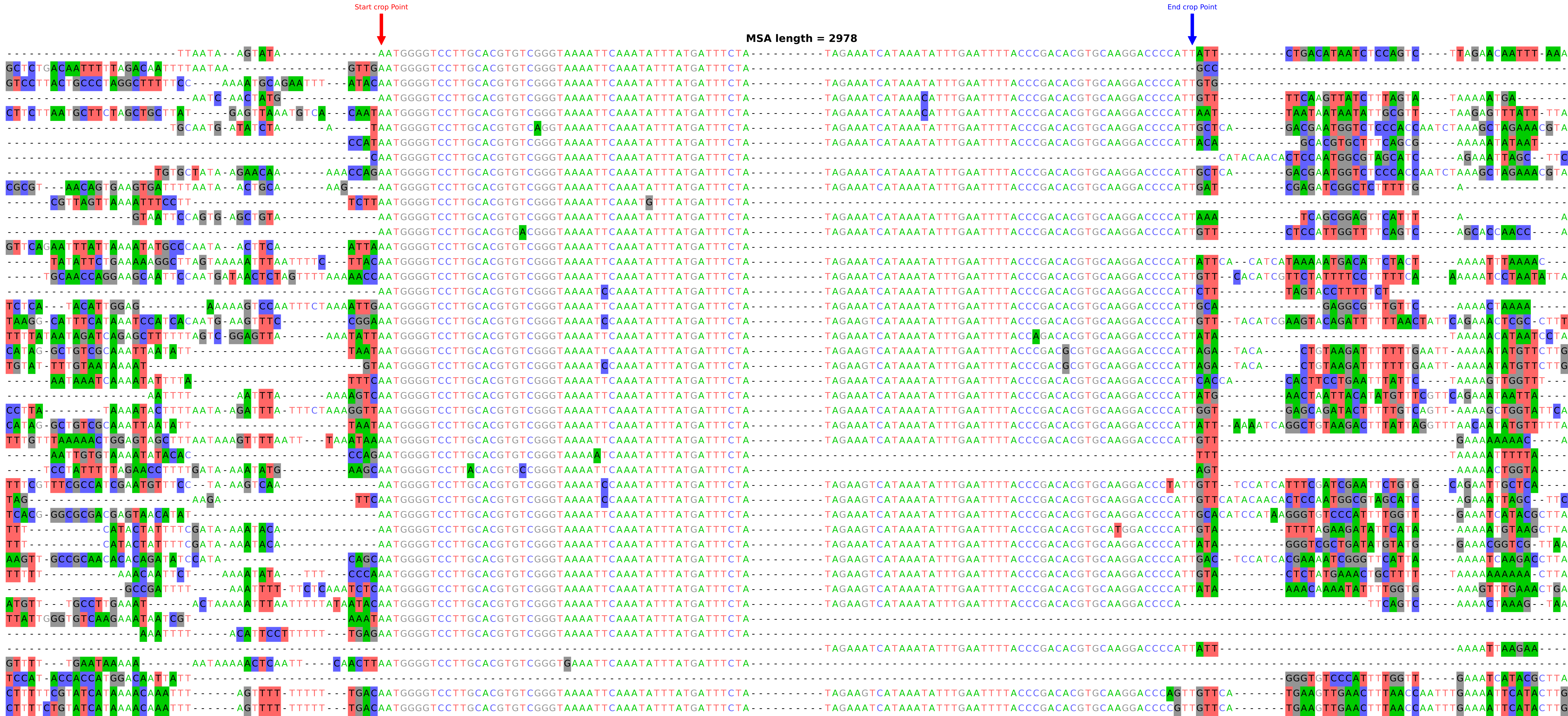
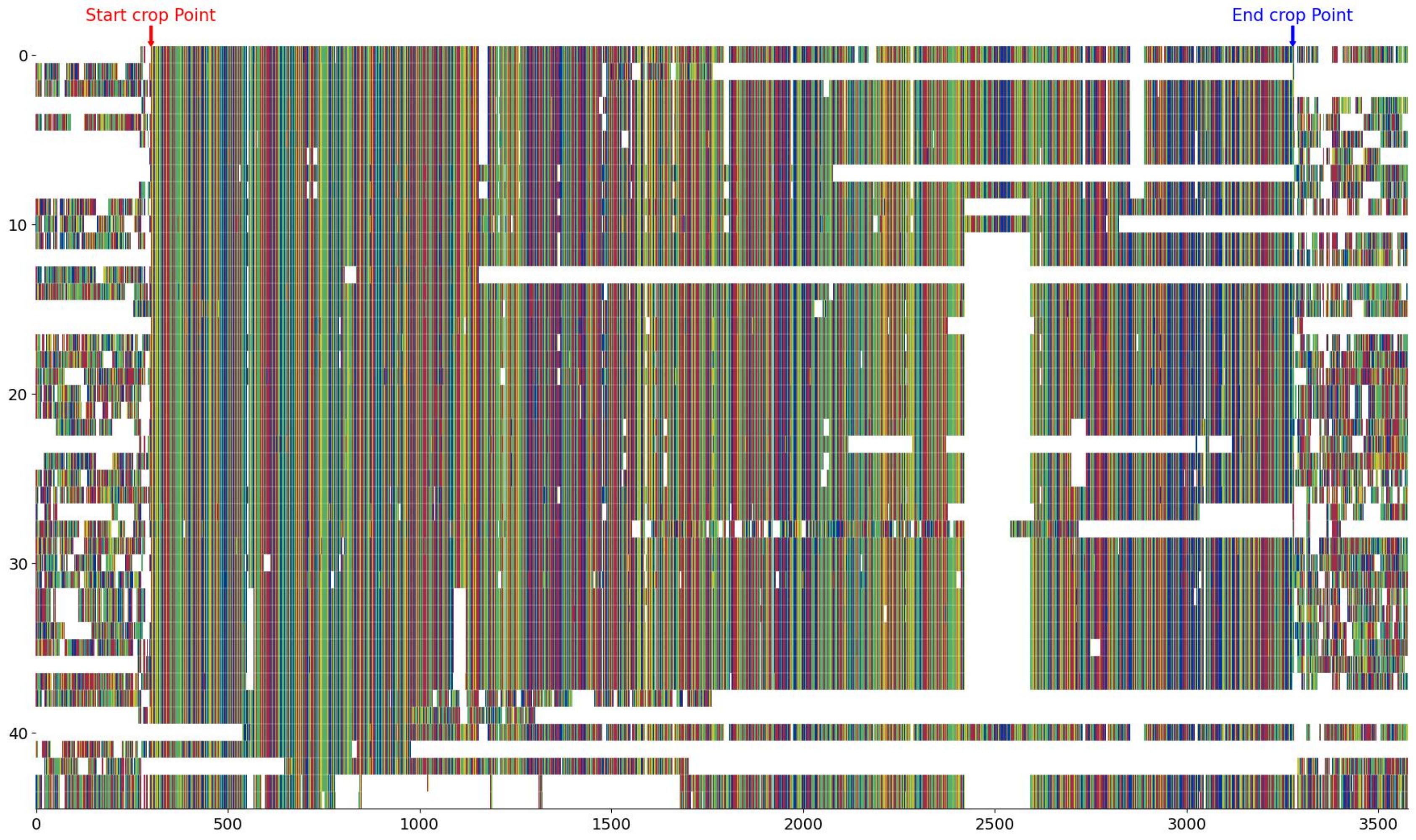
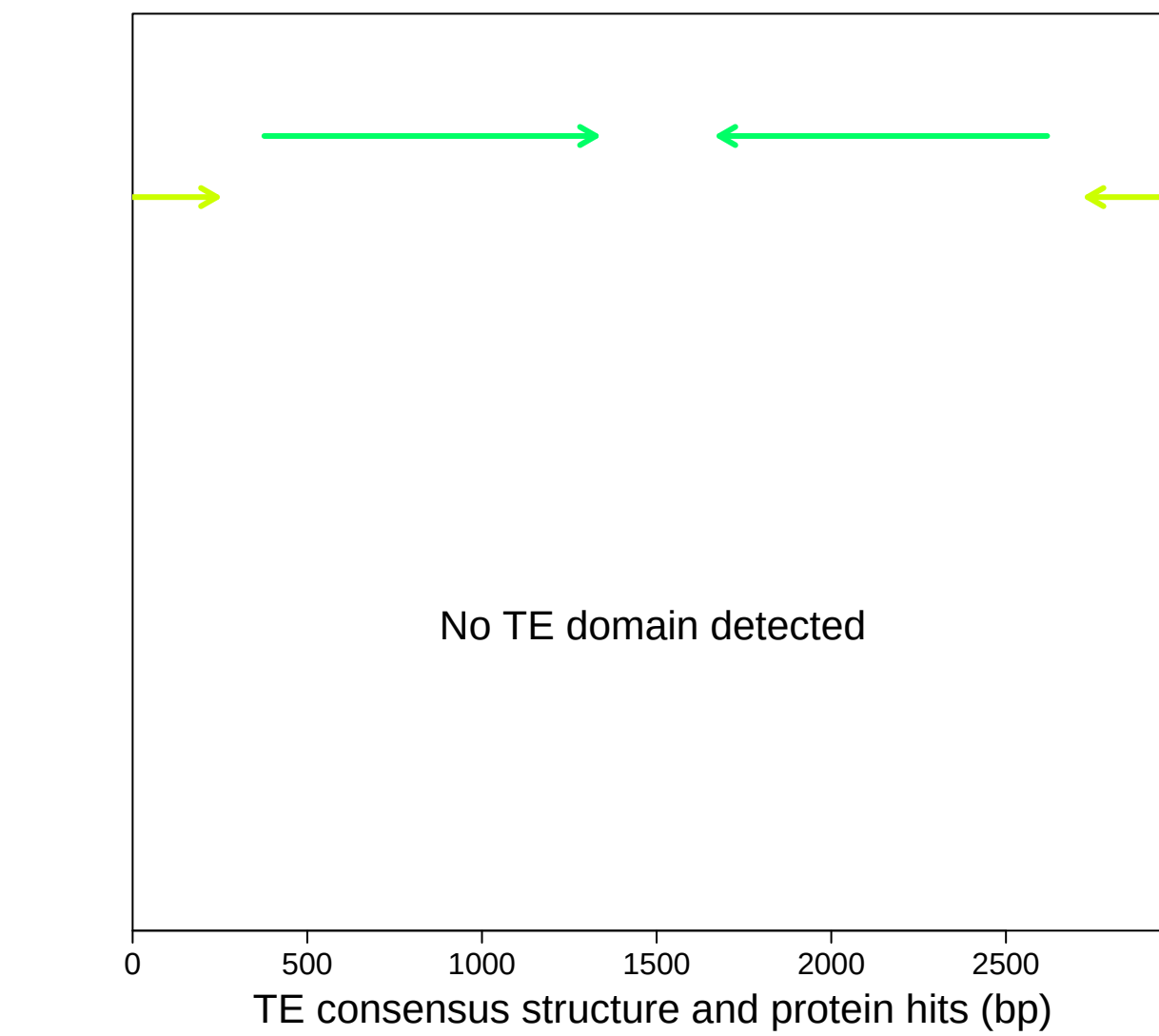
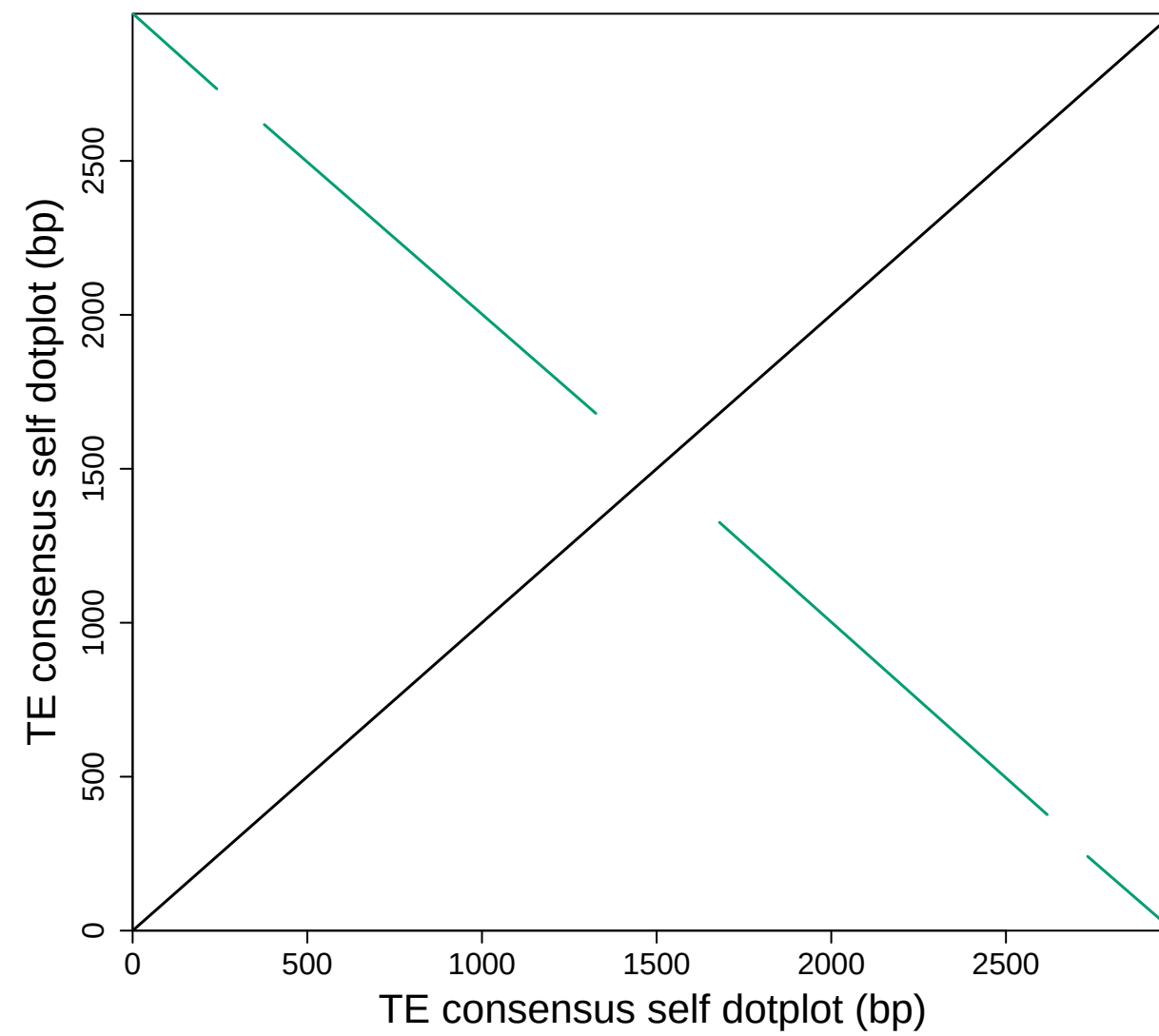
