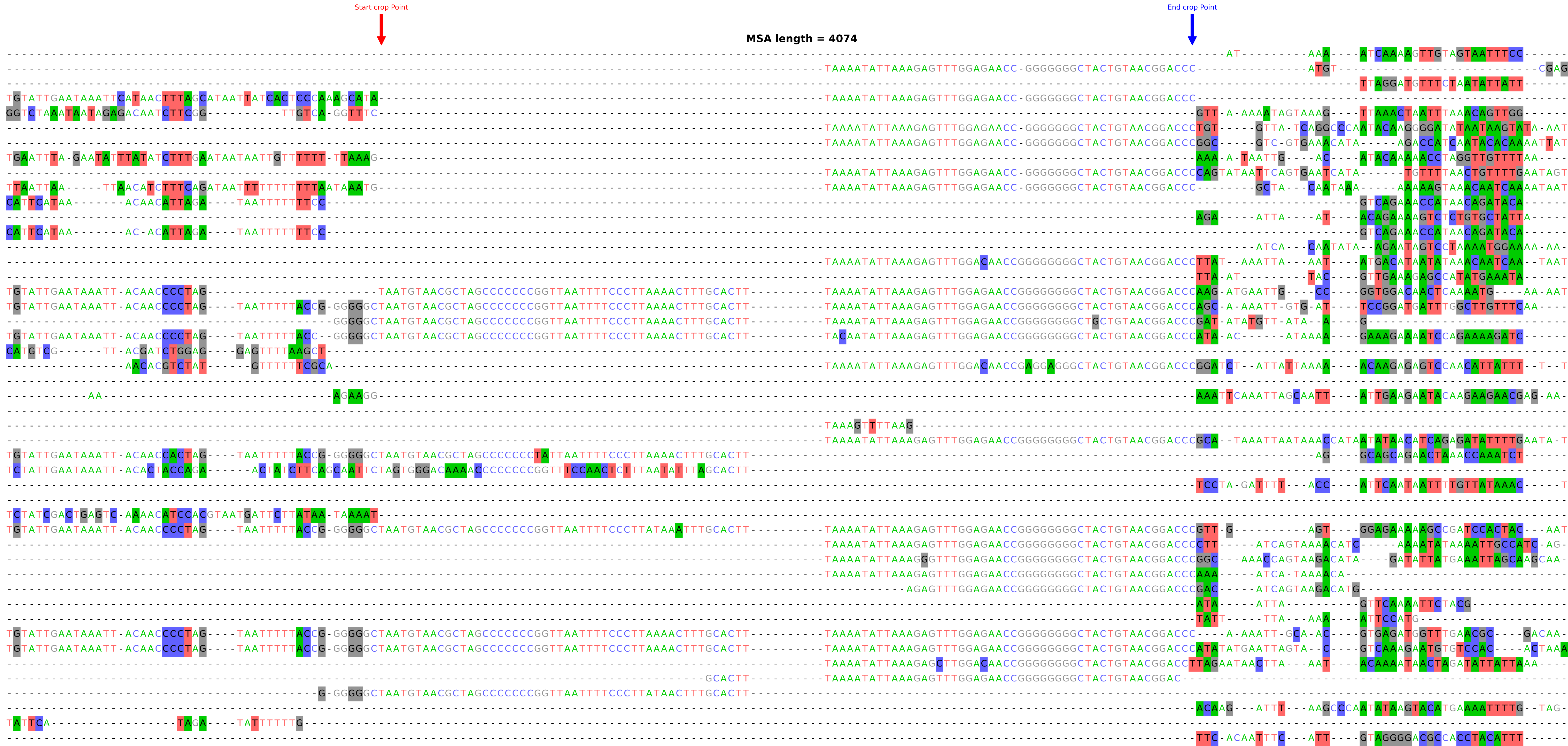
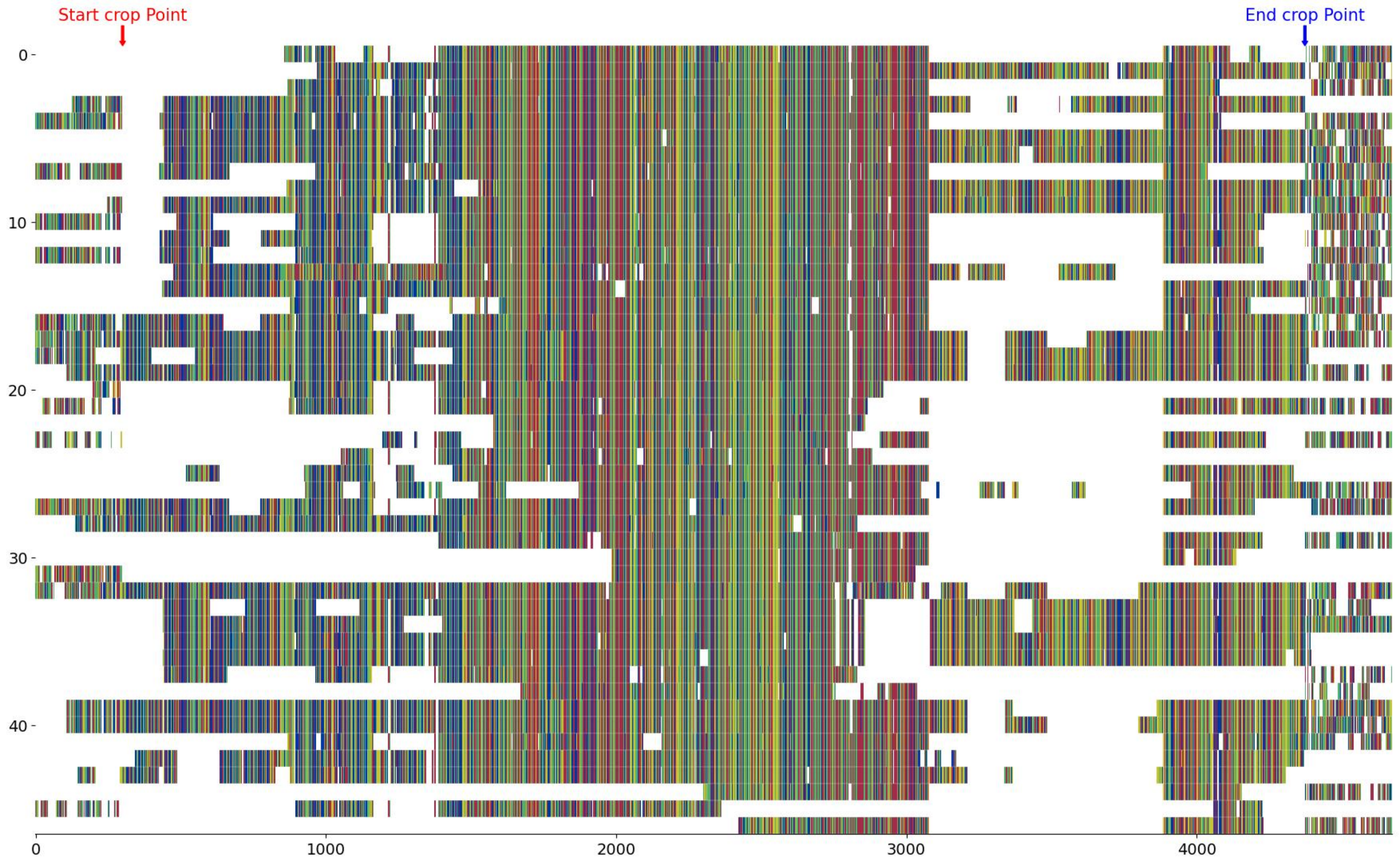
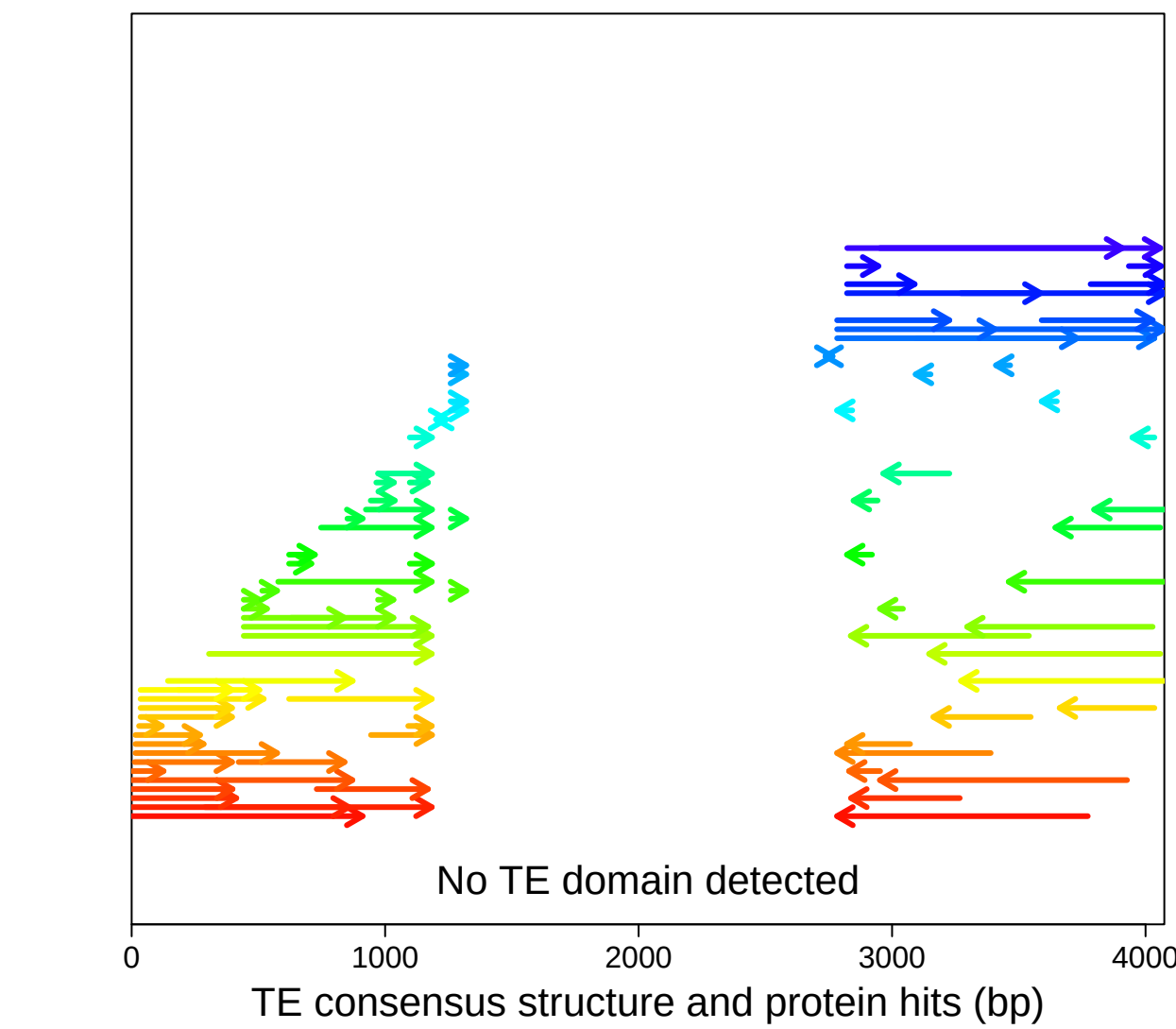
