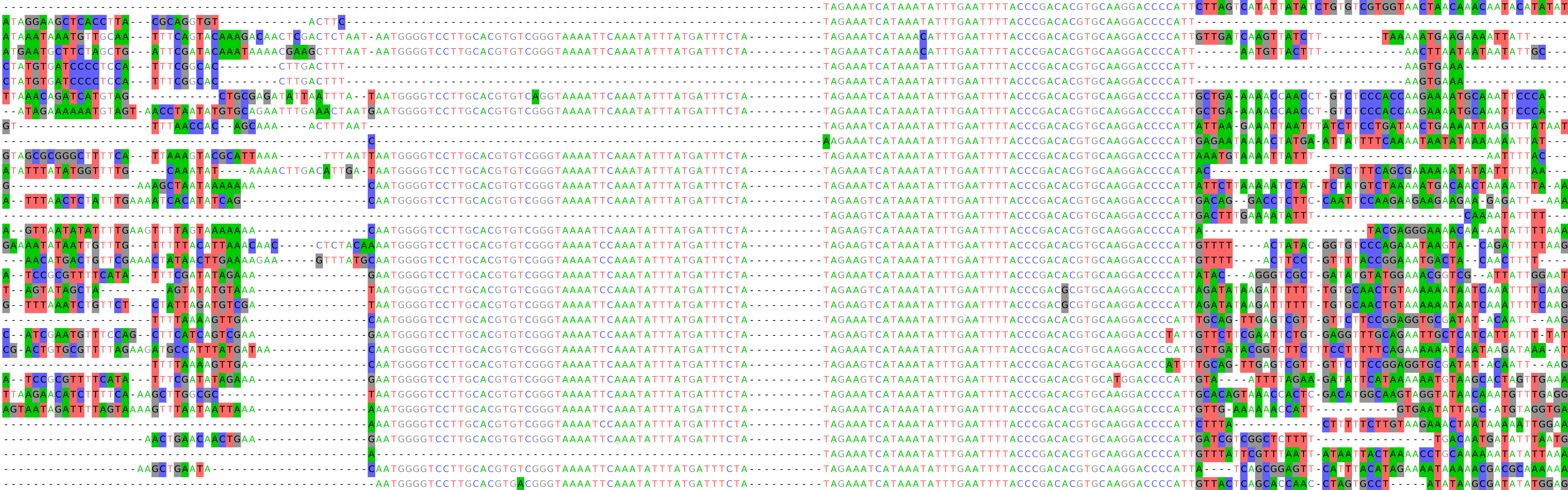


Start crop Point



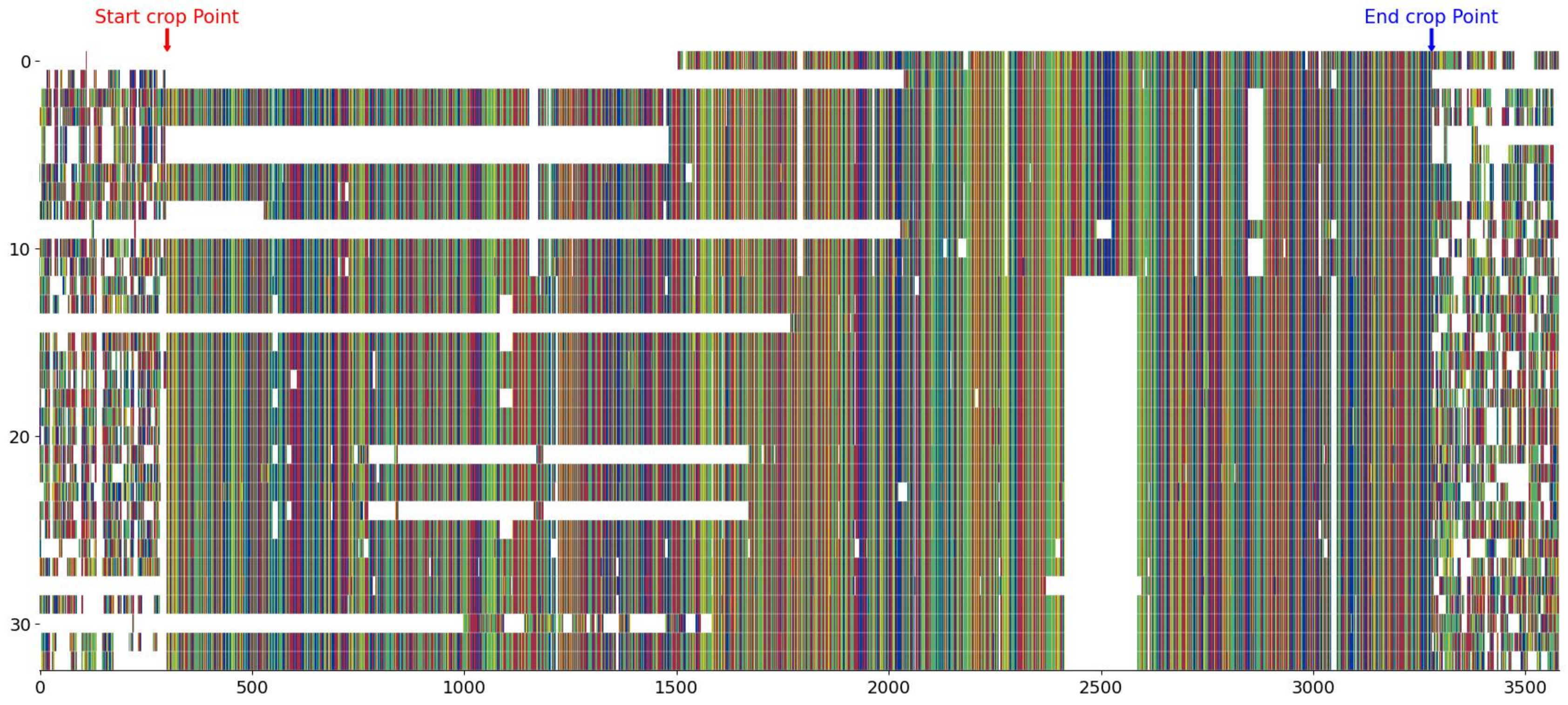
