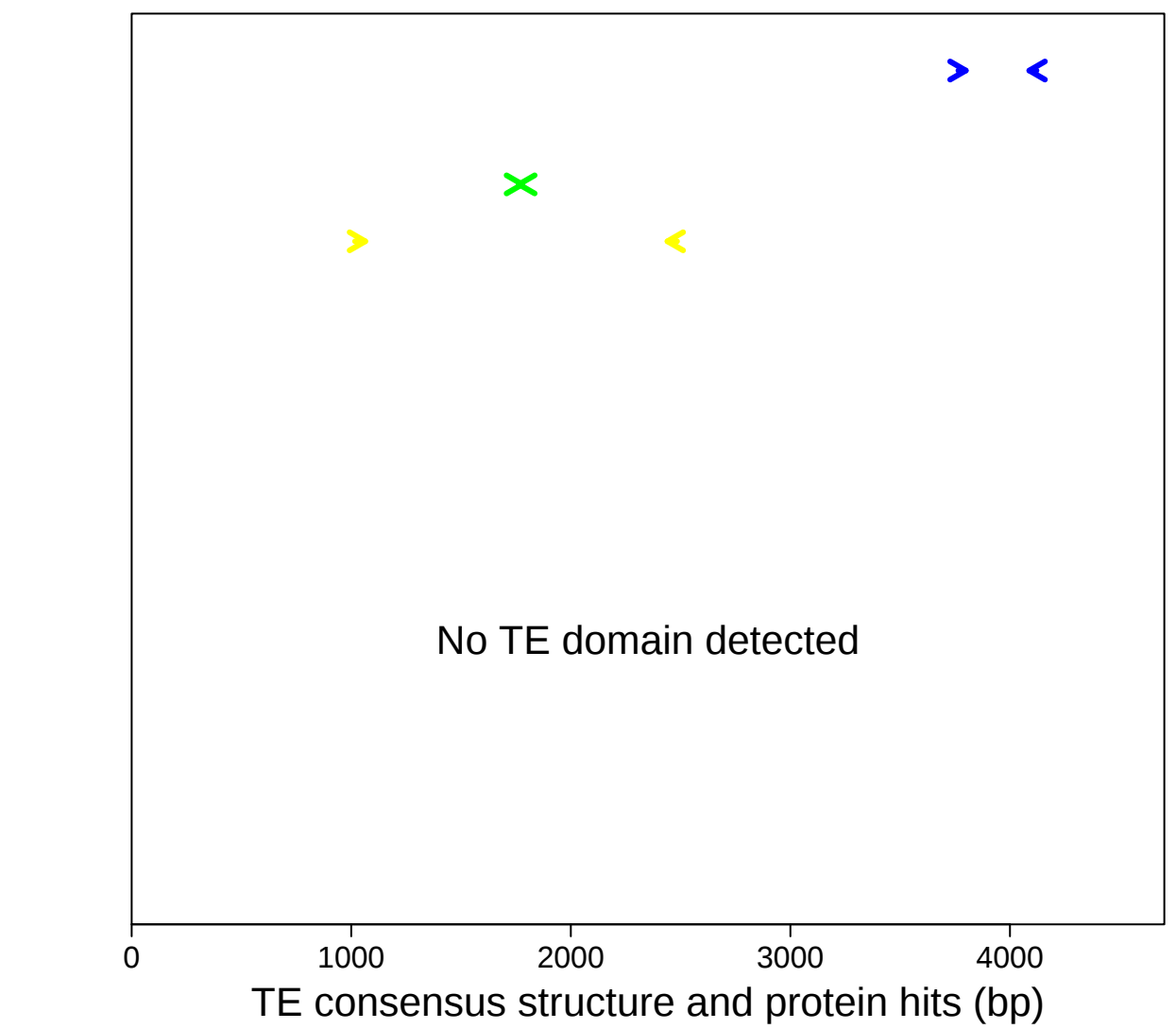
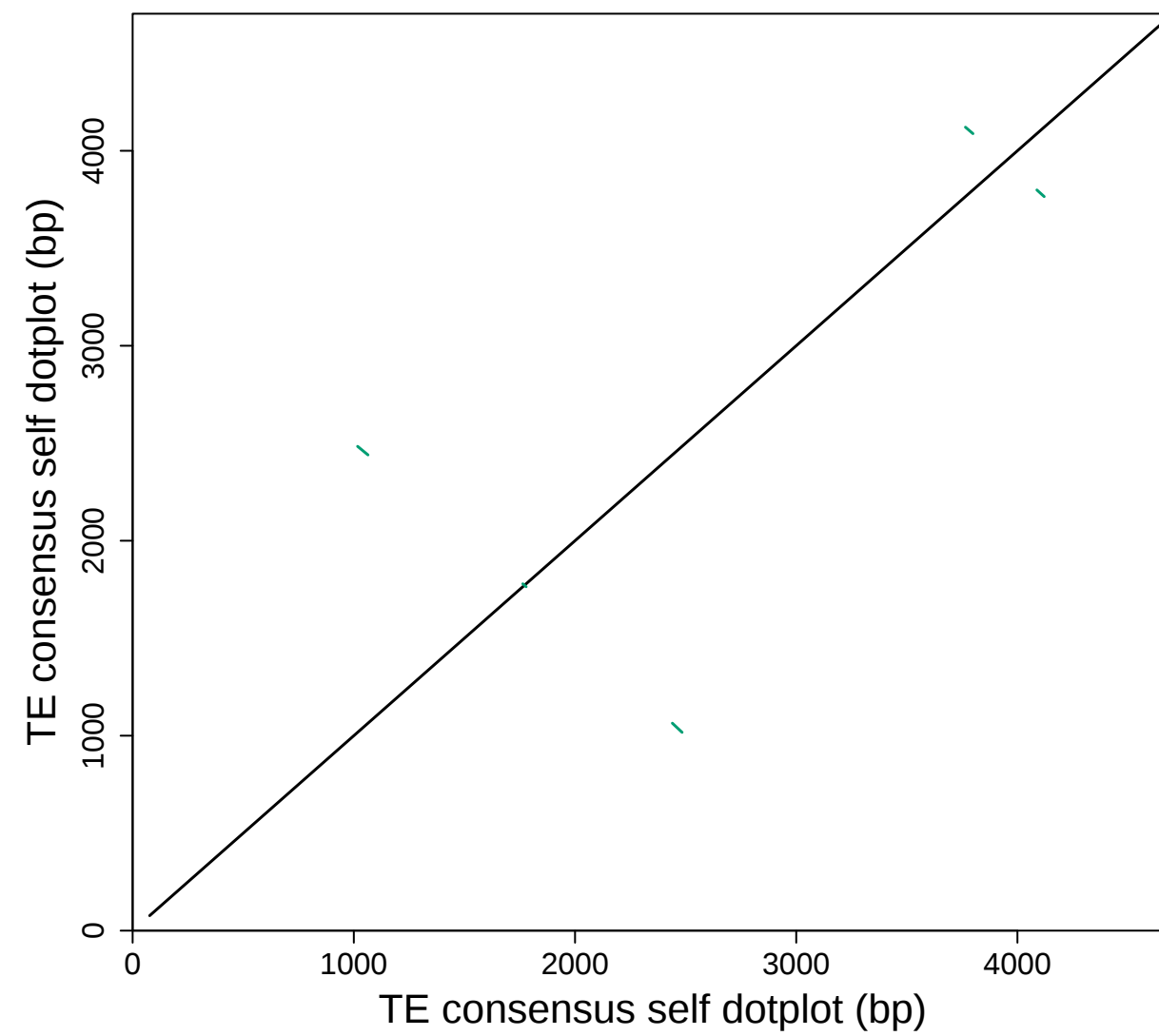
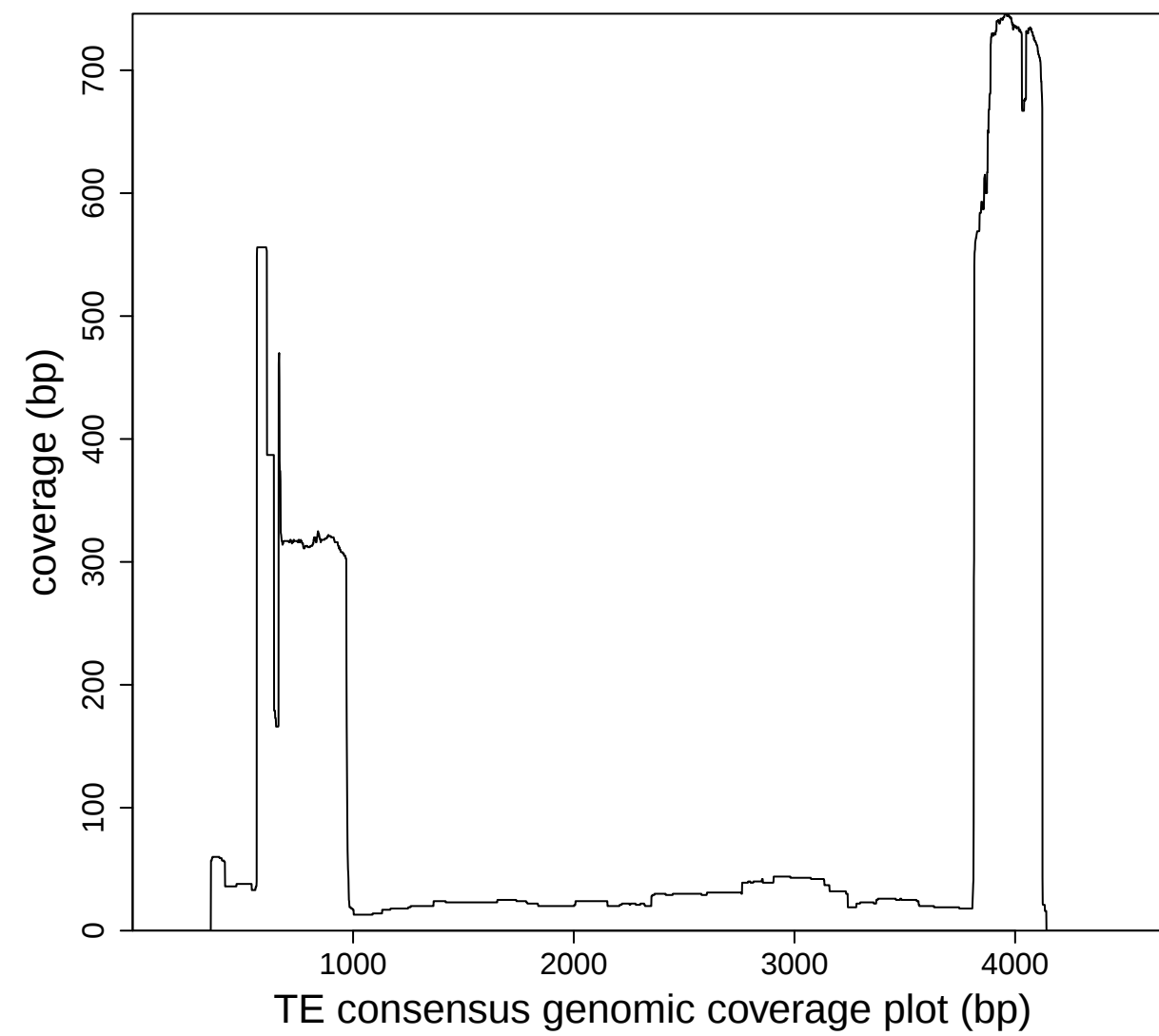