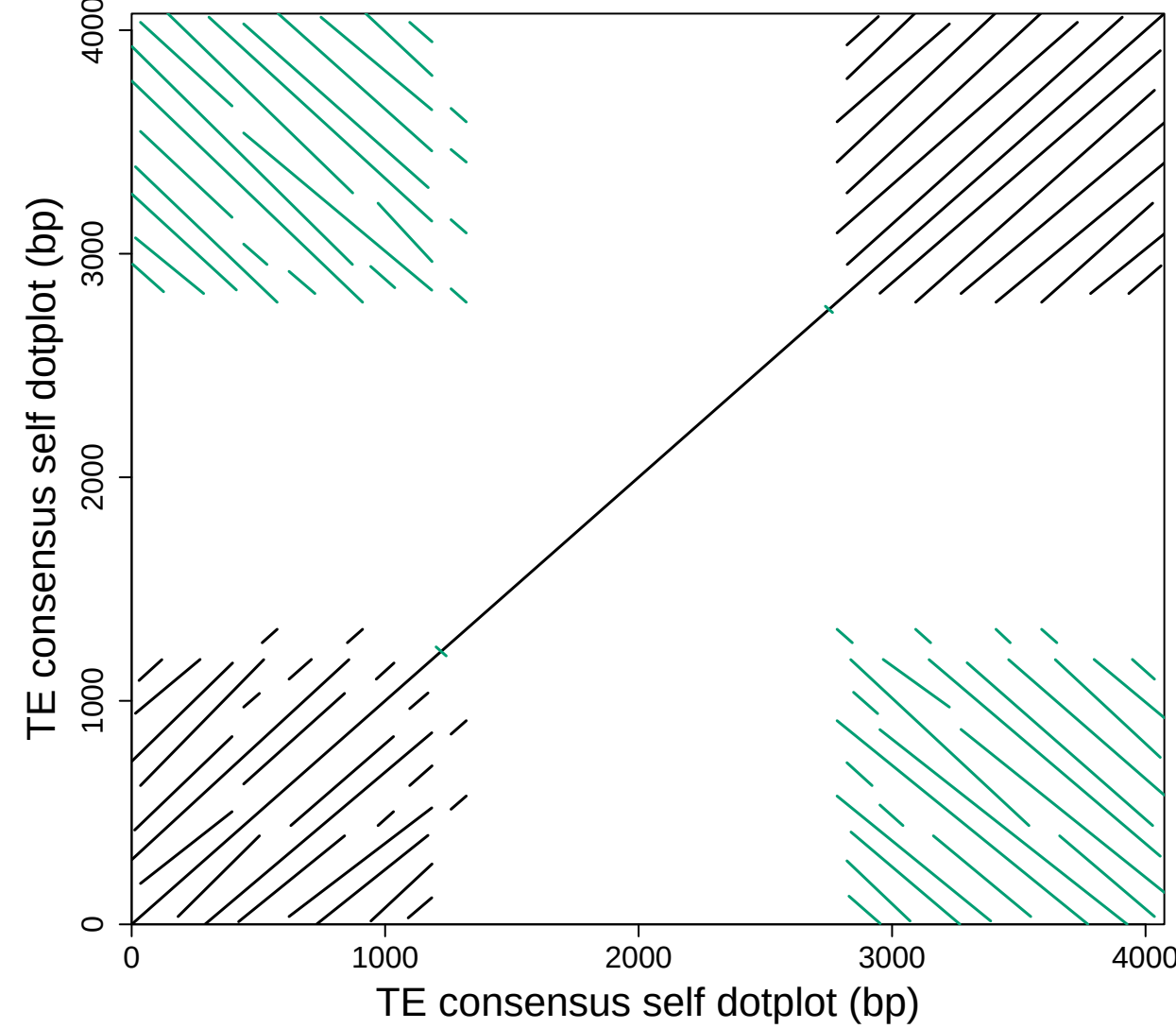
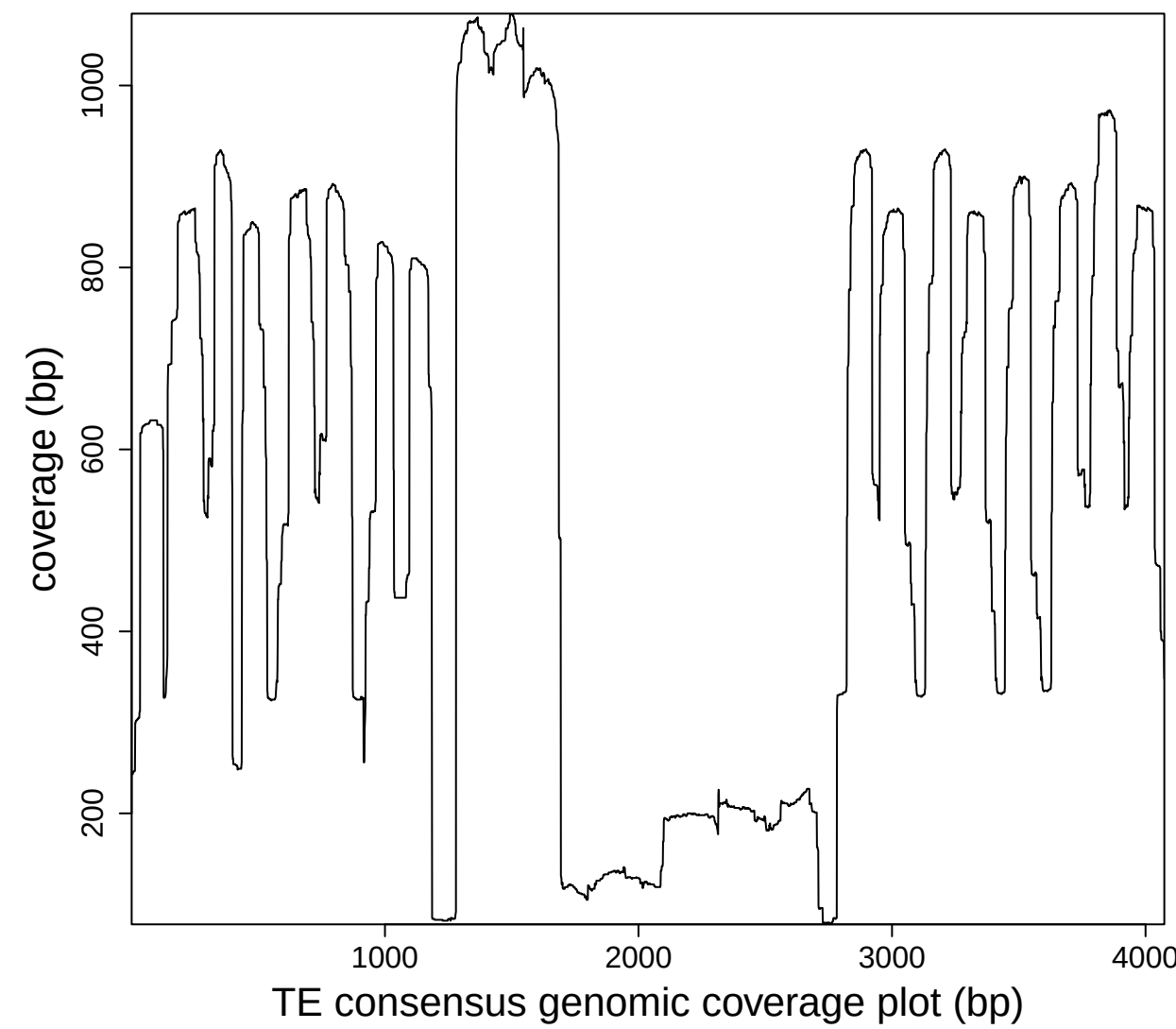
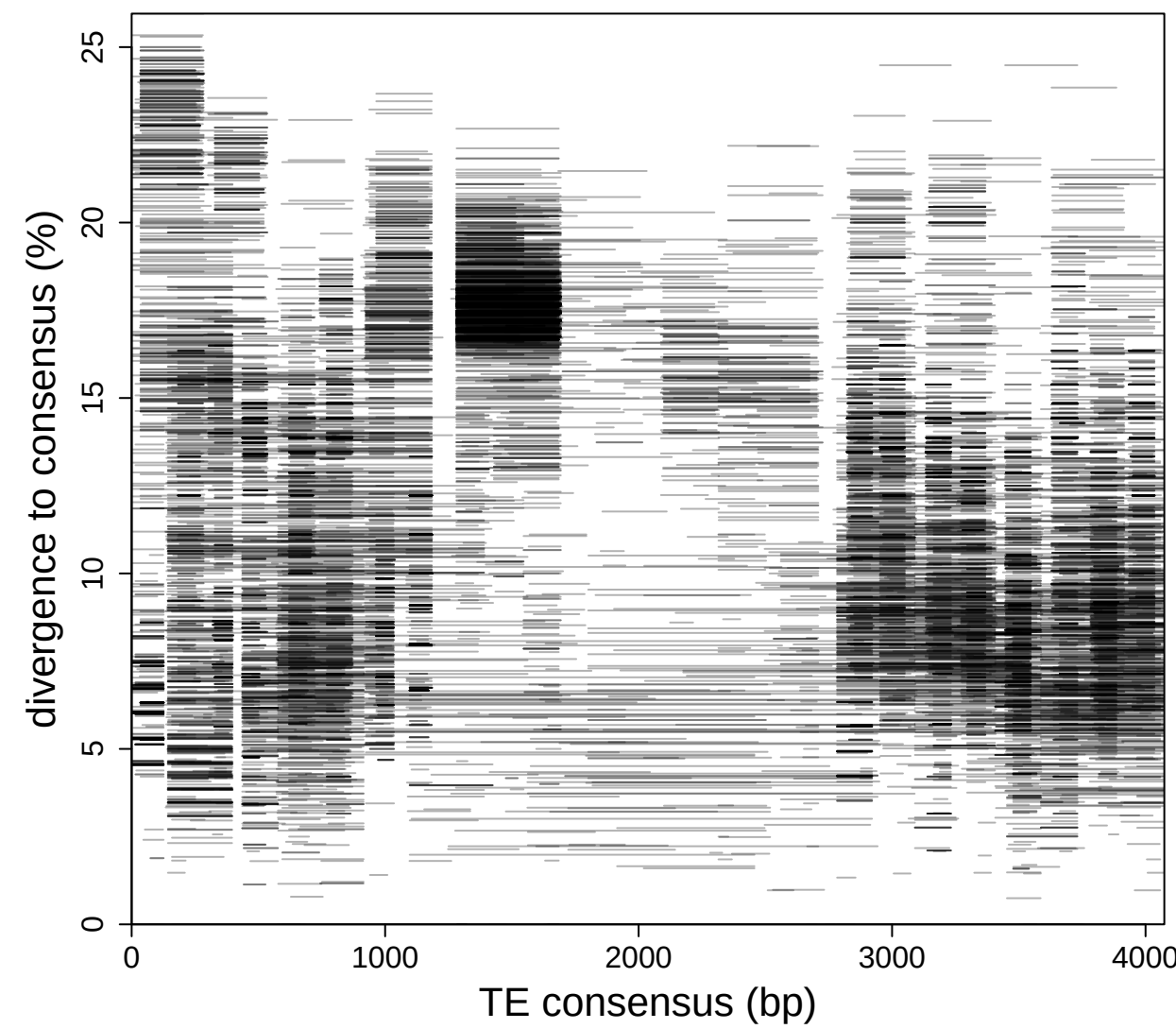