MSA length = 2980

End crop Point



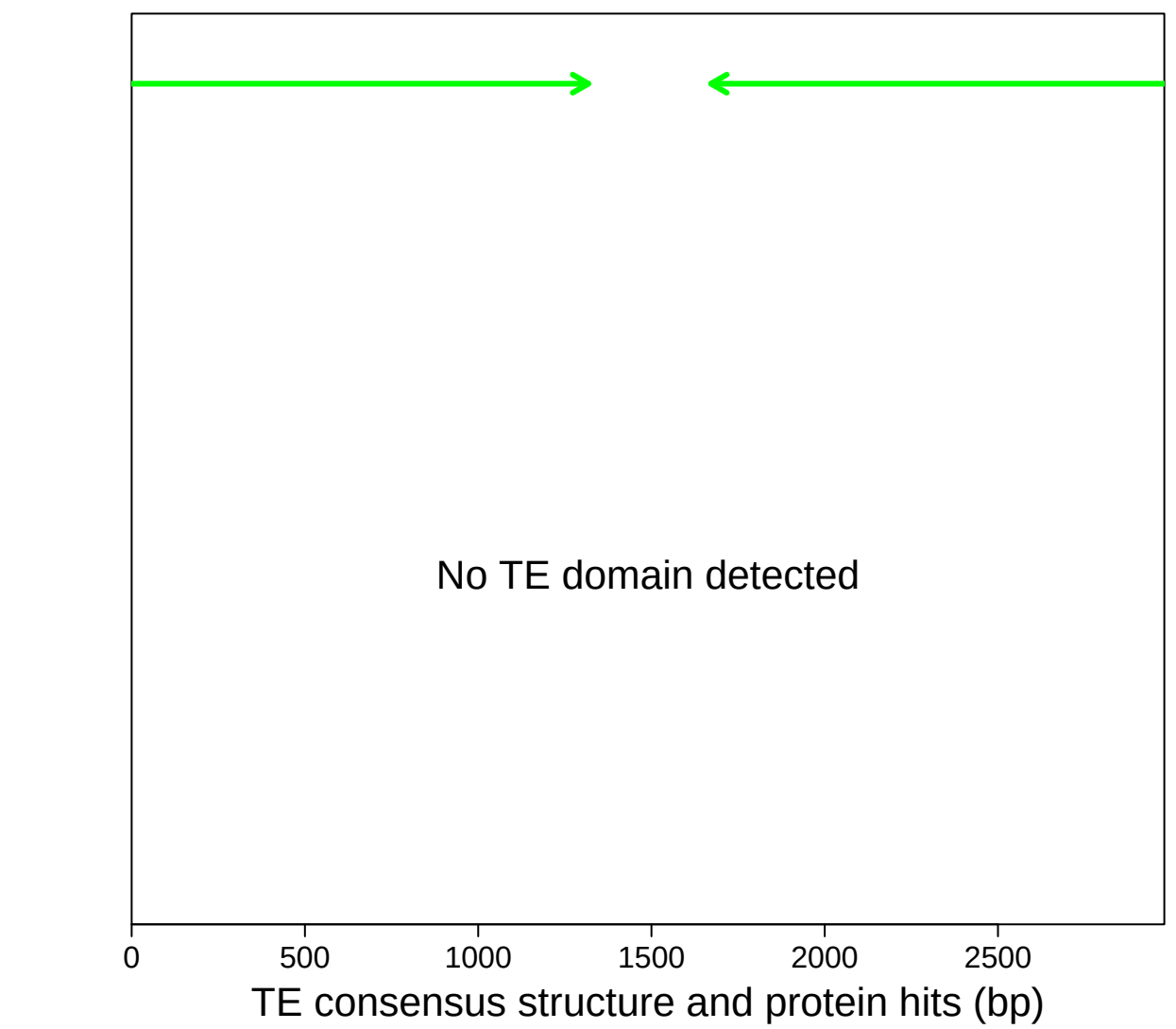
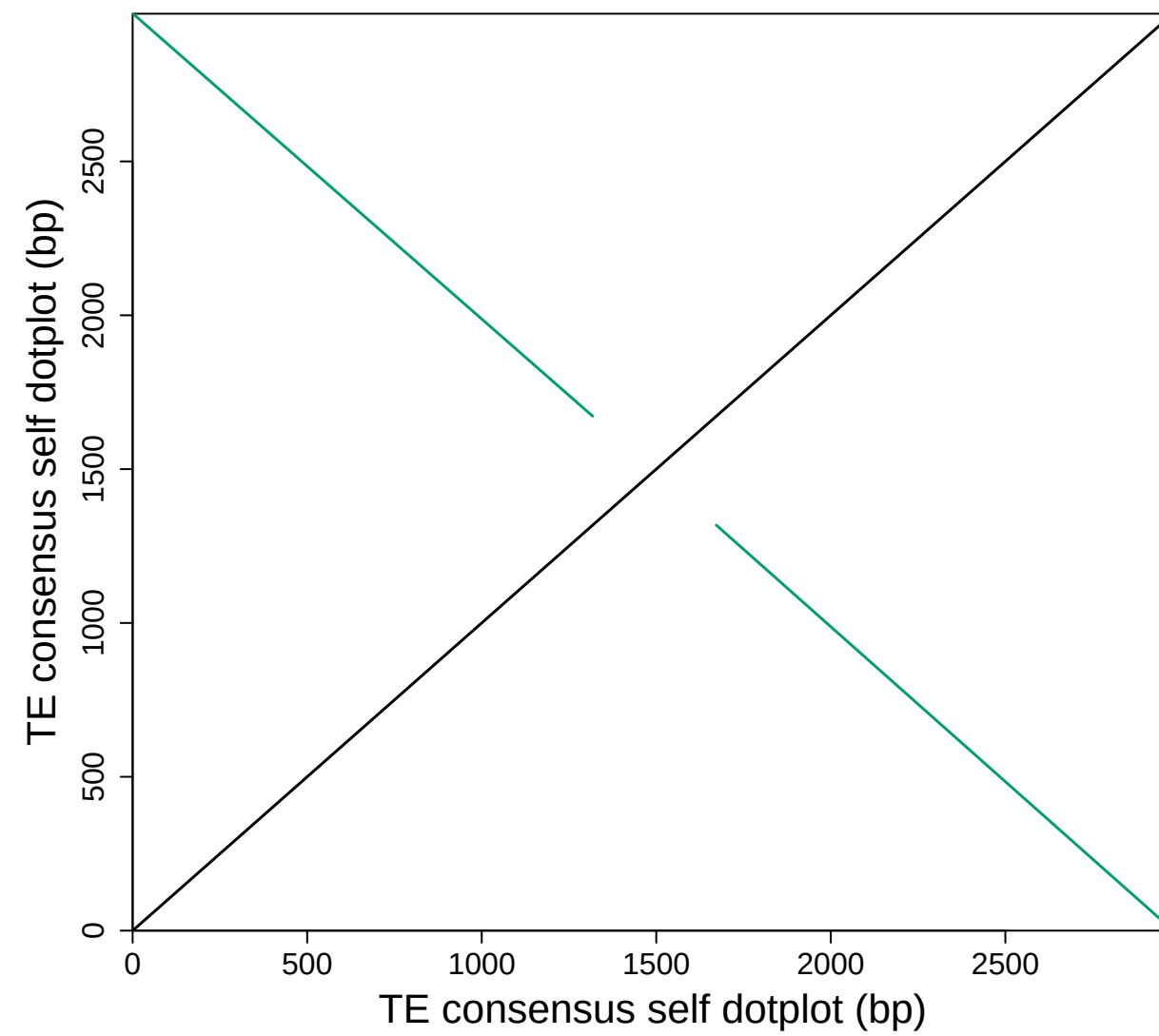