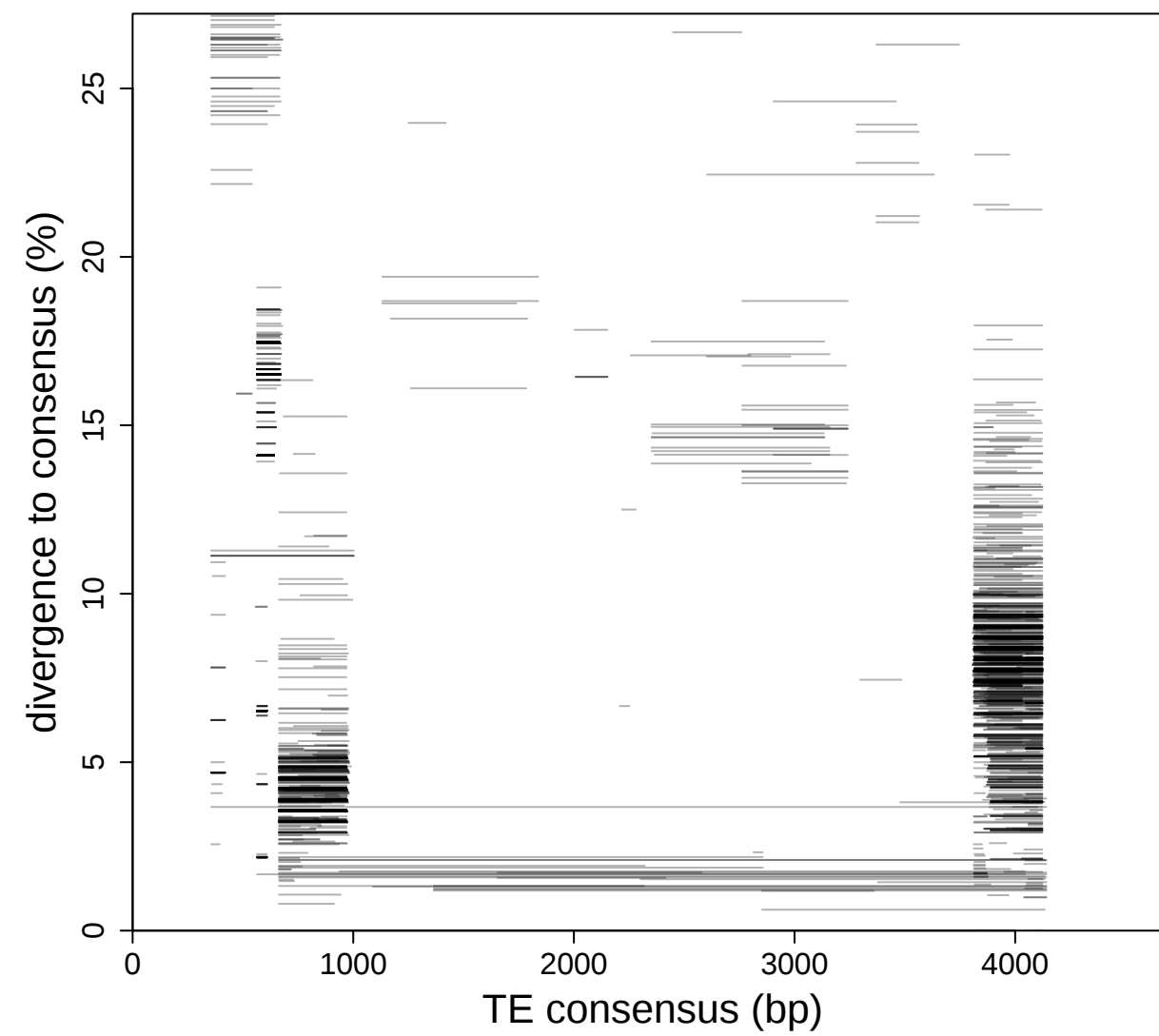
a.b.bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.f
size: 4678bp; fragments: 