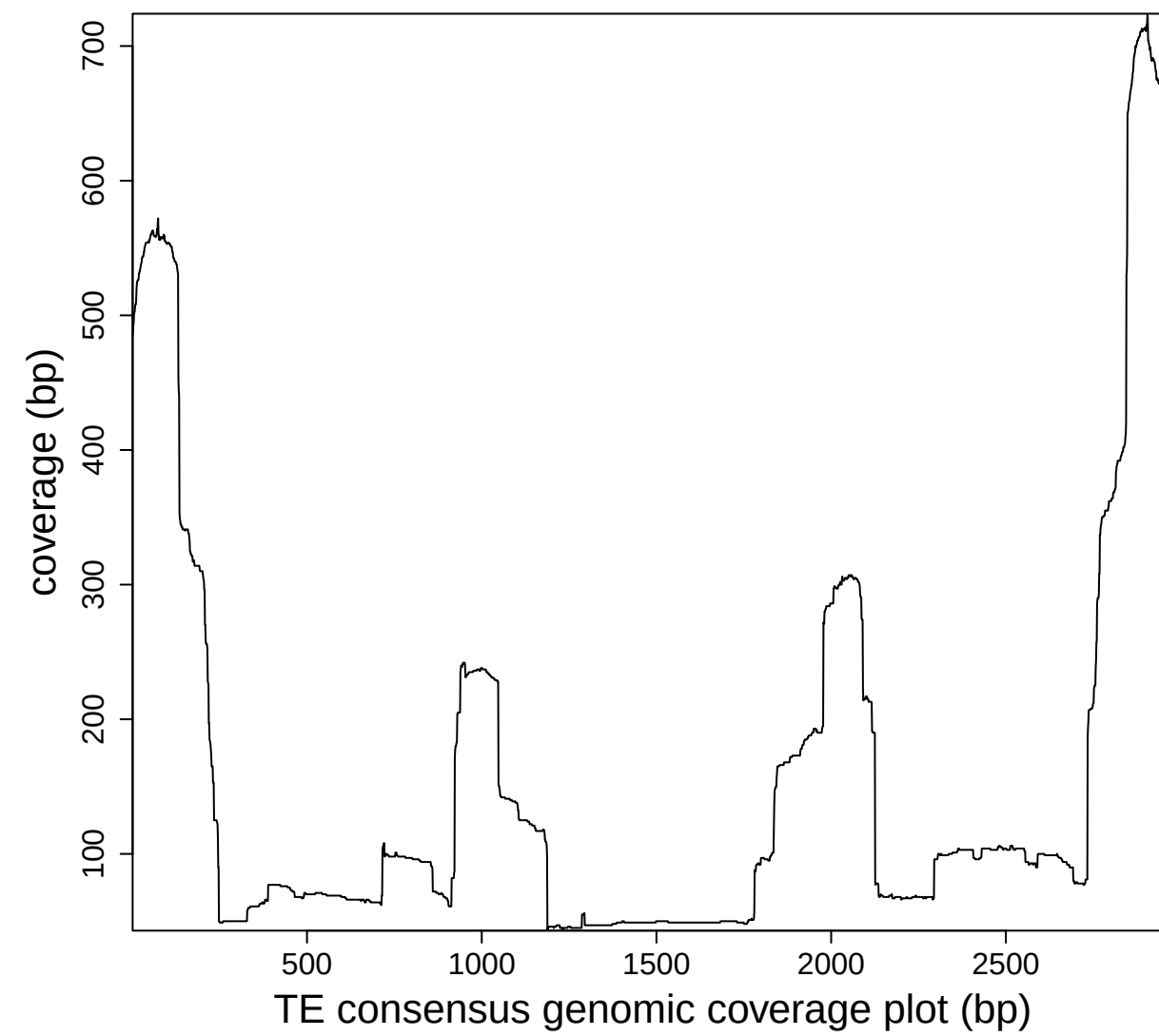
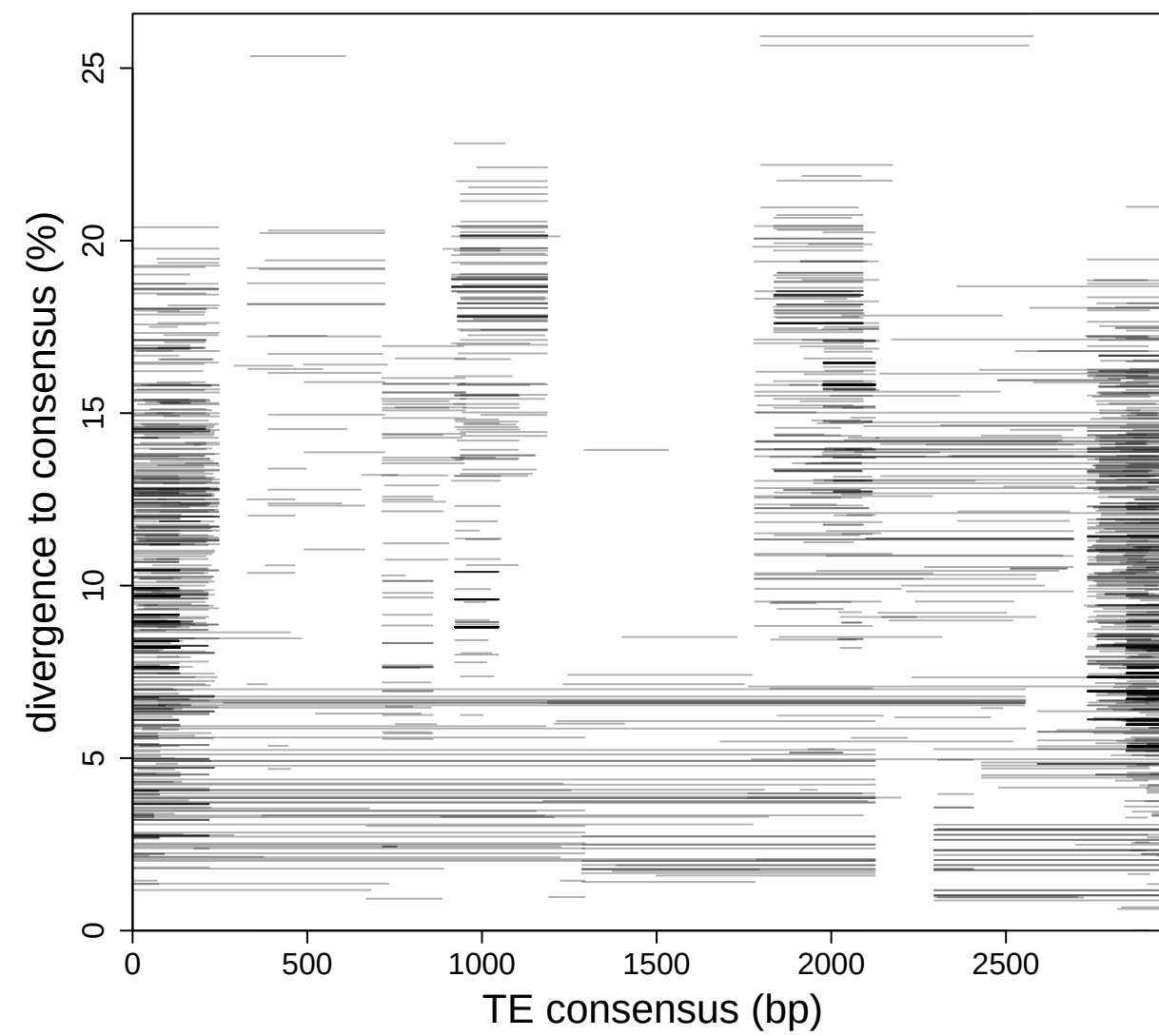






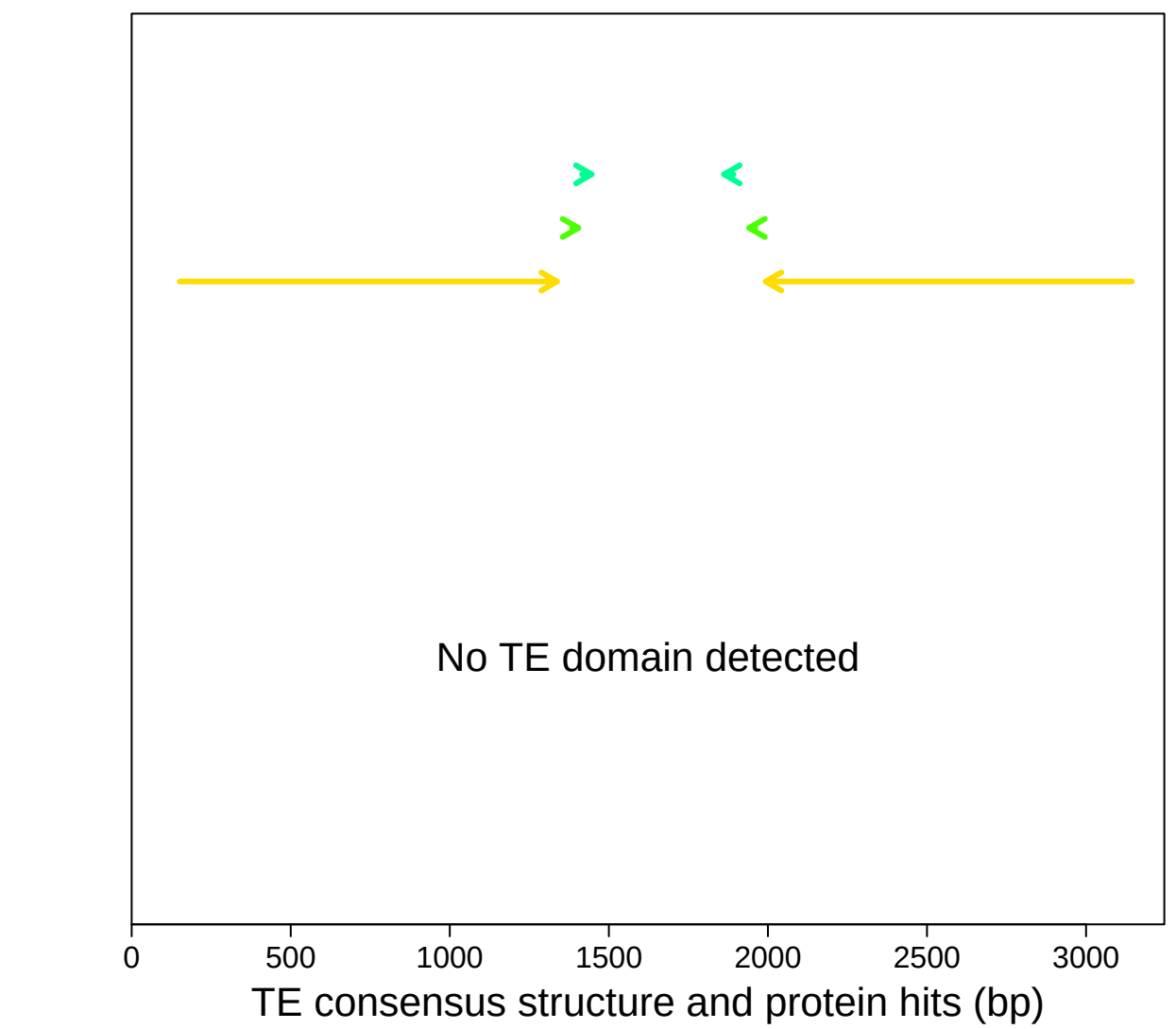
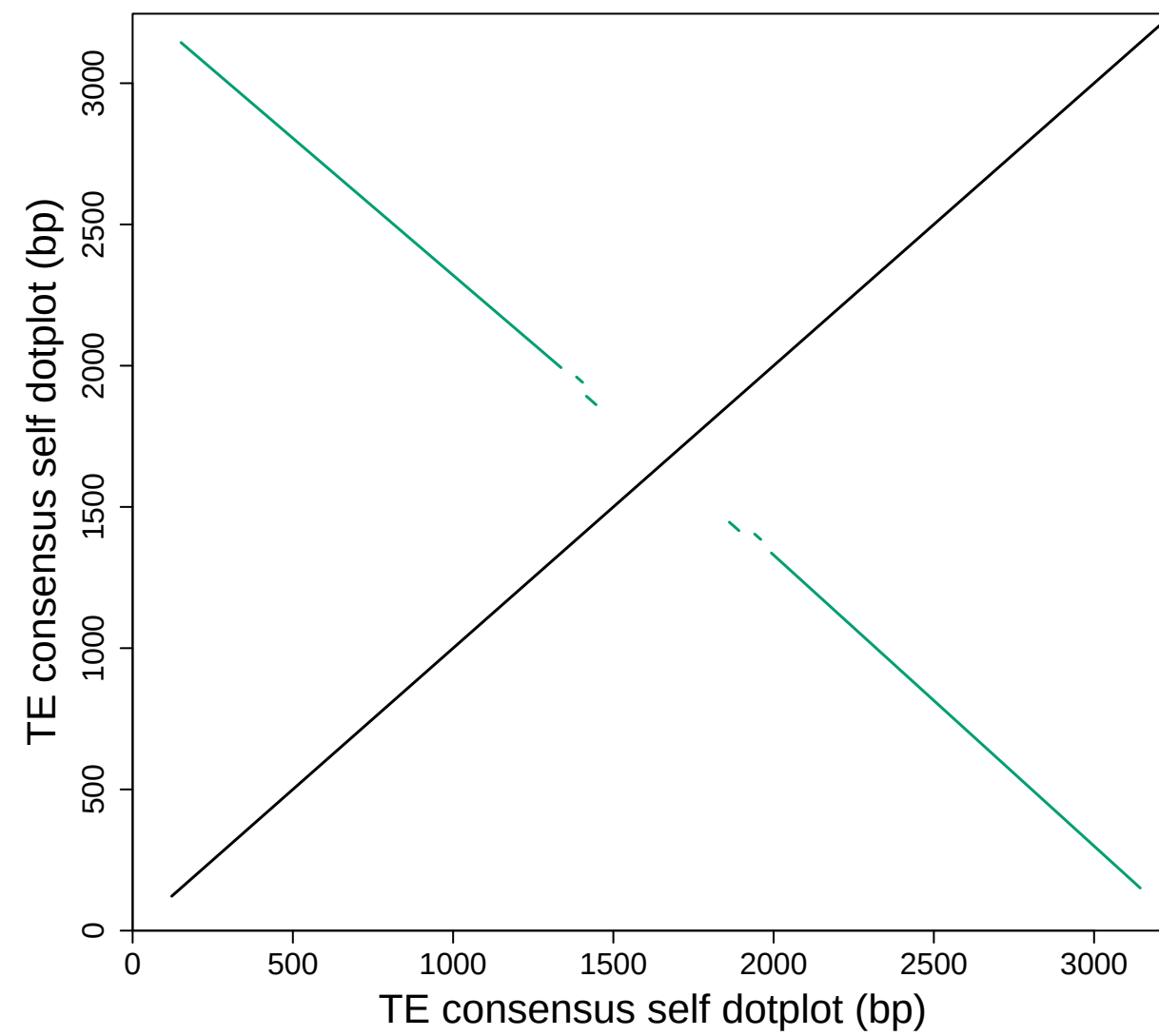