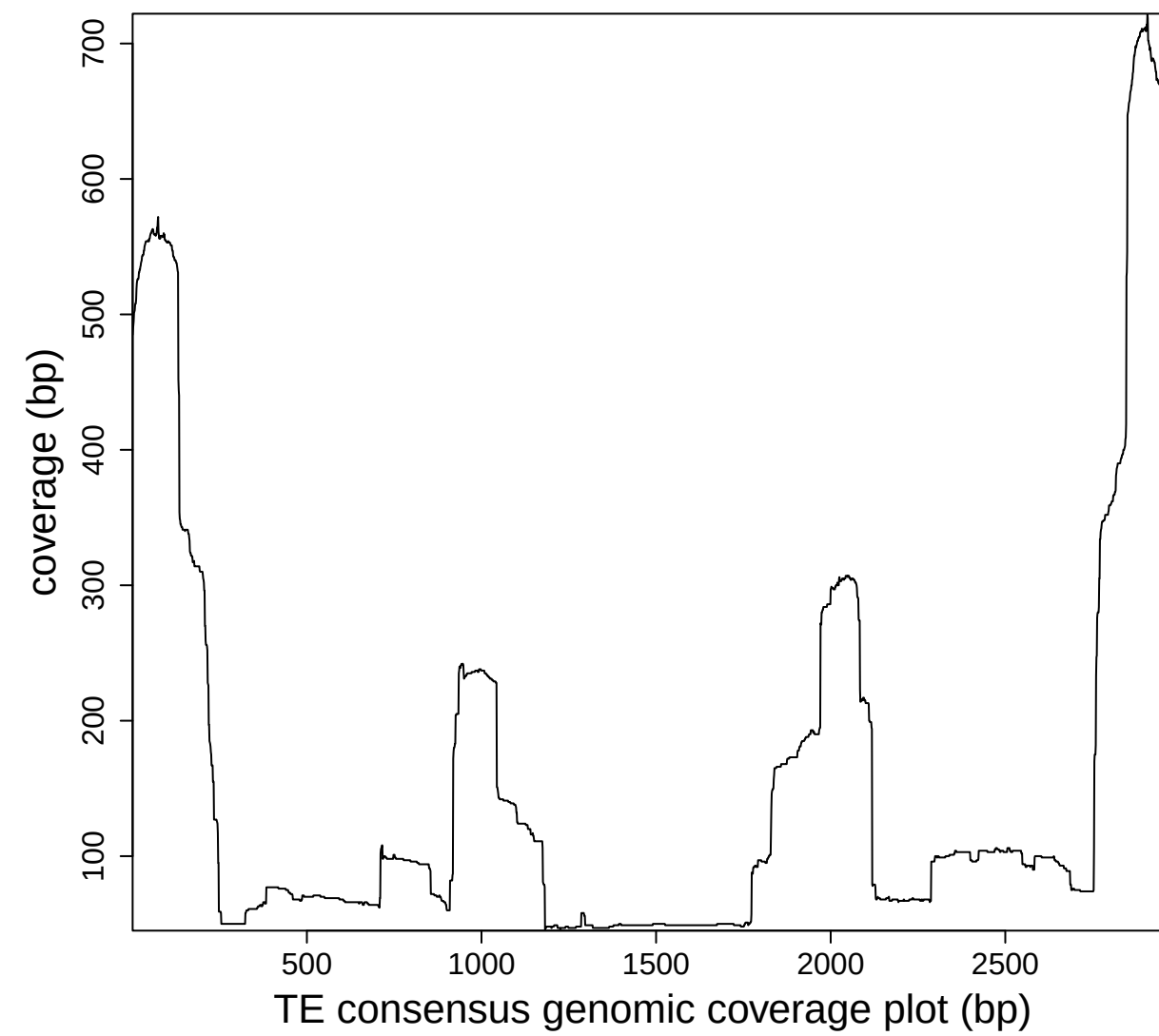
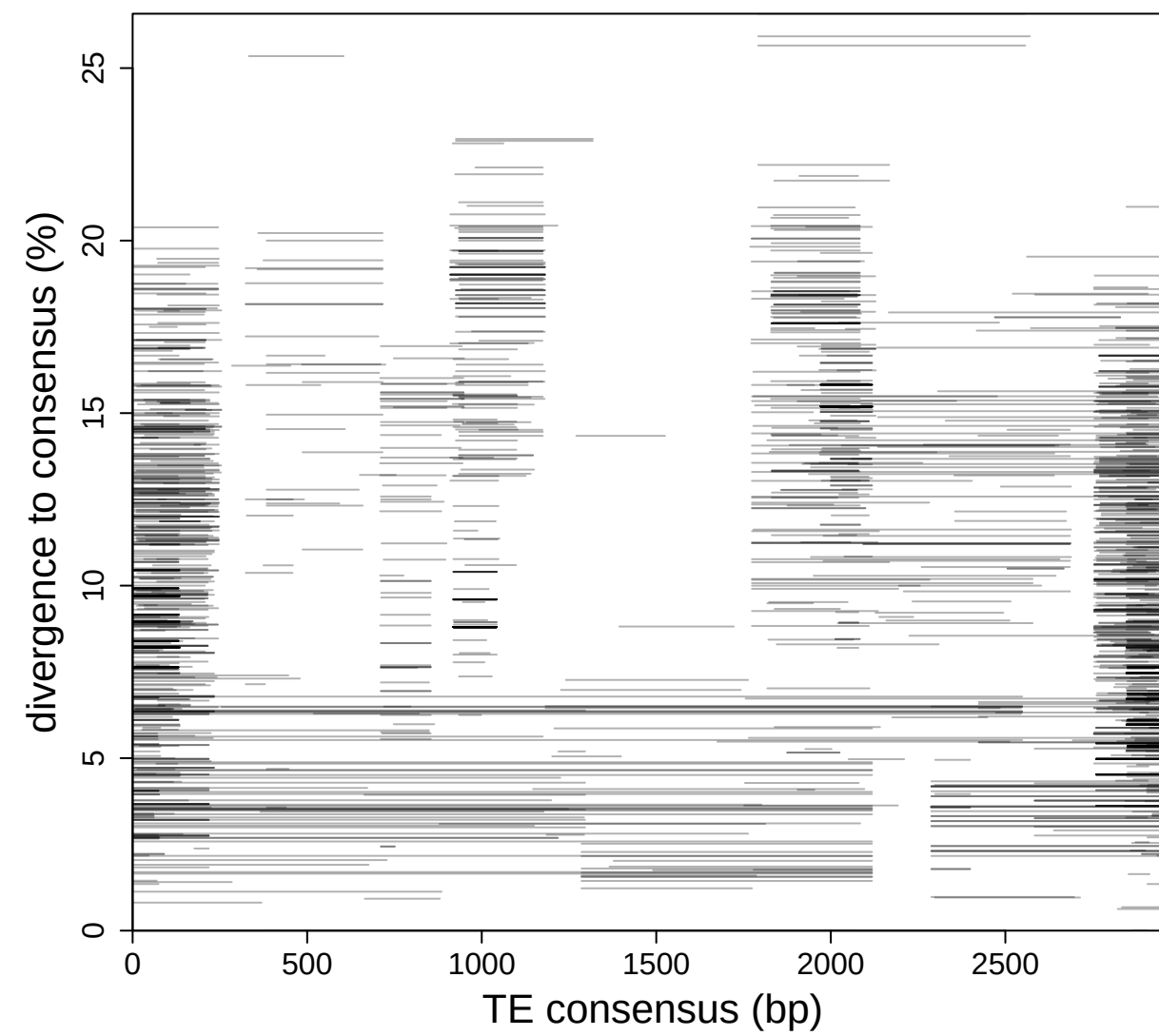
Start crop Point