3078; full length: 2 (>=4210.2bp)

After TEtrimmer 4678 bp



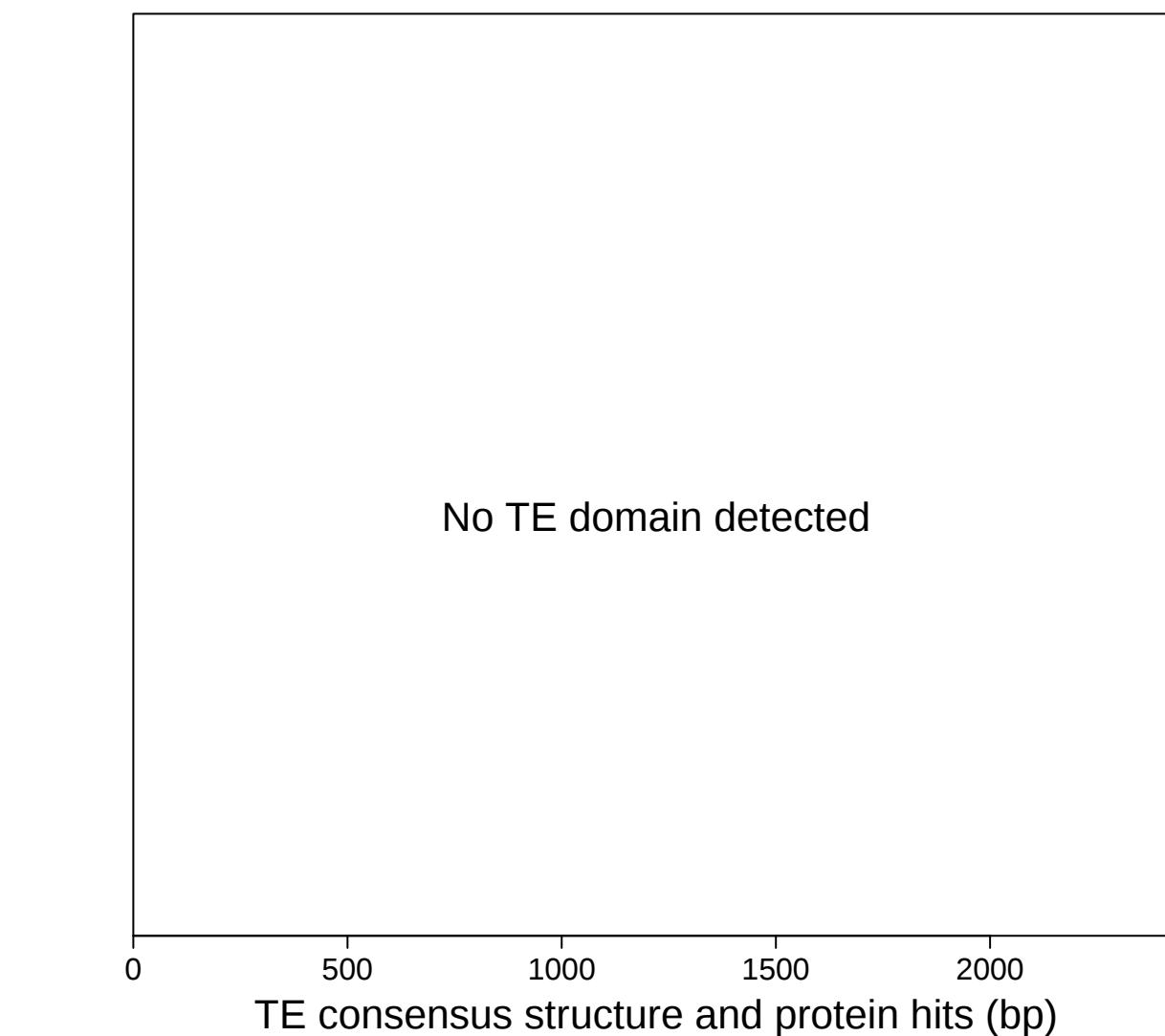