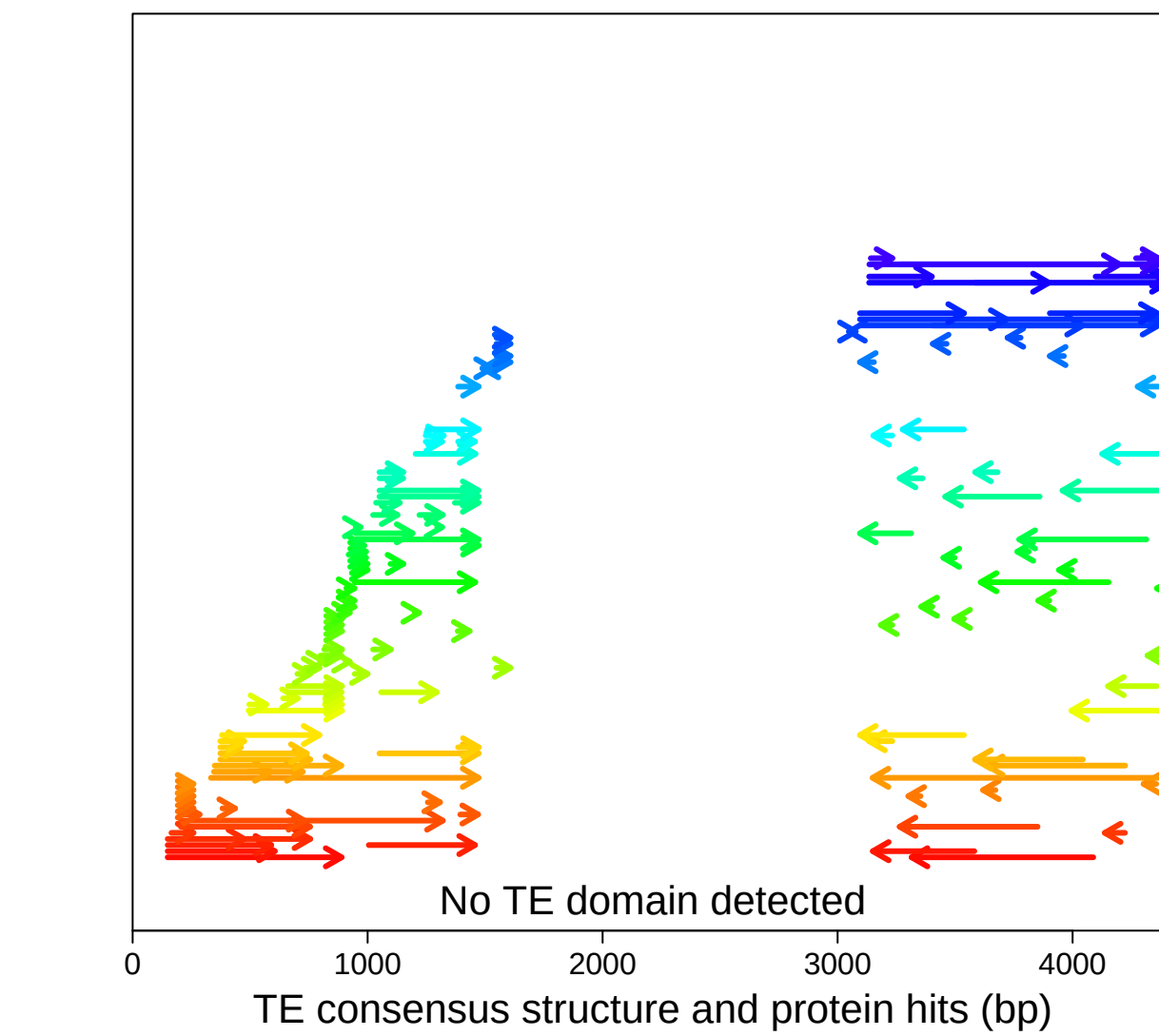
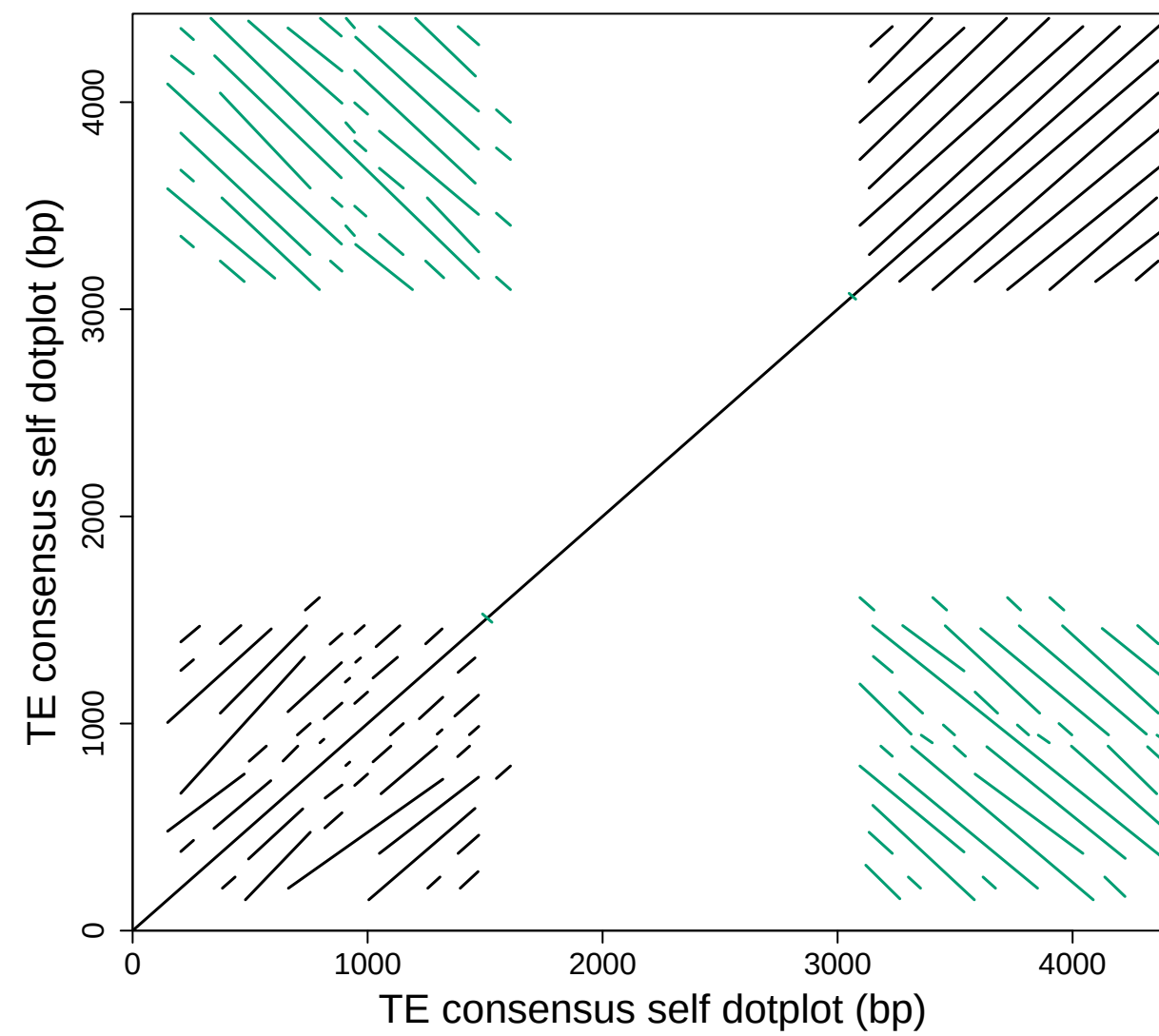
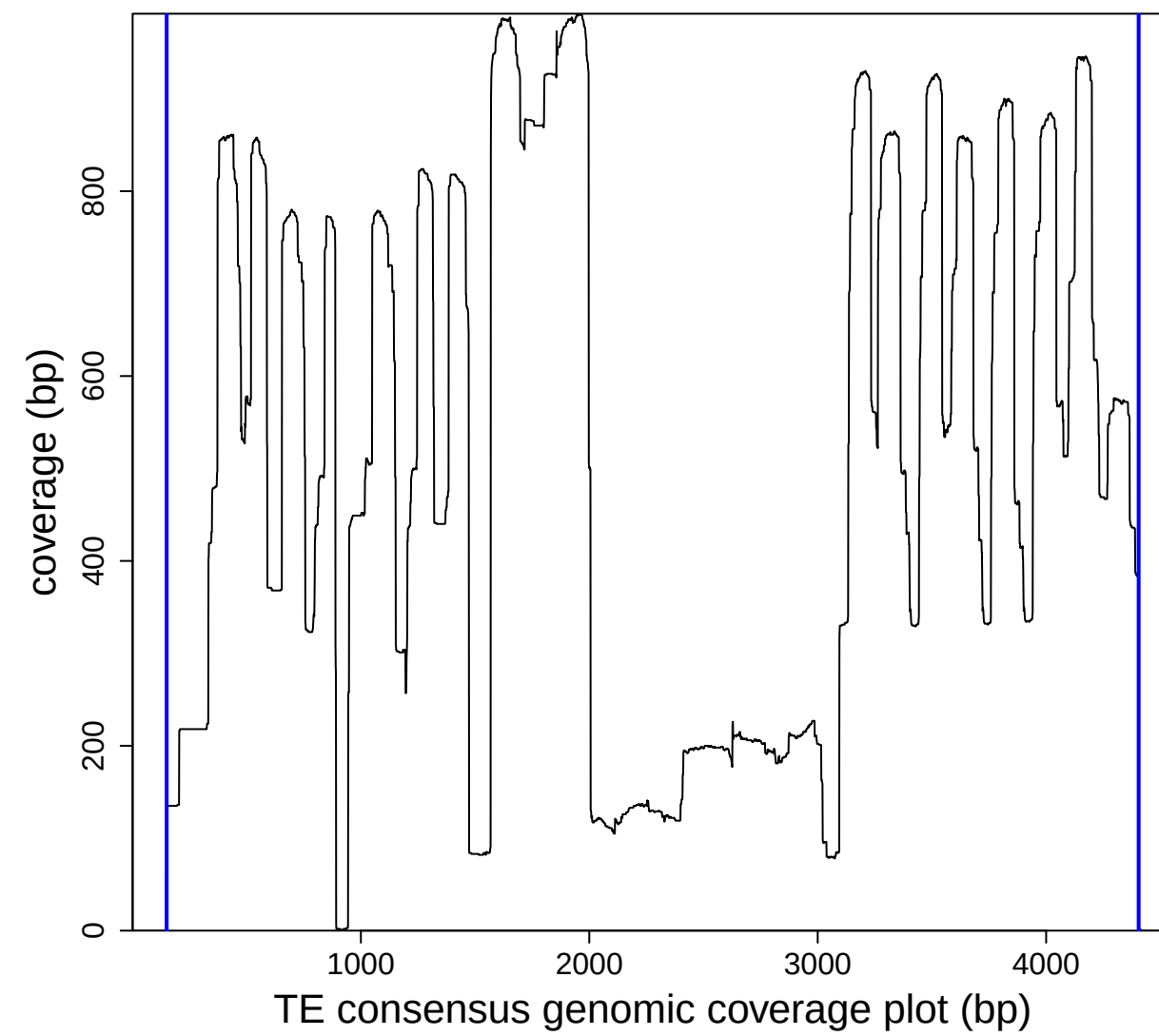