End crop Point



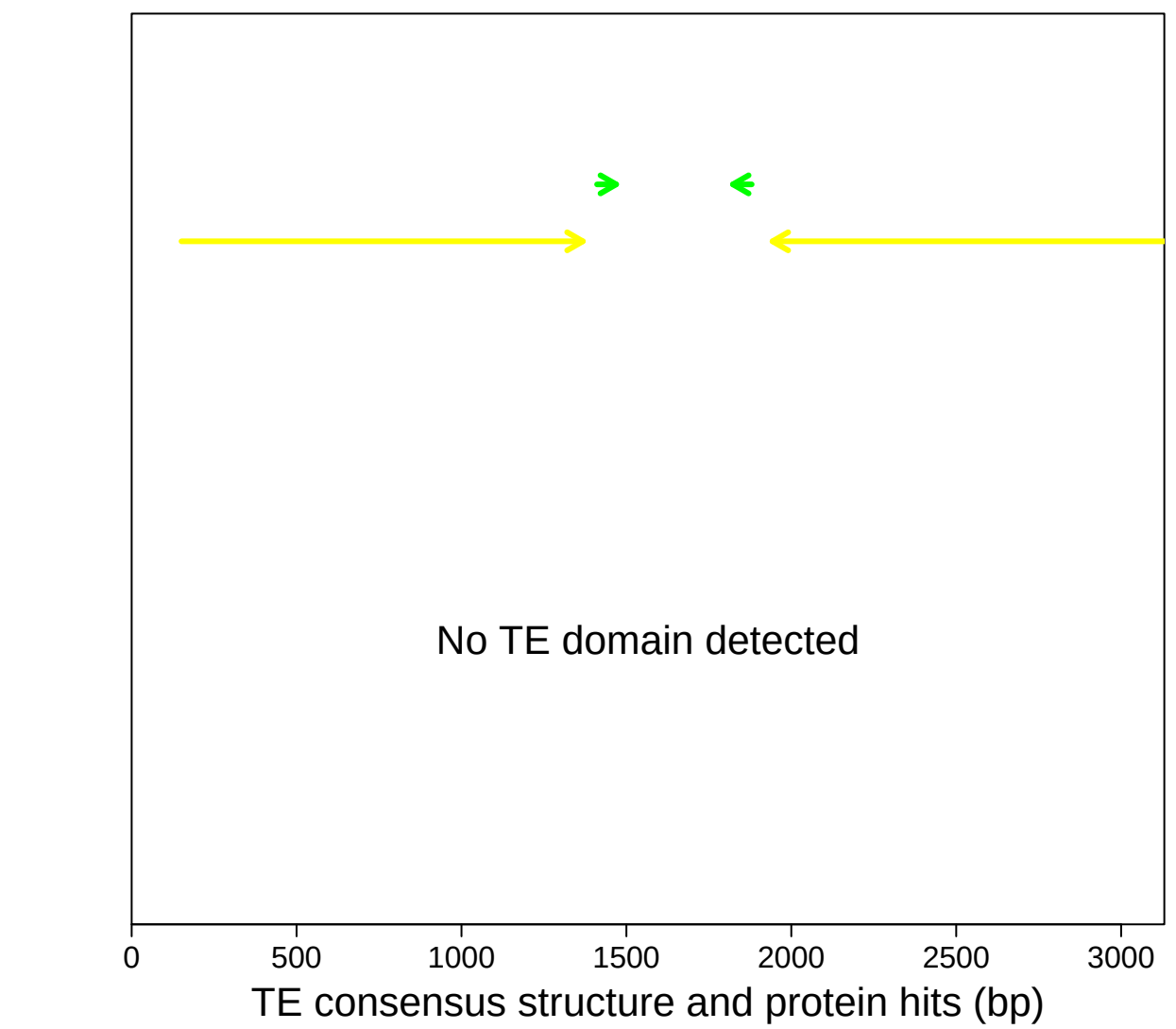
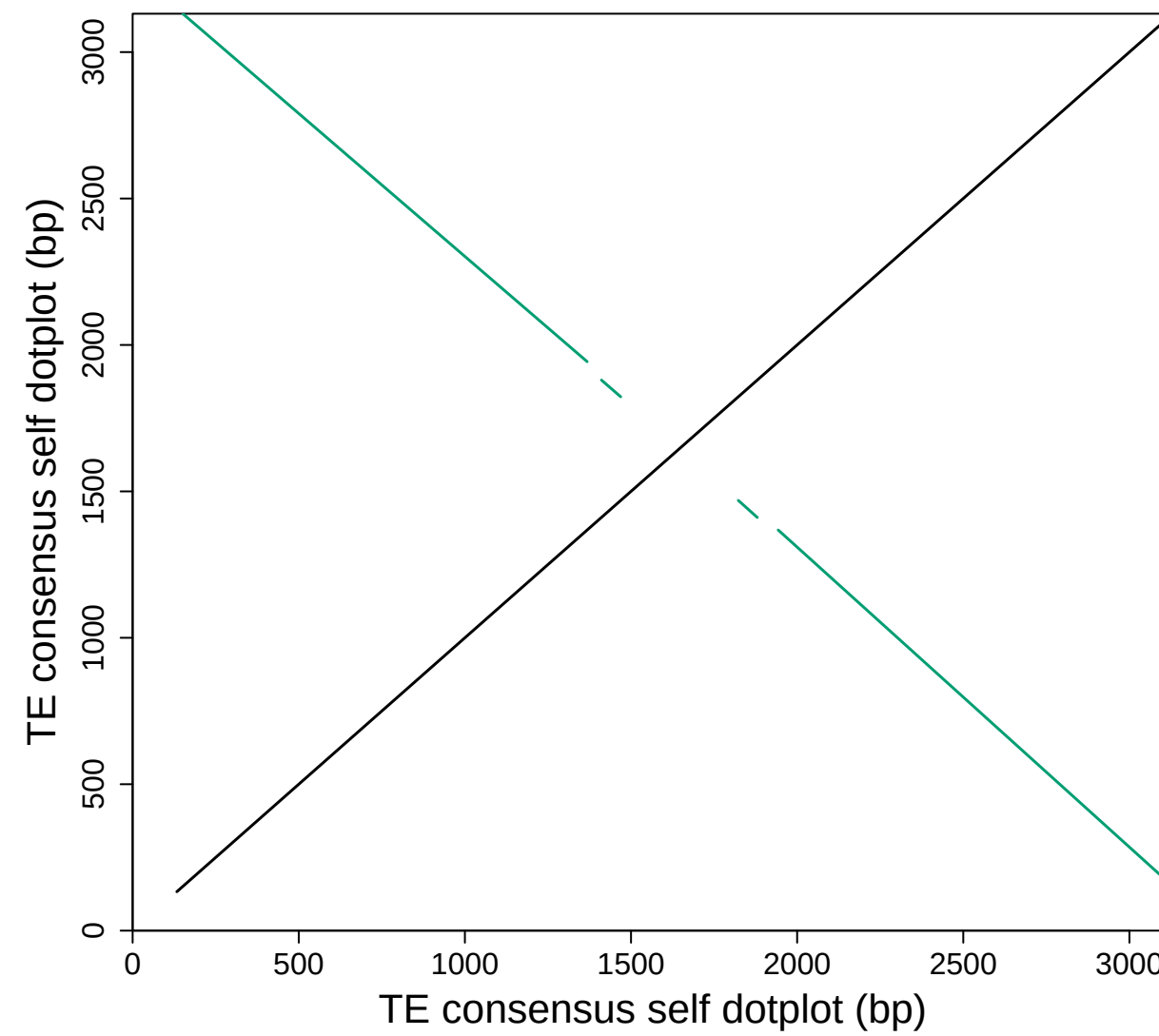
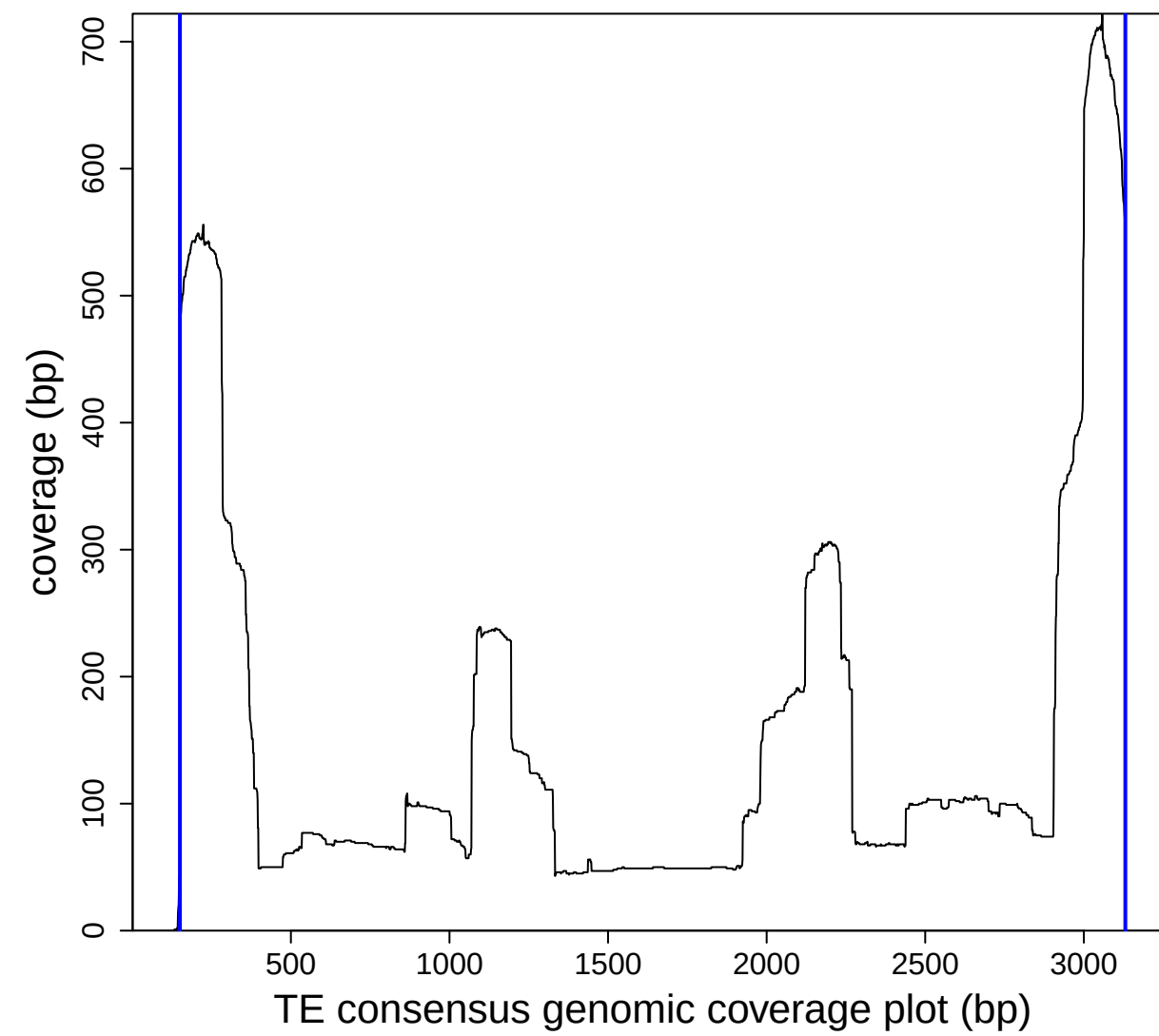