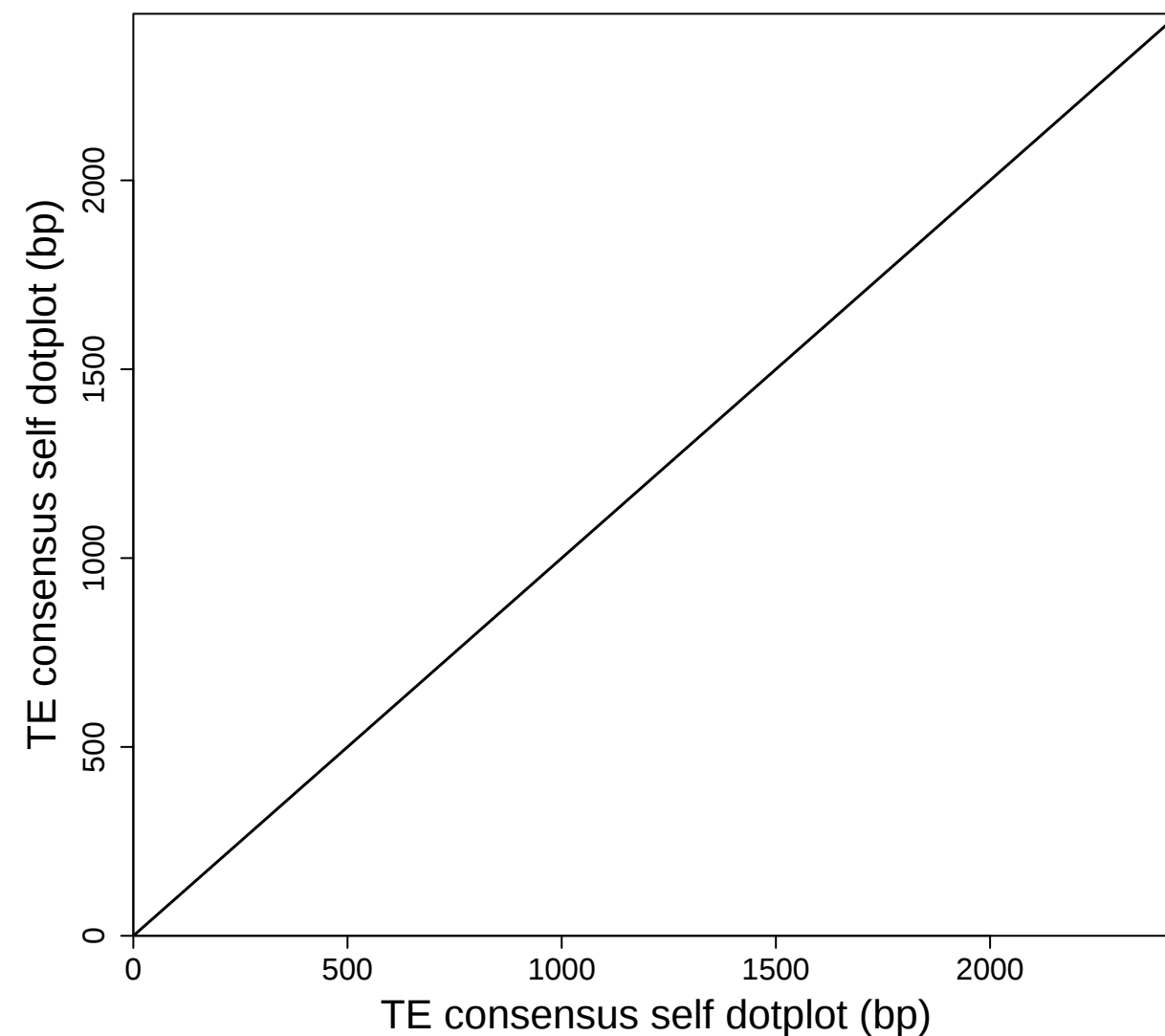
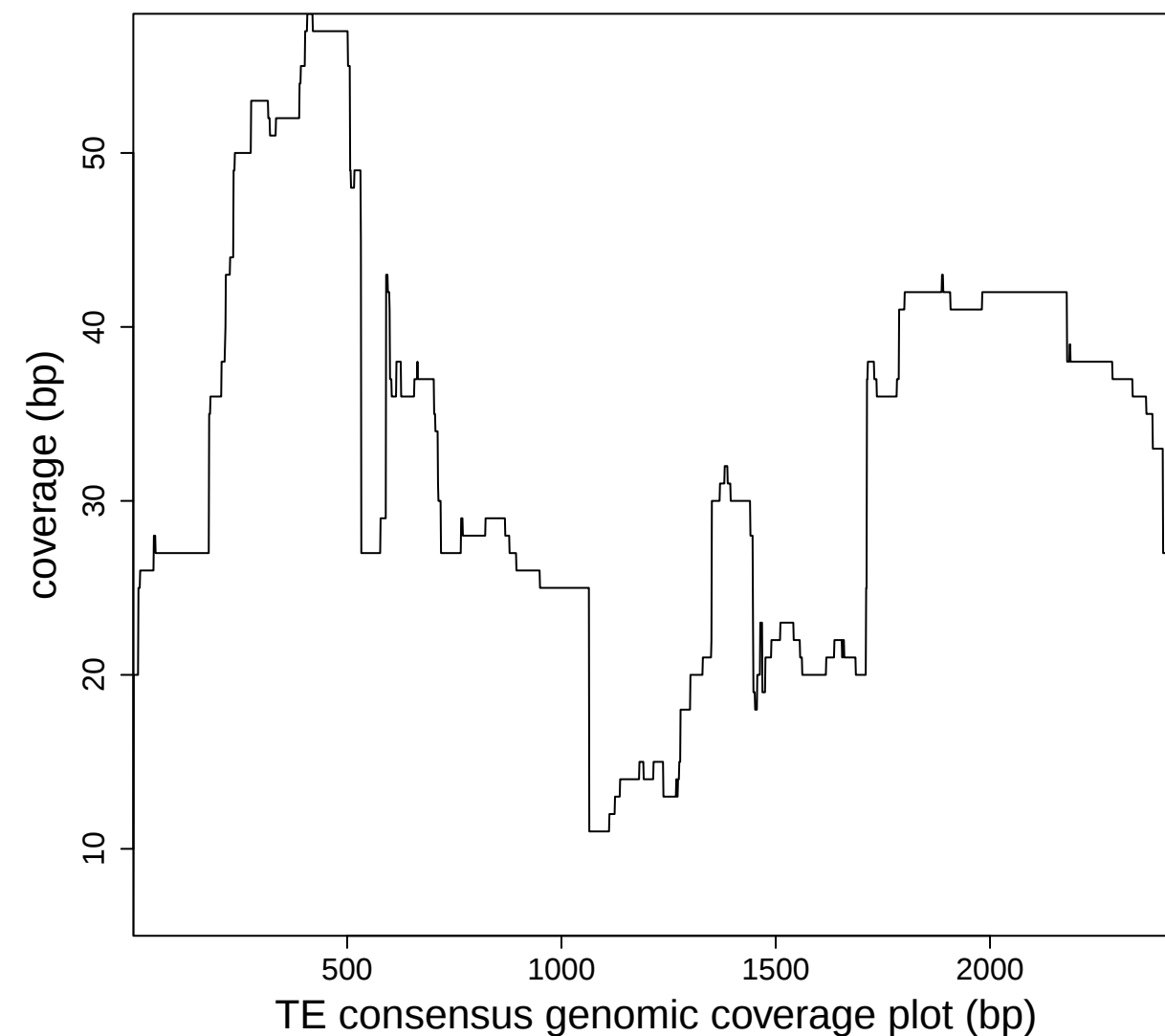
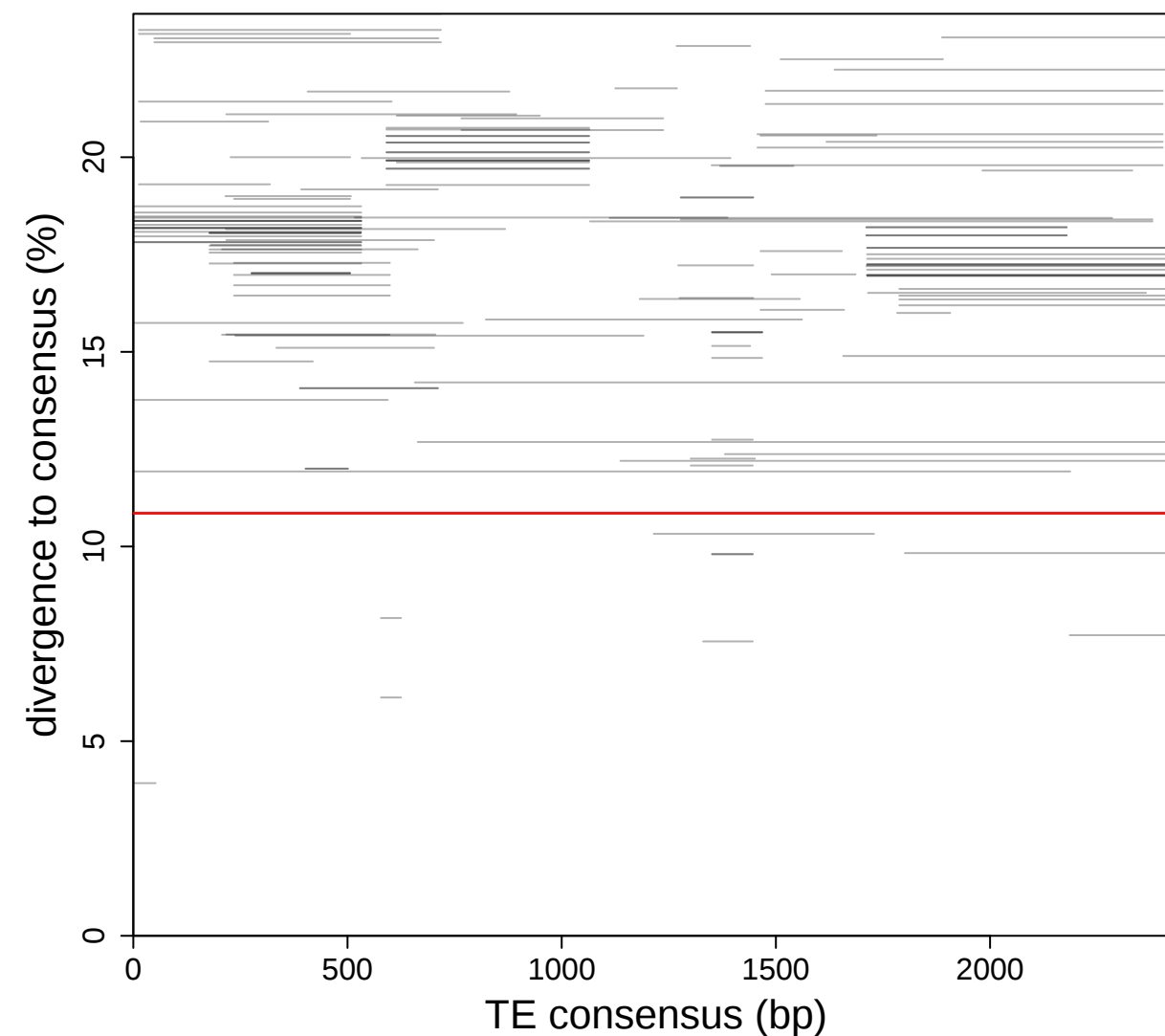