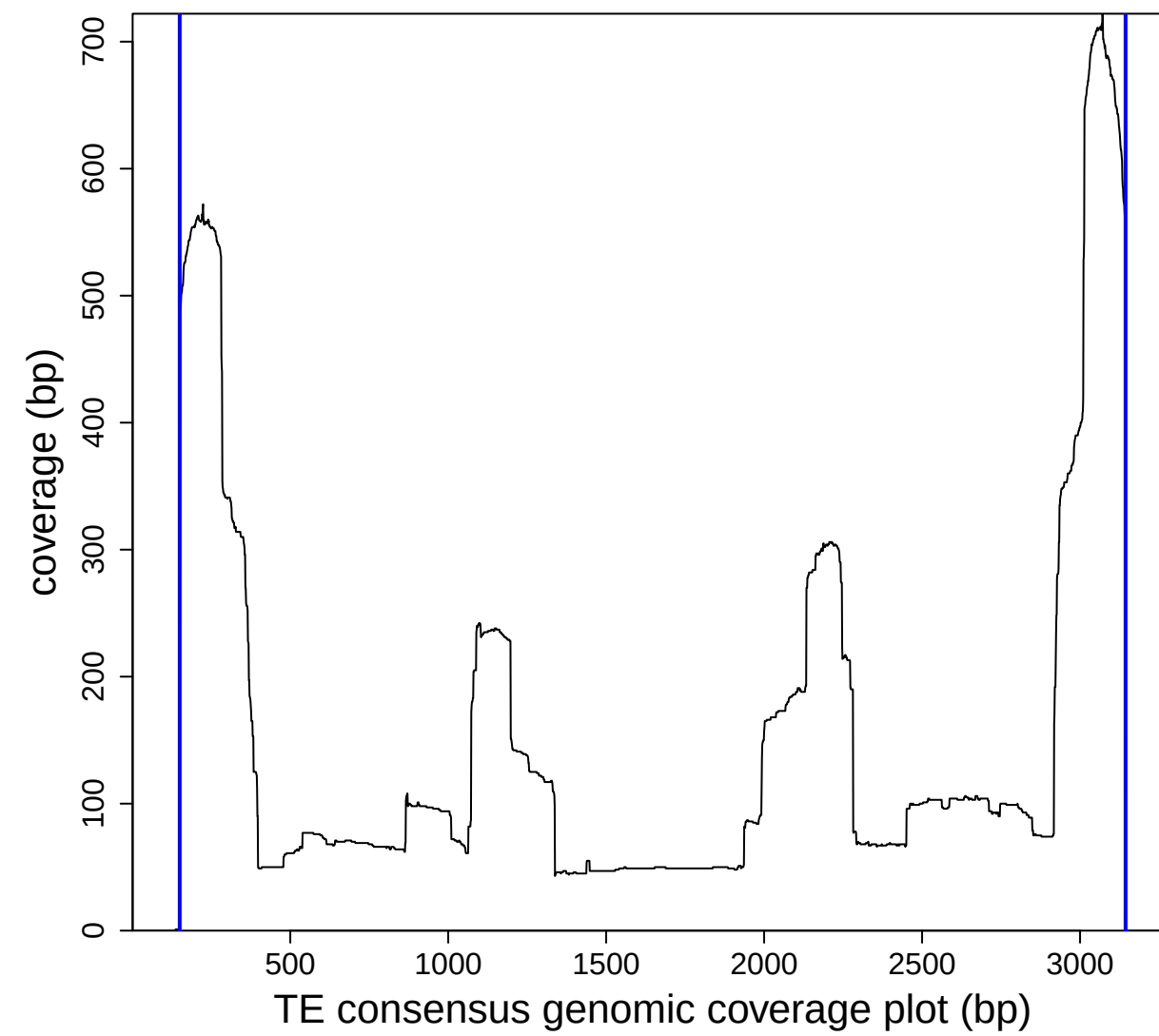
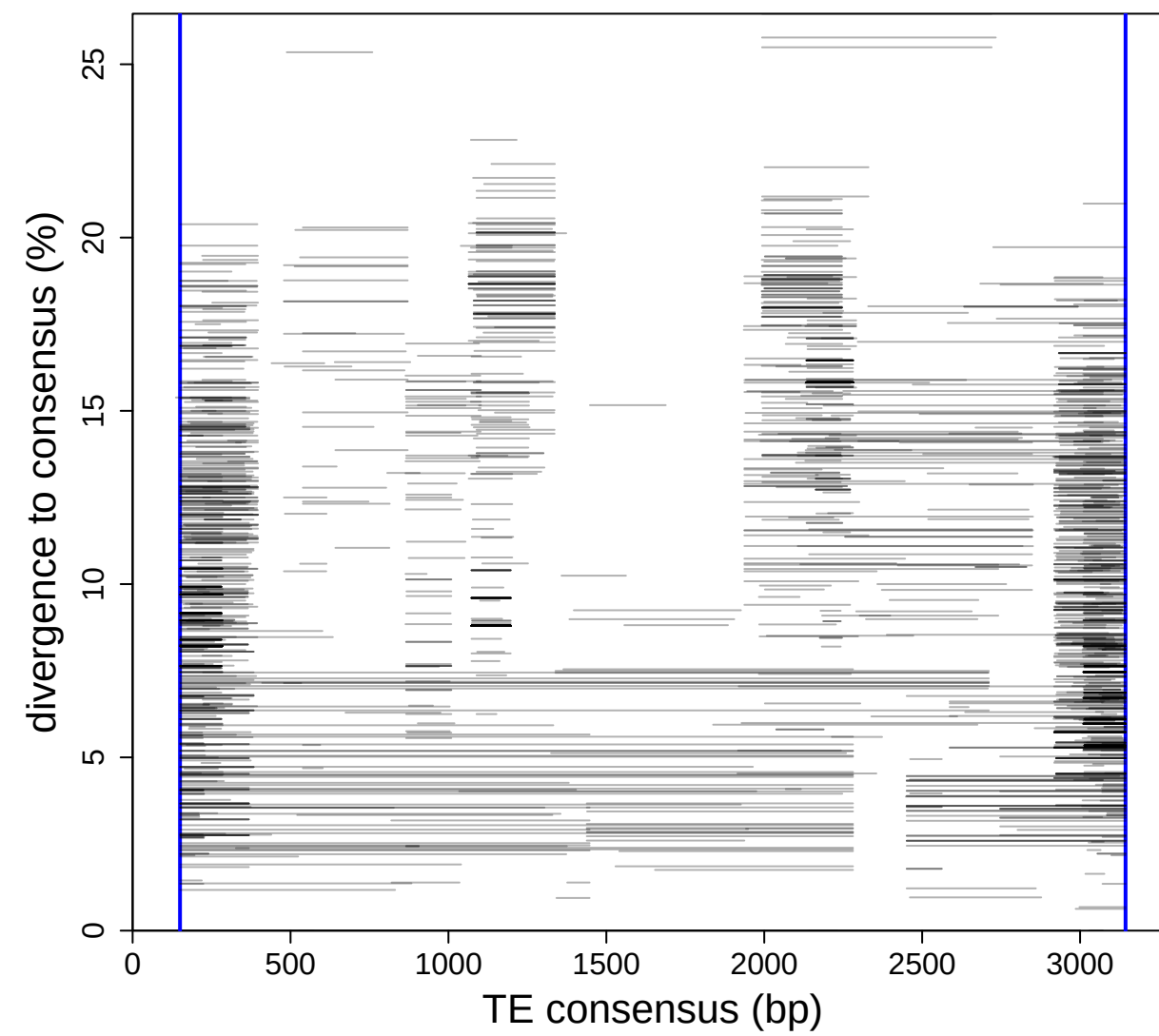
g_1.bed fm_1.bed 0 0 n.bed g_1.bed fm_2.bed 0 0 bcln.fa ε