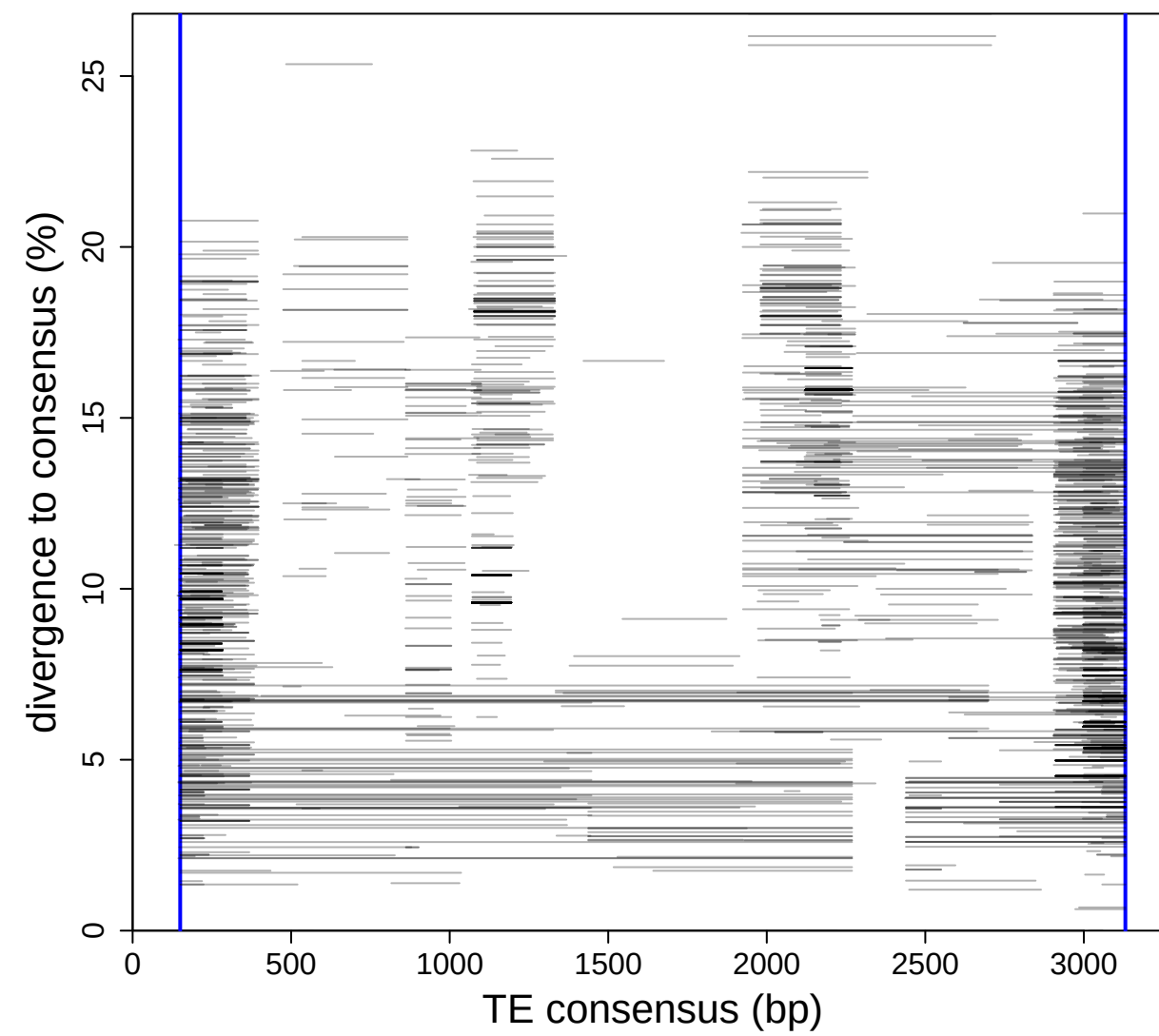
bed_uf.bed_g_2.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.fa_c
size: 2980bp; fragments: 1973; full length: 0 (>=2682bp)