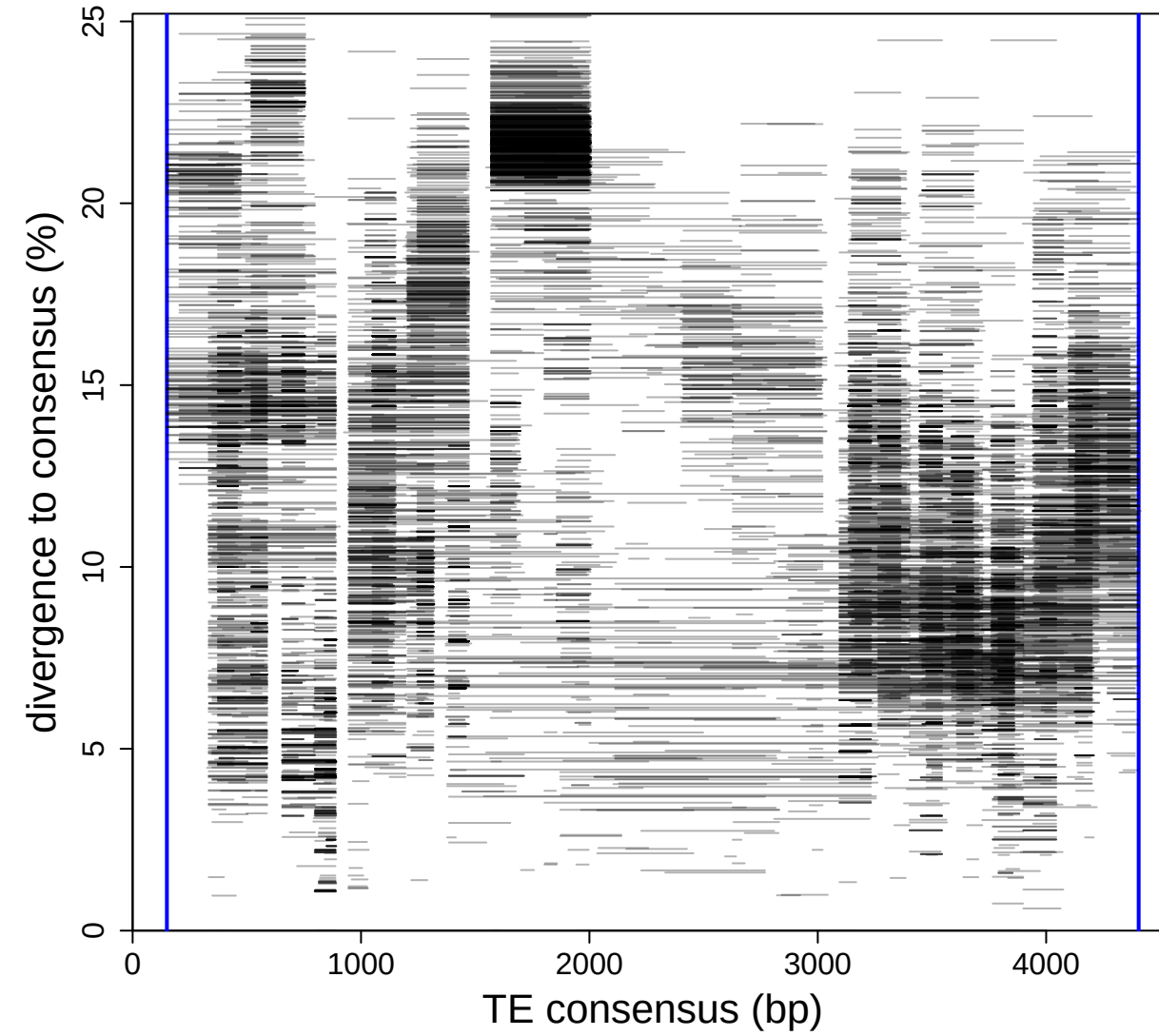
g_1.bed fm_1.bed 0 0 n.bed g_1.bed fm_2.bed 0 0 bcln.fa ε
size: 4074bp; fragments: 10196; full length: 0 (>=3666.6bp)