size: 2978bp; fragments: 1972; full length: 0 (>=2680.2bp)

After TEtrimmer 2978 bp



f.bed_g_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bc
size: 3294bp; fragments: 1971; full length: 0 (>=2964.6bp)

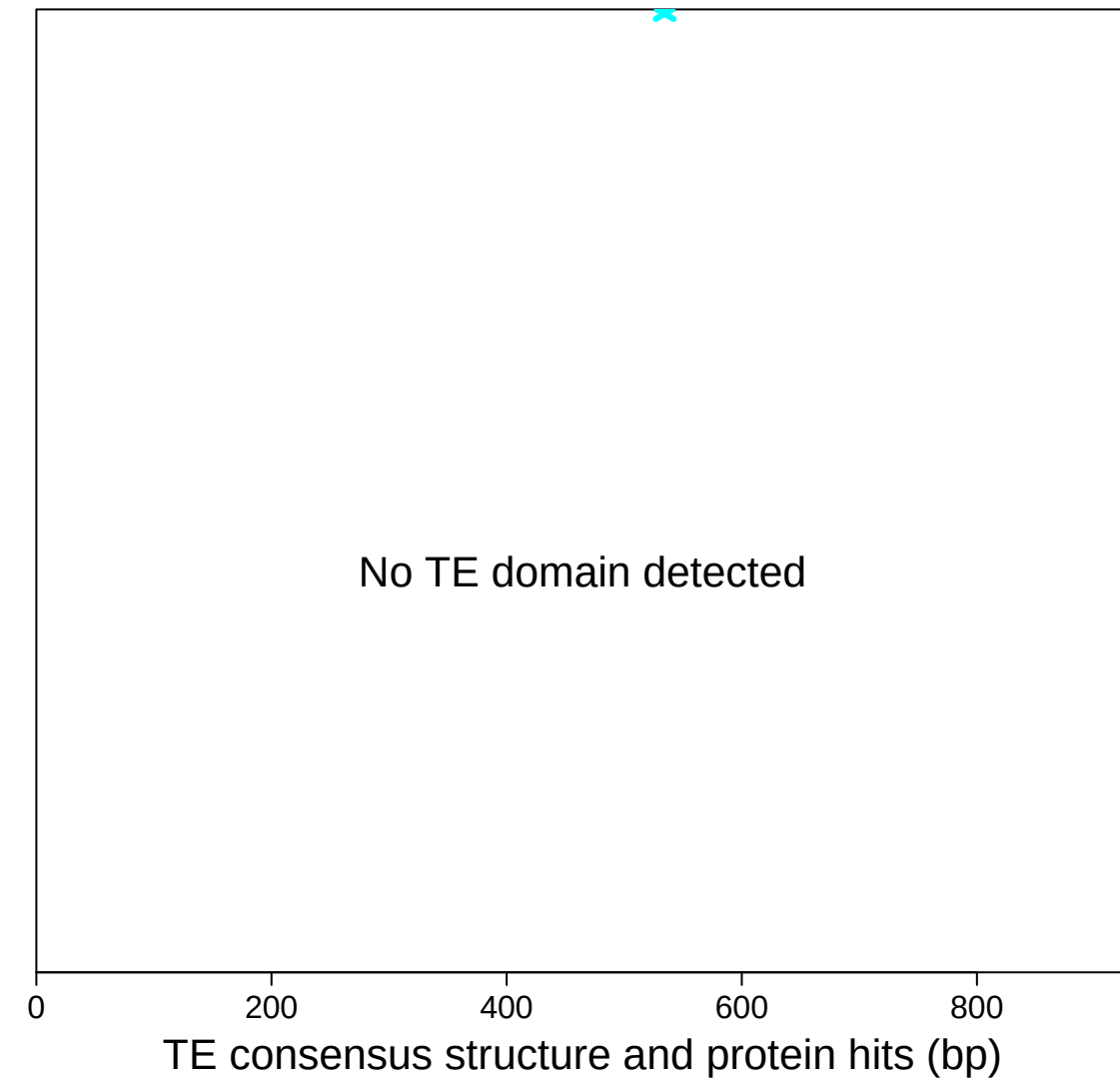
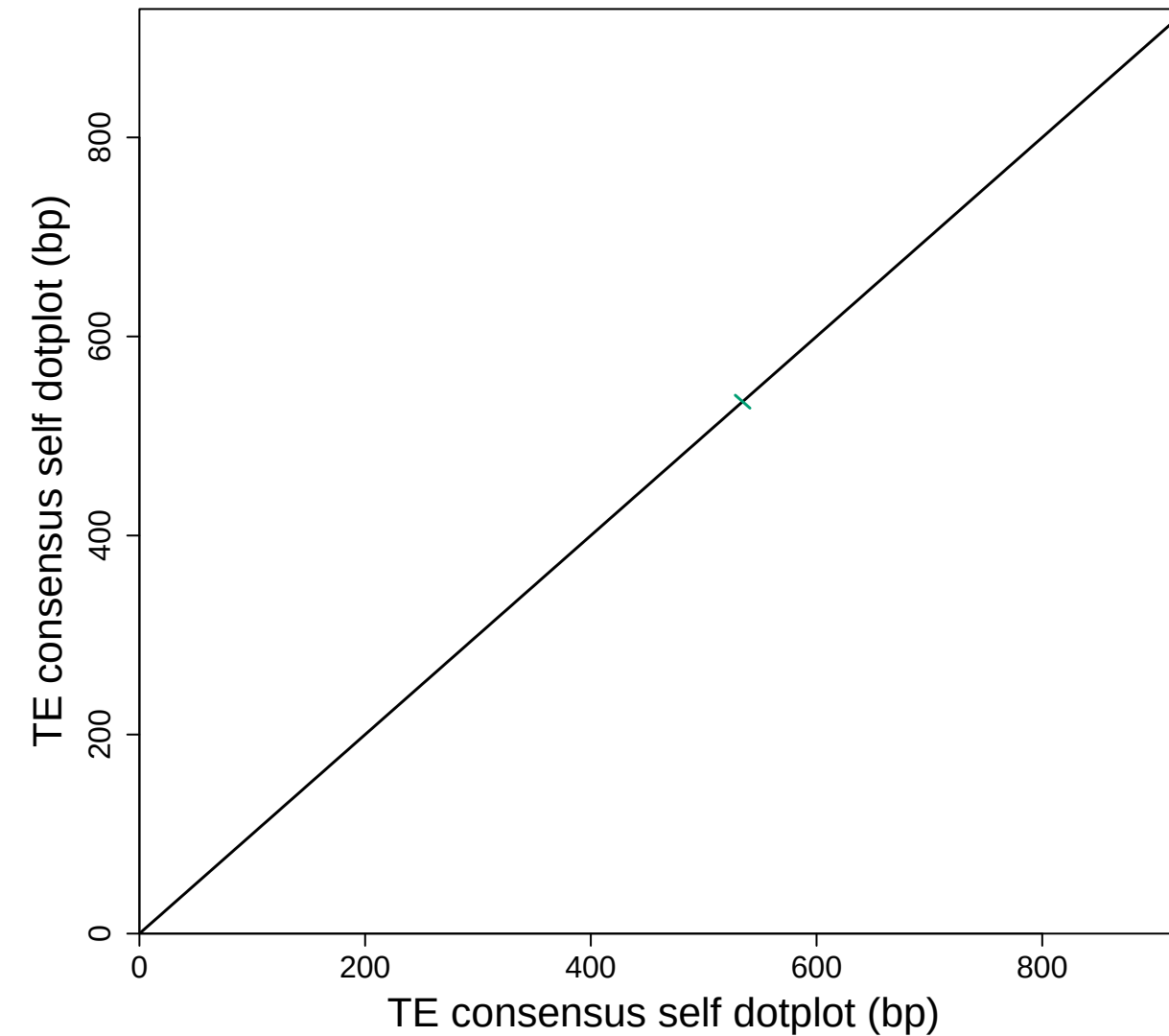
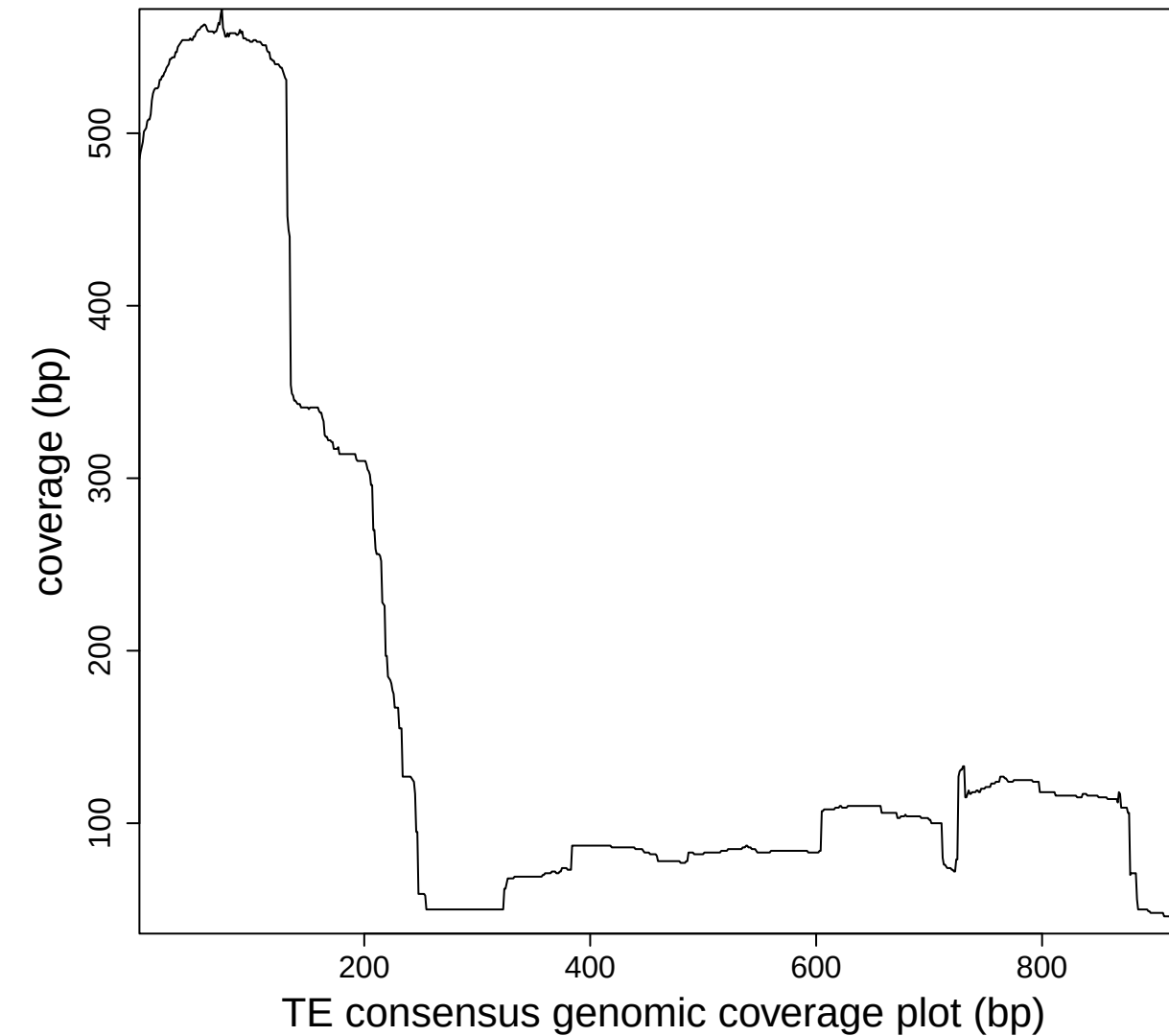