After TEtrimmer 2980 bp



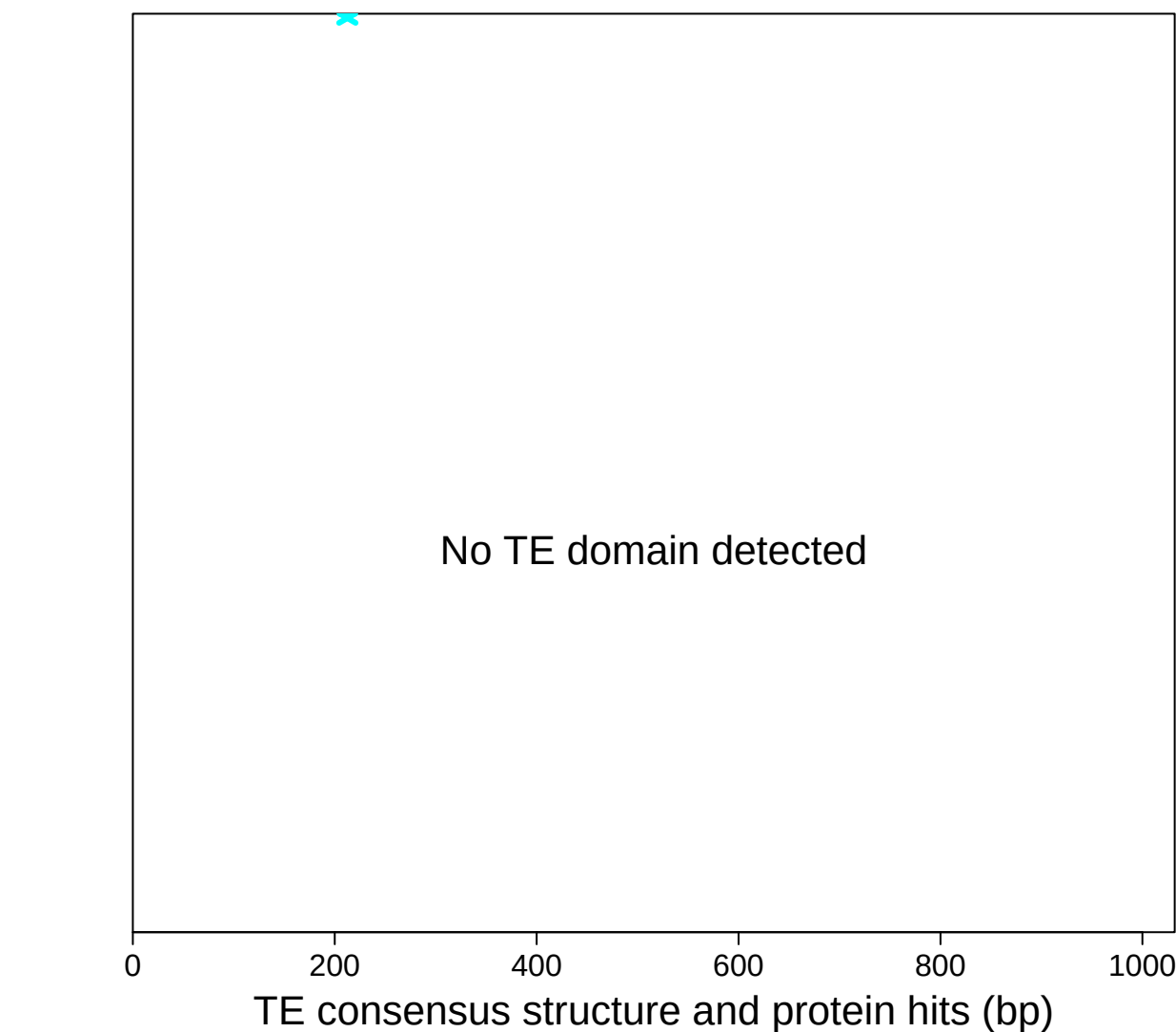
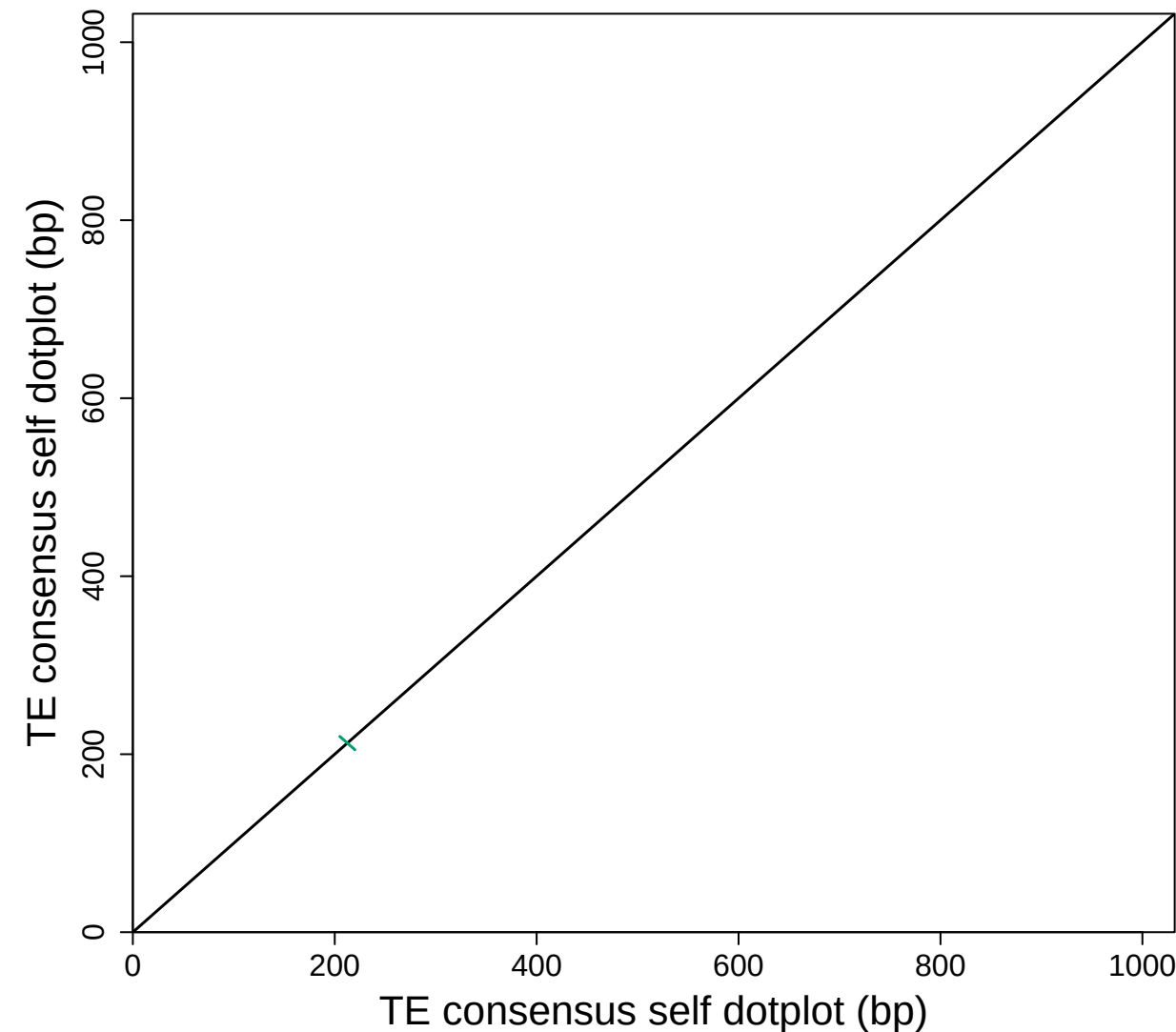
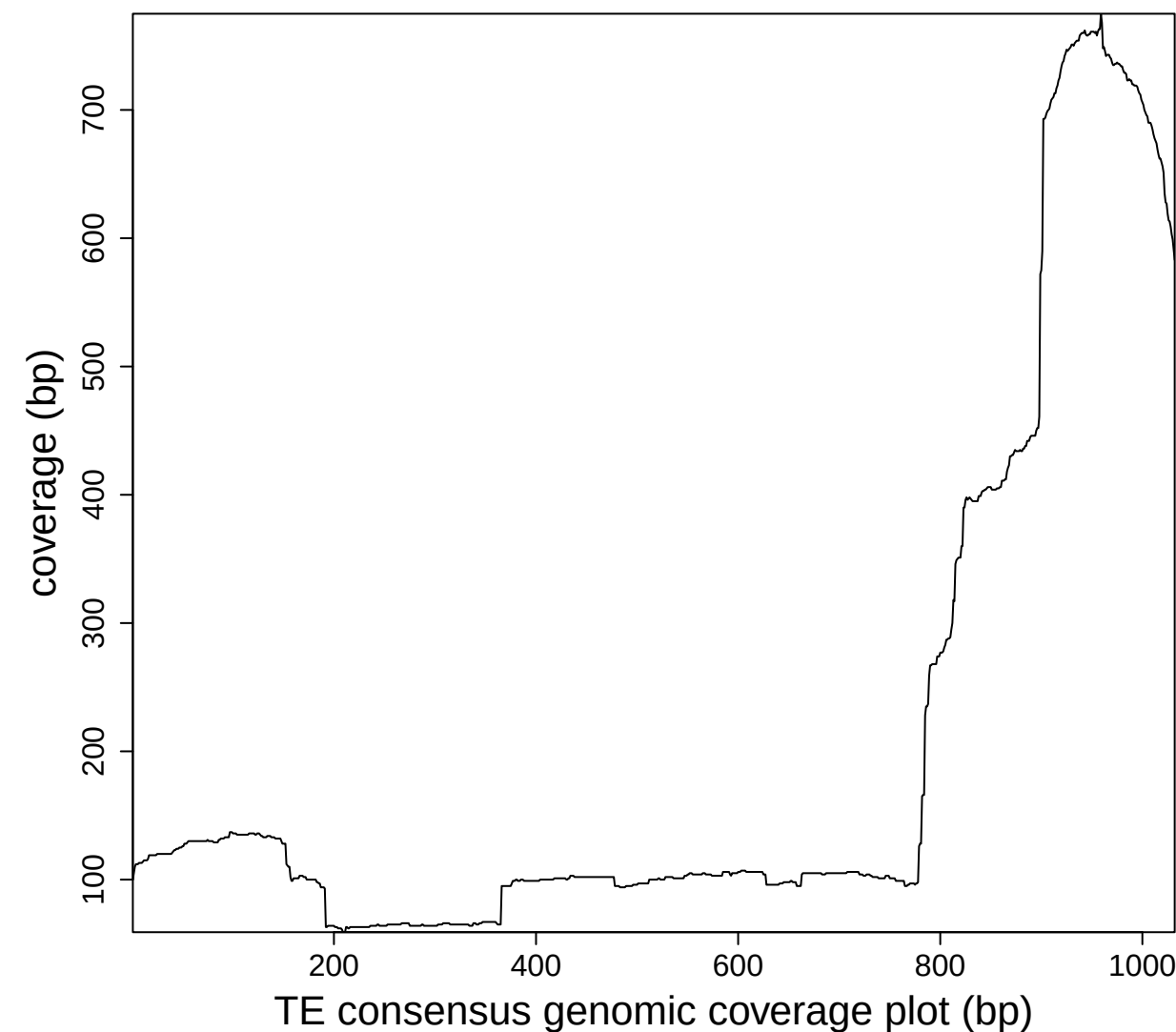