asta.b.bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_c
size: 4715bp; fragments: 1885; full length: 0 (>=4243.5bp)

After TEtrimmer Extended plot Blue lines are boundaries

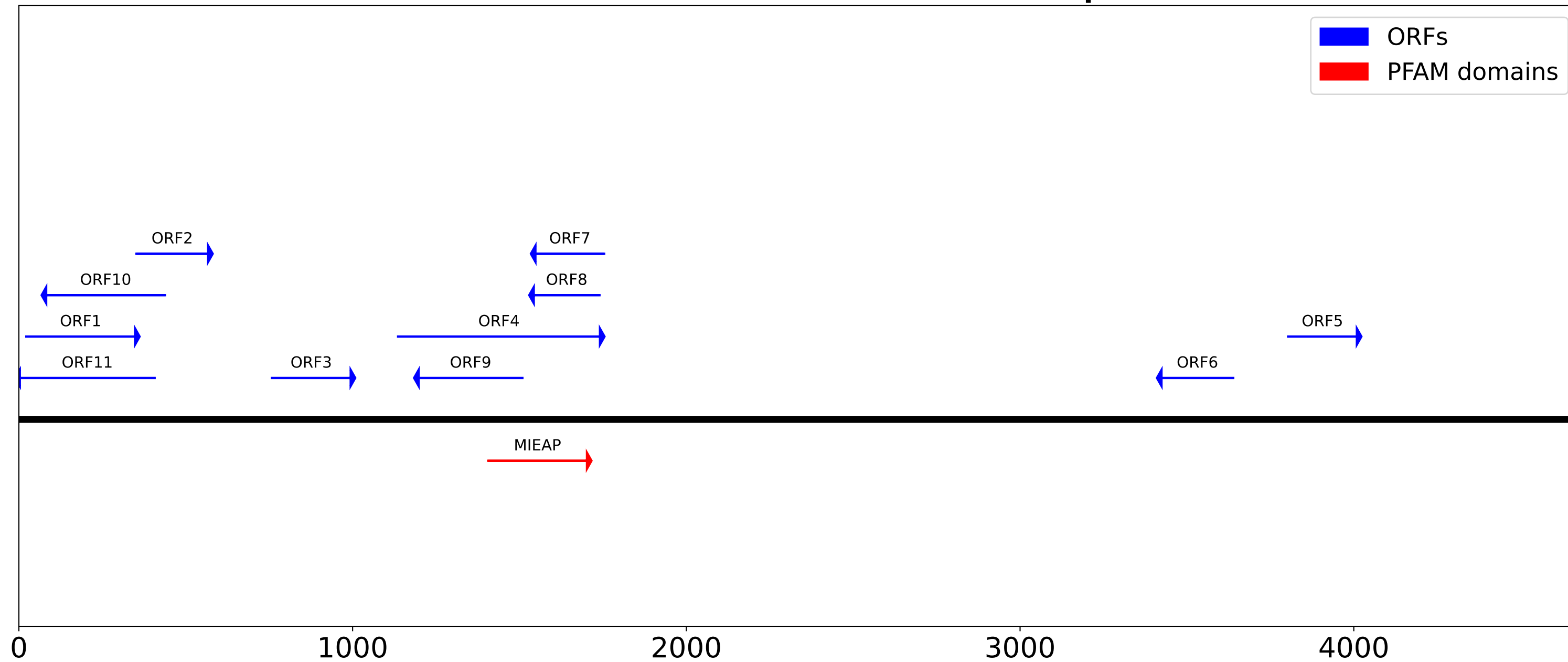


TE: rnd_5_family_9737
size: 2441bp; fragments: 