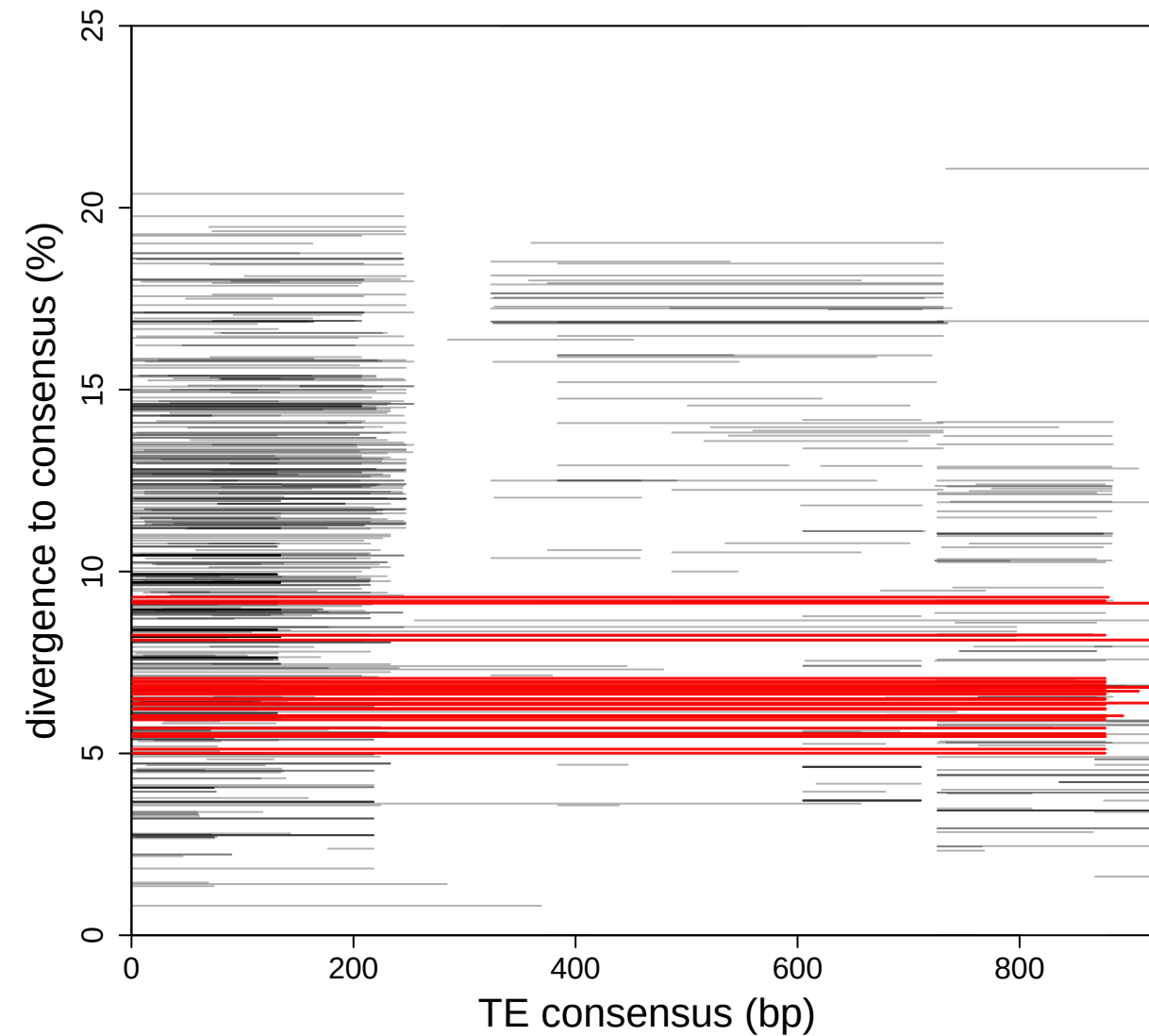
After TEtrimmer Extended plot Blue lines are boundaries



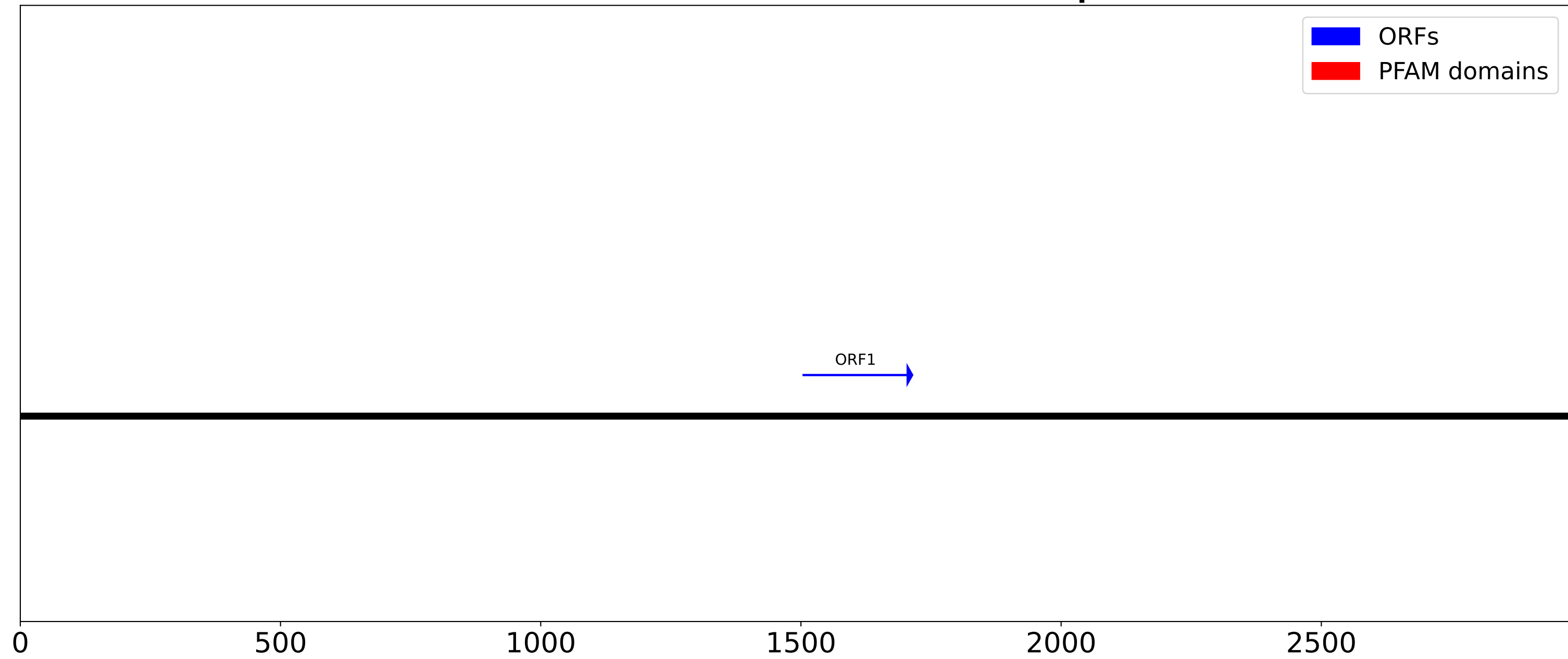
Before TEtrimmer 929 bp