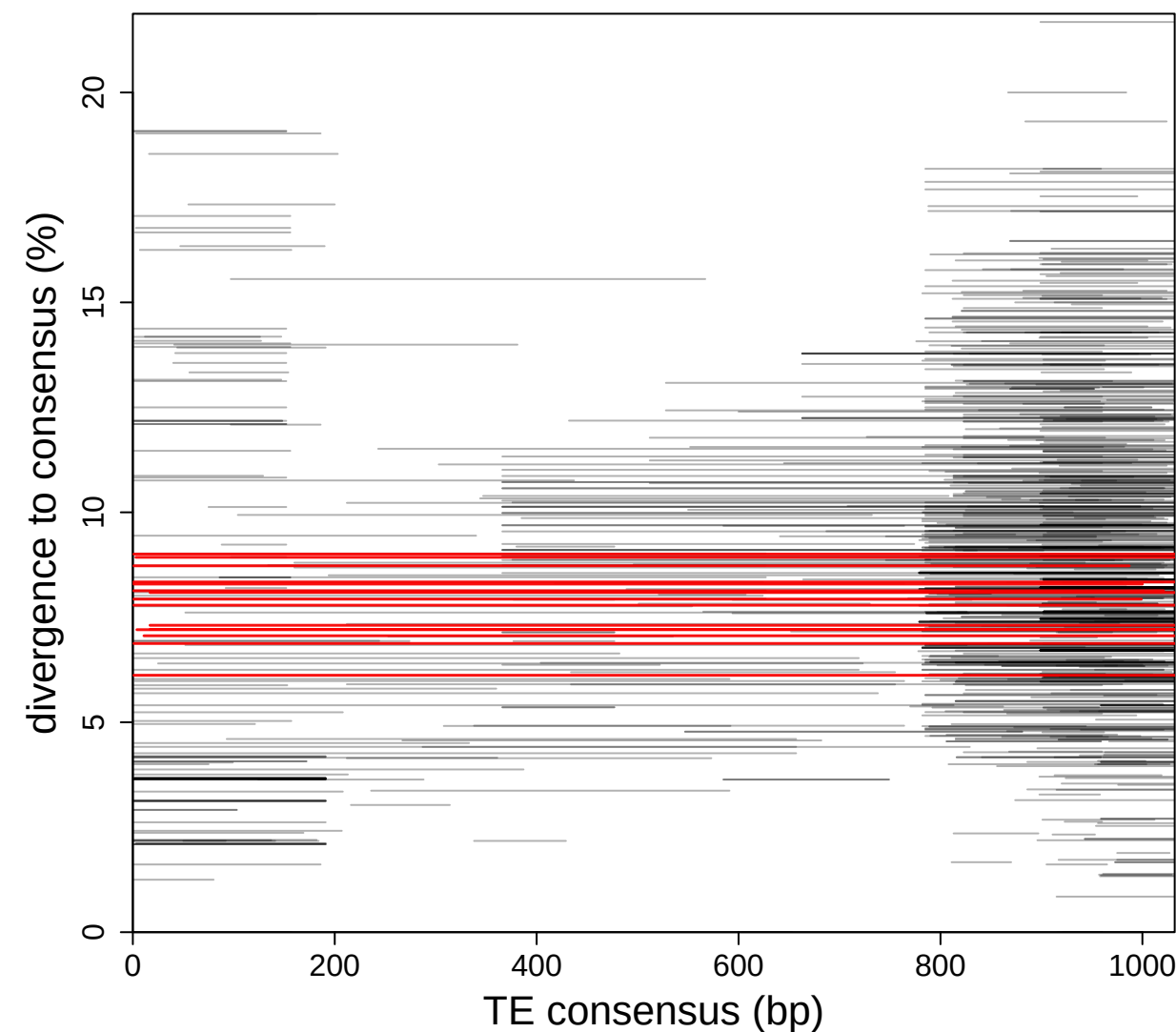
ly_7933.fasta.b.bed uf.bed g 2.bed fm 1.bed 0 0 bcln.fa aln.f
size: 3281bp; fragments: 1948; full length: 0 (>=2952.9bp)