After TEtrimmer 4074 bp



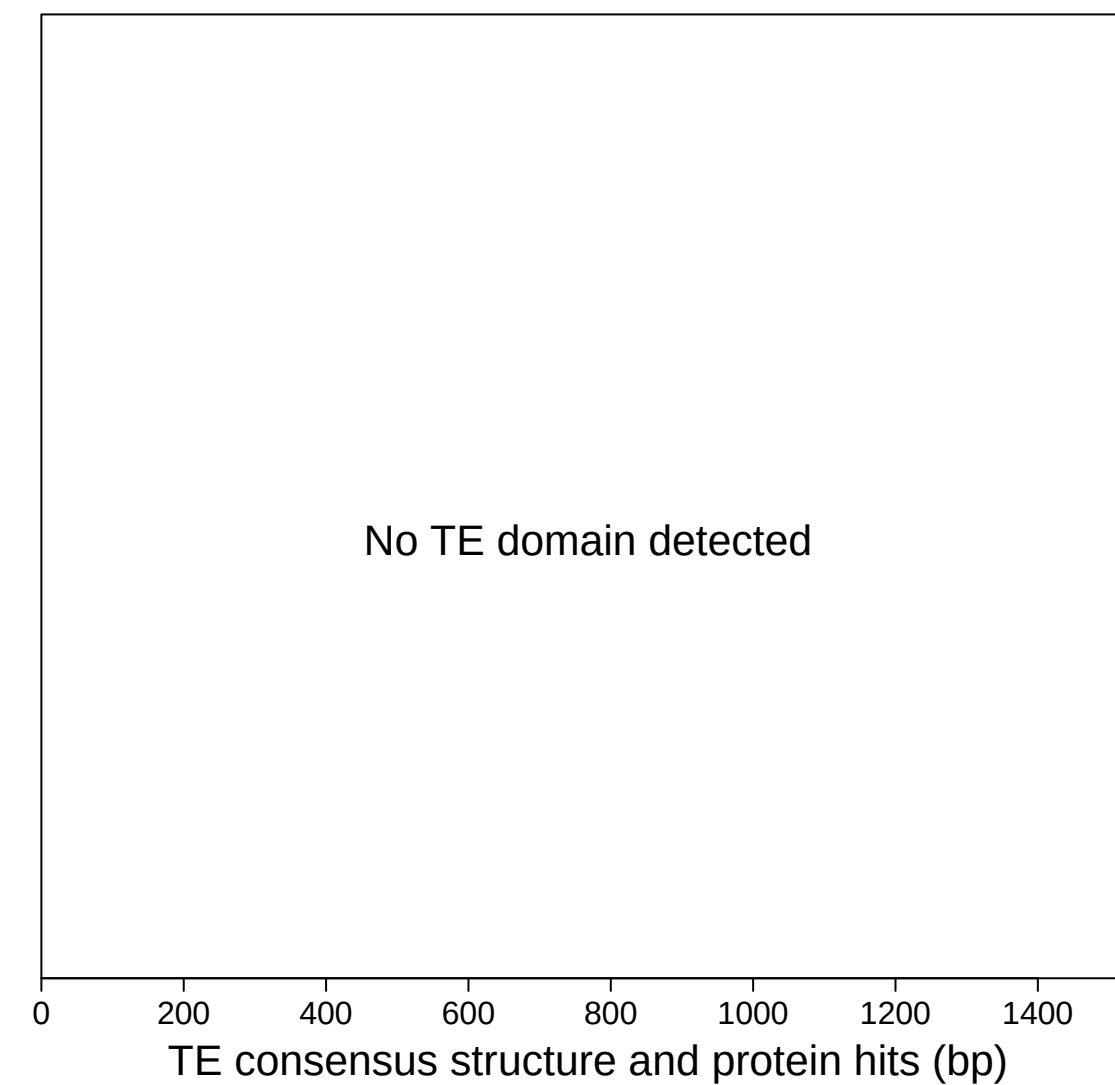
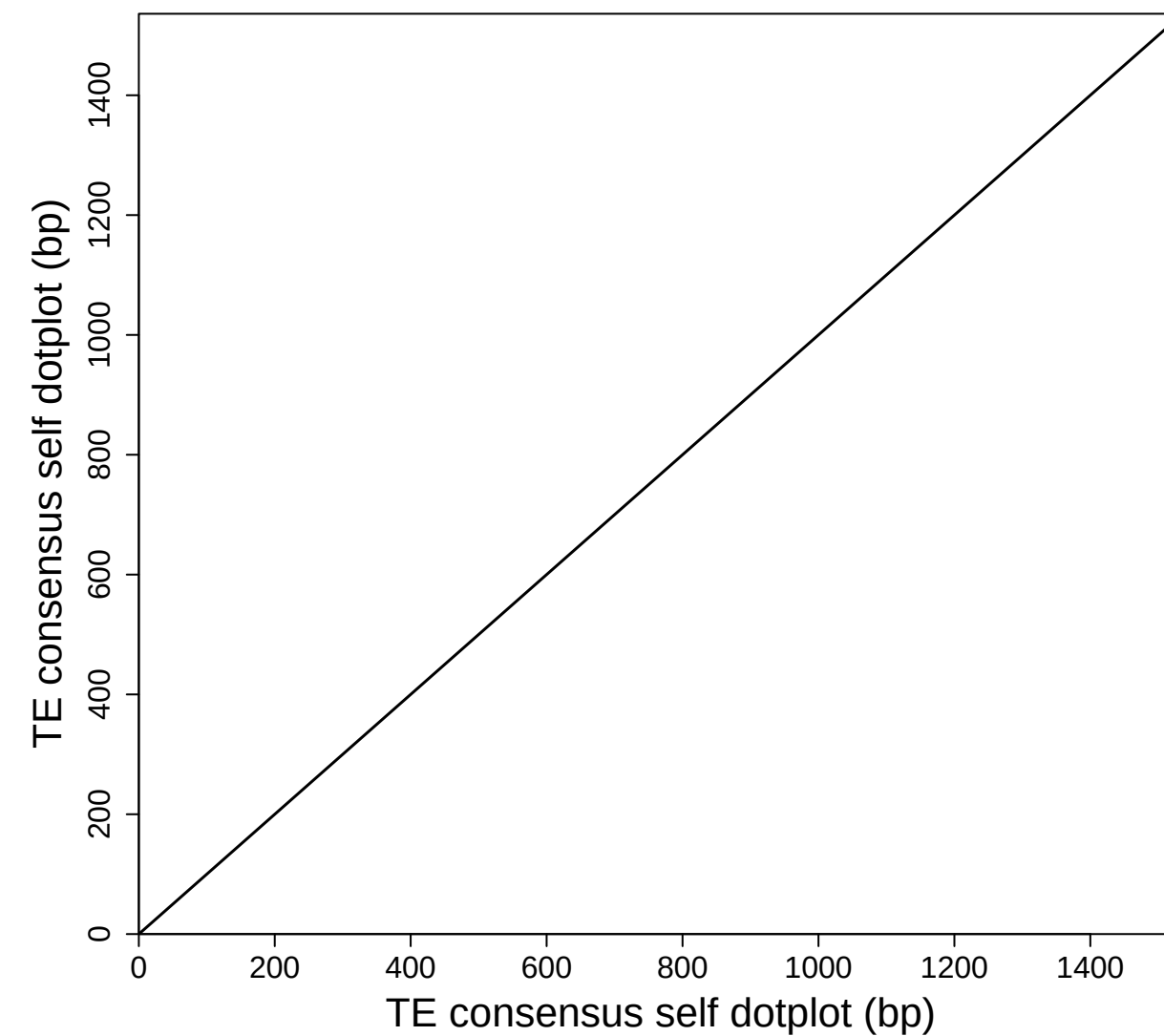