151; full length: 1 (>=2196.9bp)

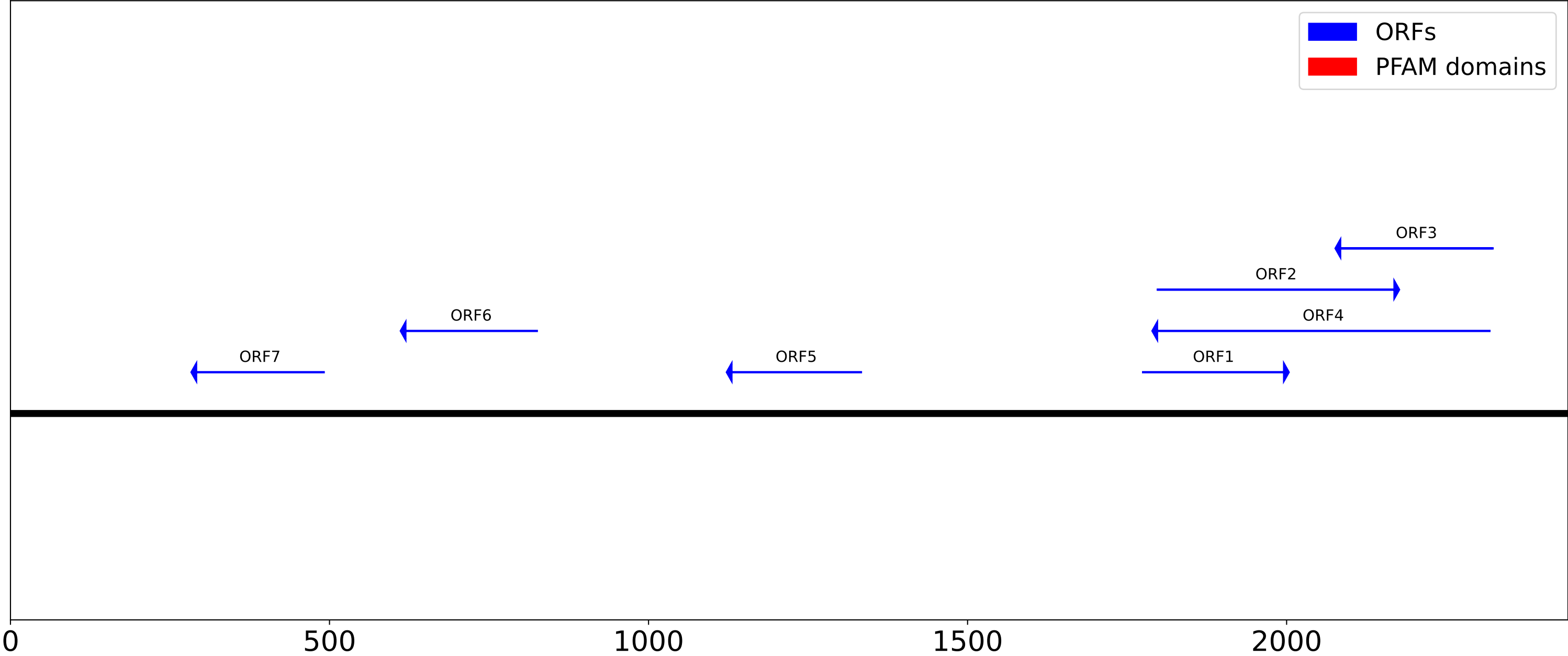
Before TEtrimmer 2441 bp