TE: rnd_5_family_1477

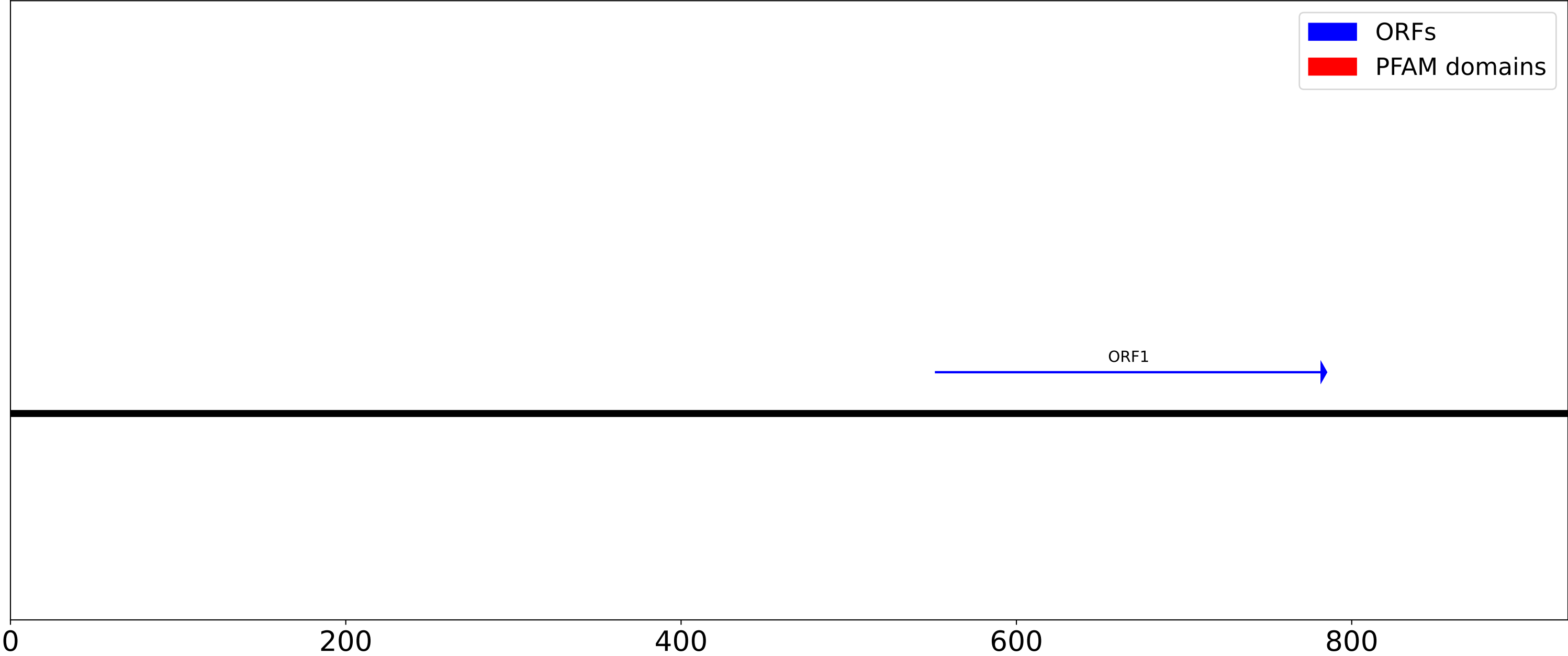
size: 929bp; fragments: 786; full length: 36 (≥ 836.1 bp)



After TEtrimmer ORF and PFAM domain plot