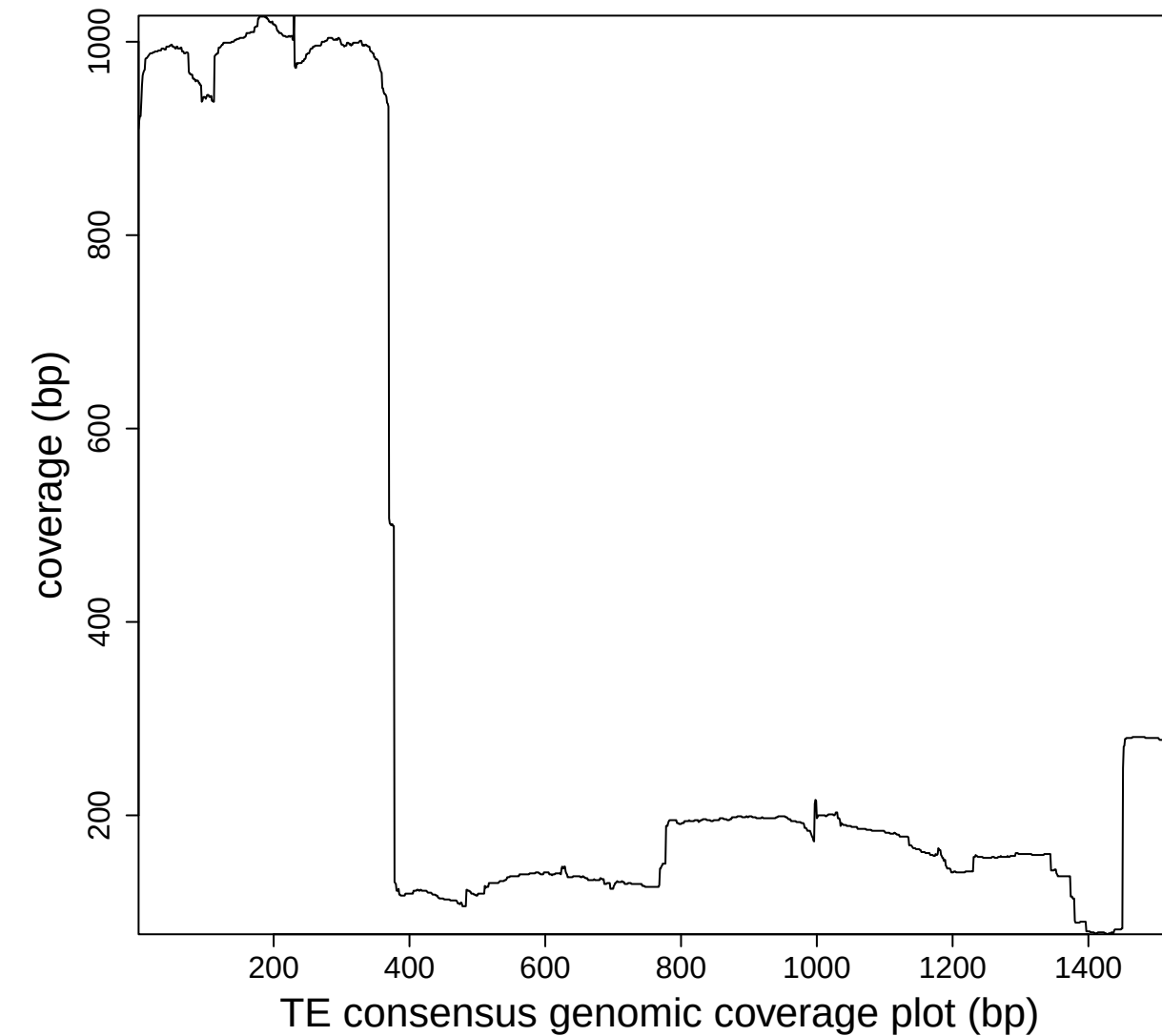
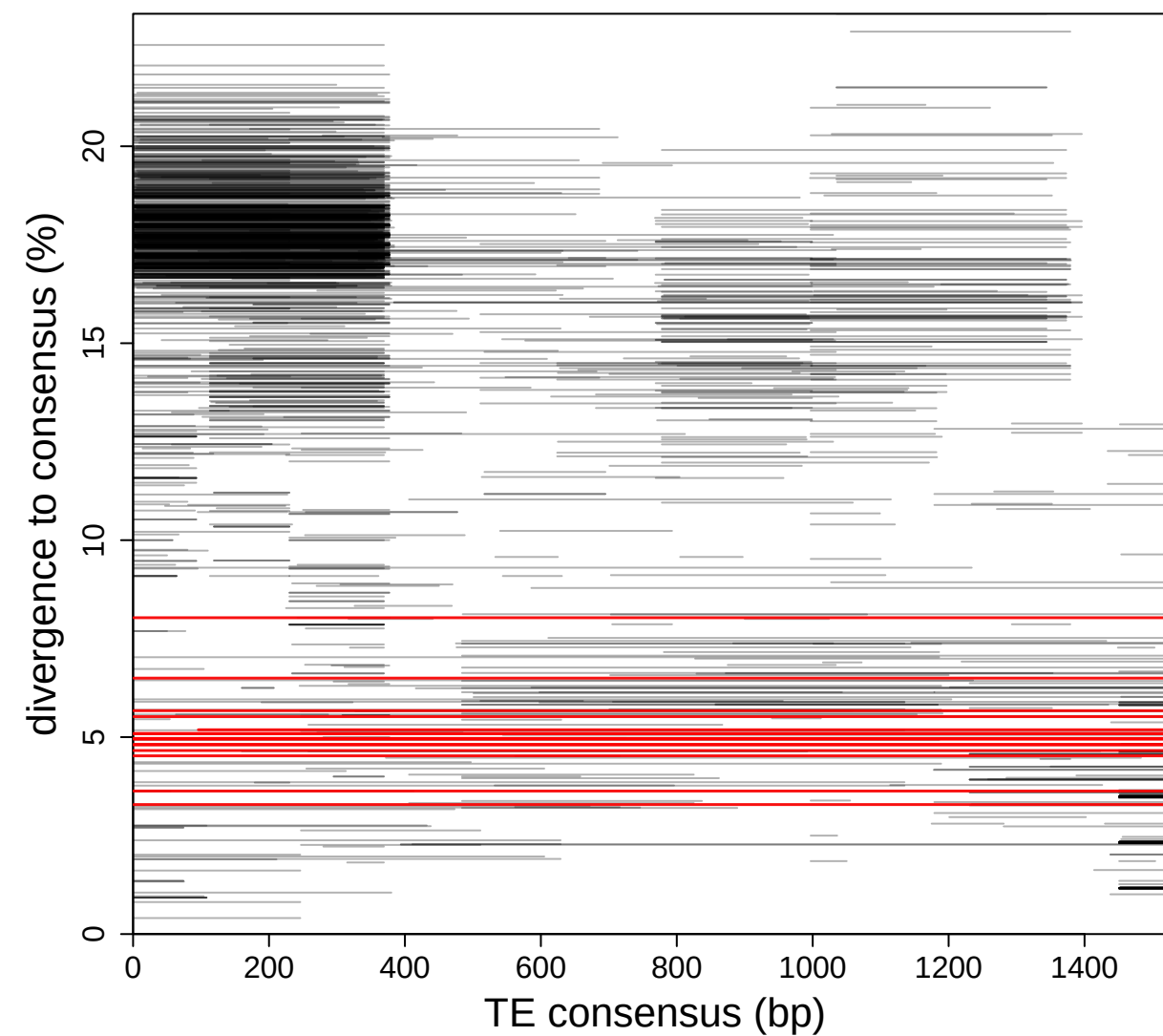
f.bed_g_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bc