After TEtrimmer ORF and PFAM domain plot



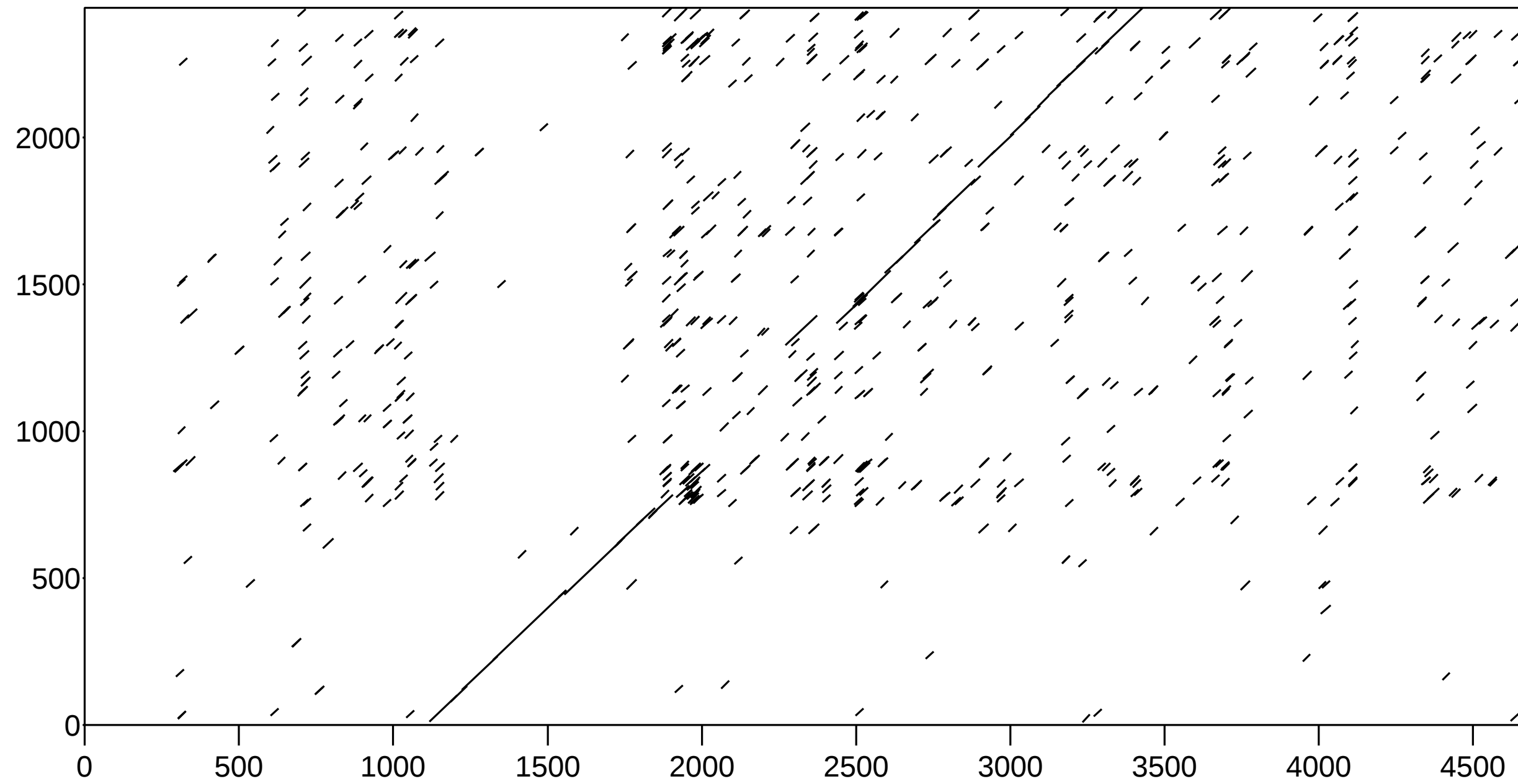
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)