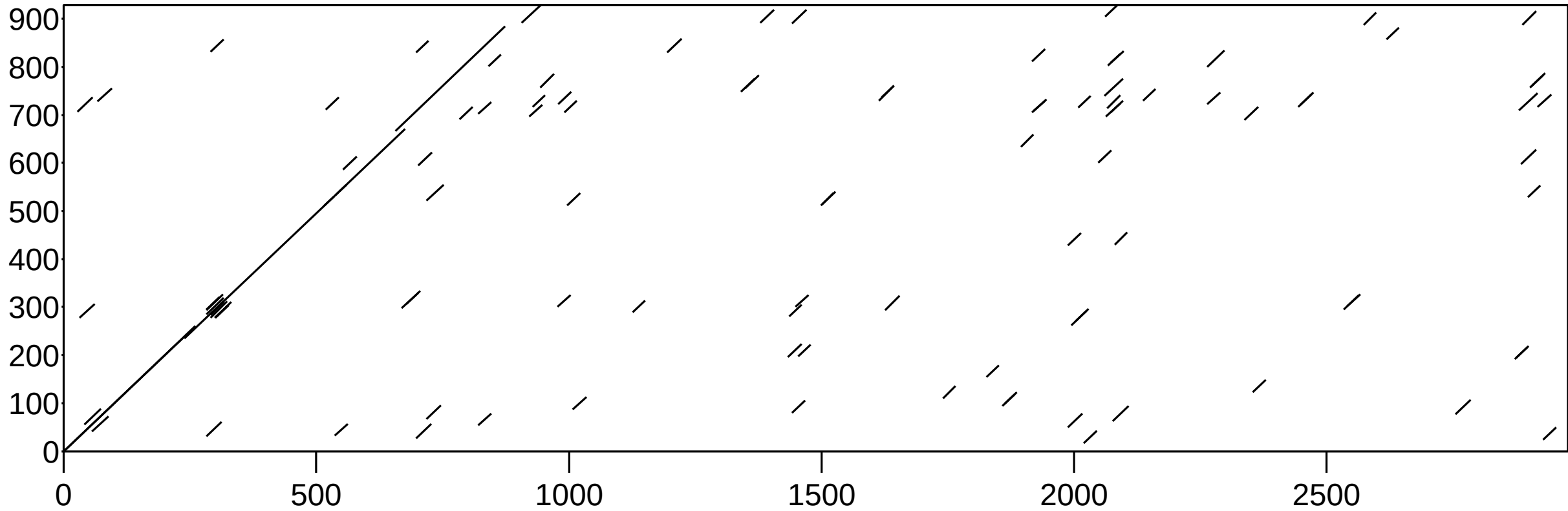
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

: consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)