After TEtrimmer Extended plot Blue lines are boundaries

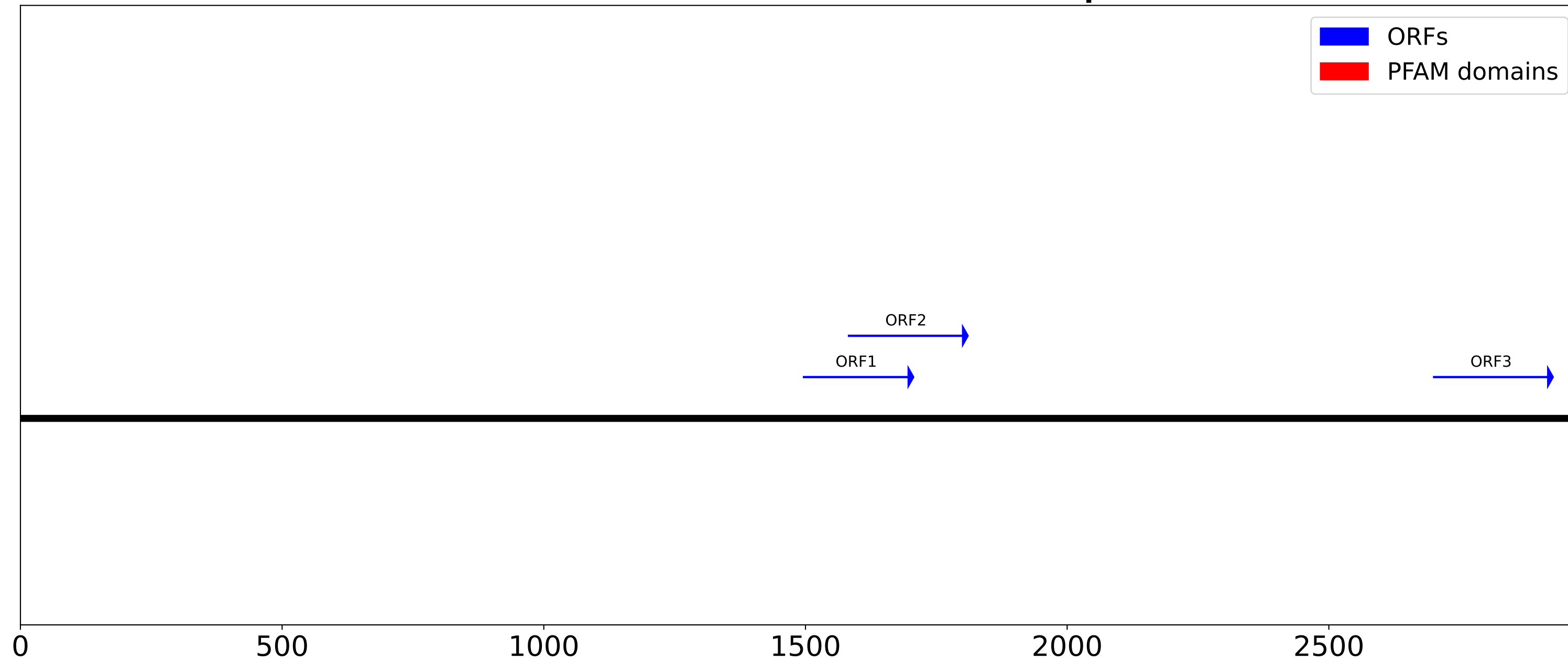


TE: rnd_5_family_7933
size: 1032bp; fragments: 994; full length: 15 (≥ 928.8 bp)

Before TEtrimmer 1032 bp



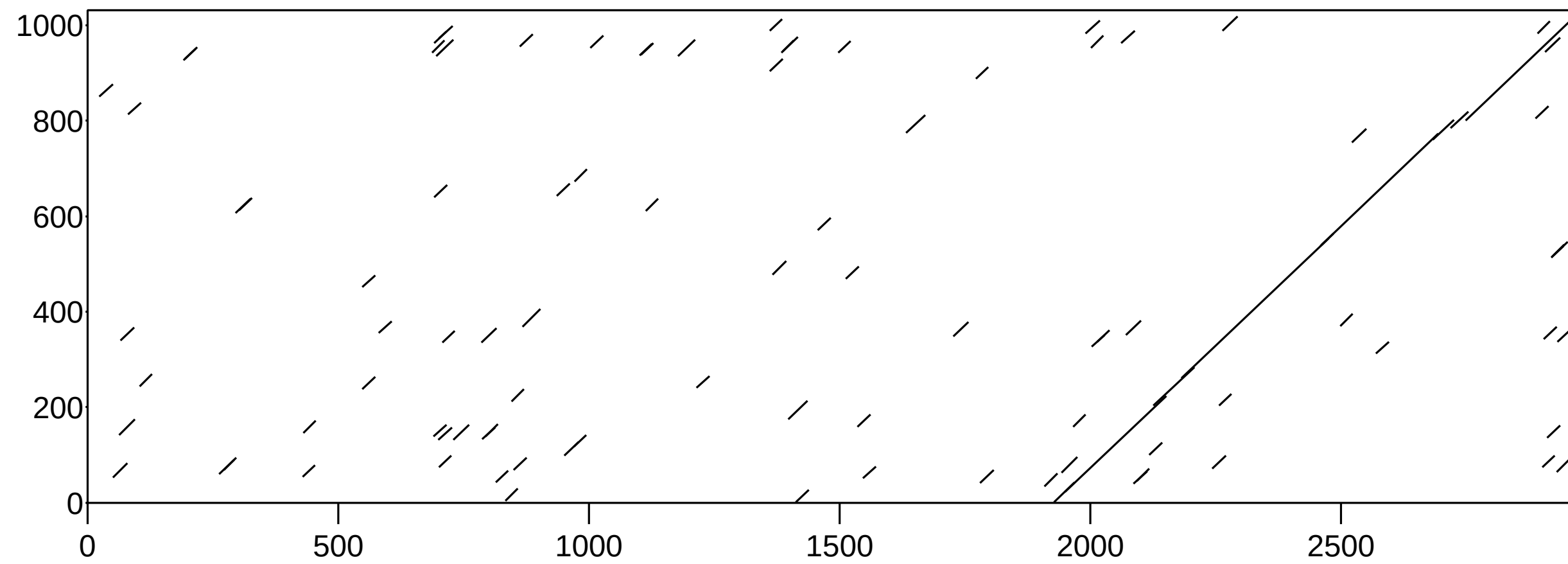
After TEtrimmer ORF and PFAM domain plot



Dotplot

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

E consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)