size: 4555bp; fragments: 9610; full length: 0 (>=4099.5bp)

After TEtrimmer Extended plot Blue lines are boundaries

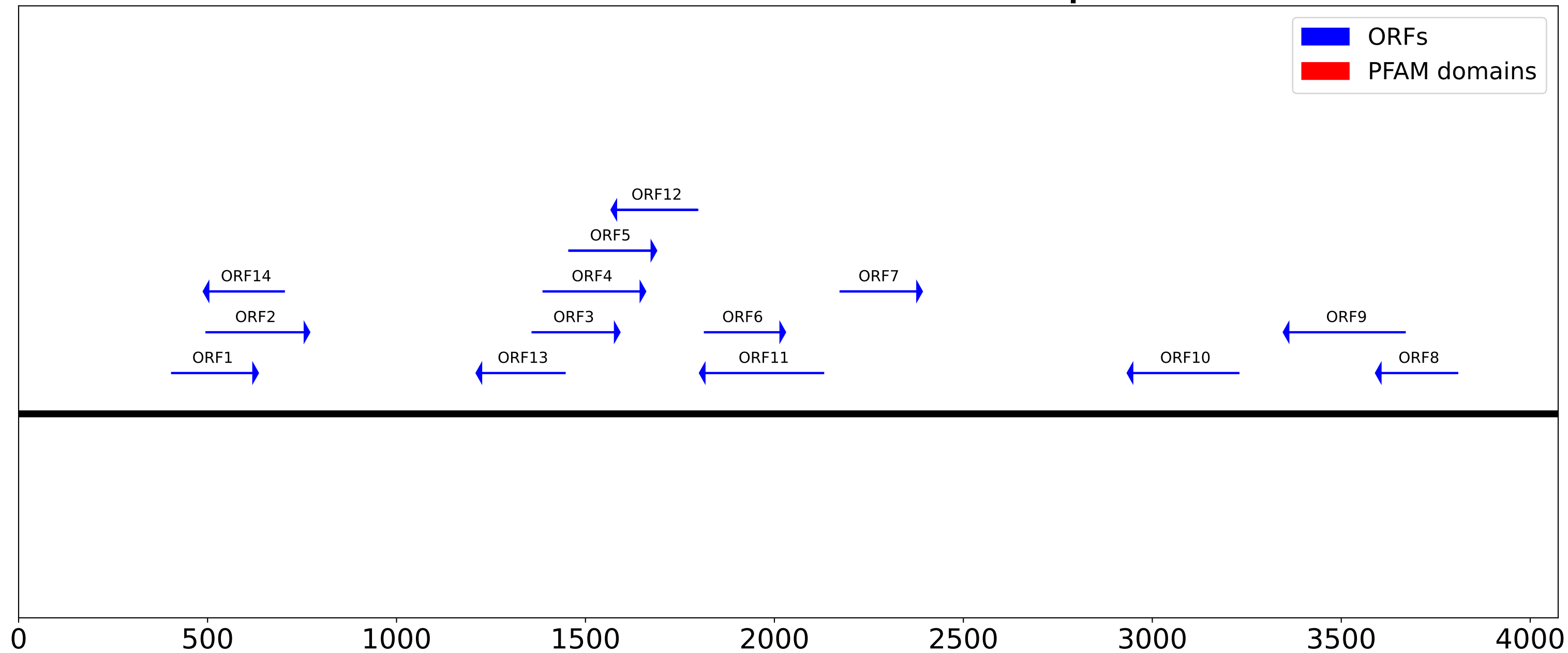


TE: rnd_5_family_5626
size: 1536bp; fragments: 1690; full length: 16 (≥ 1382.4 bp)

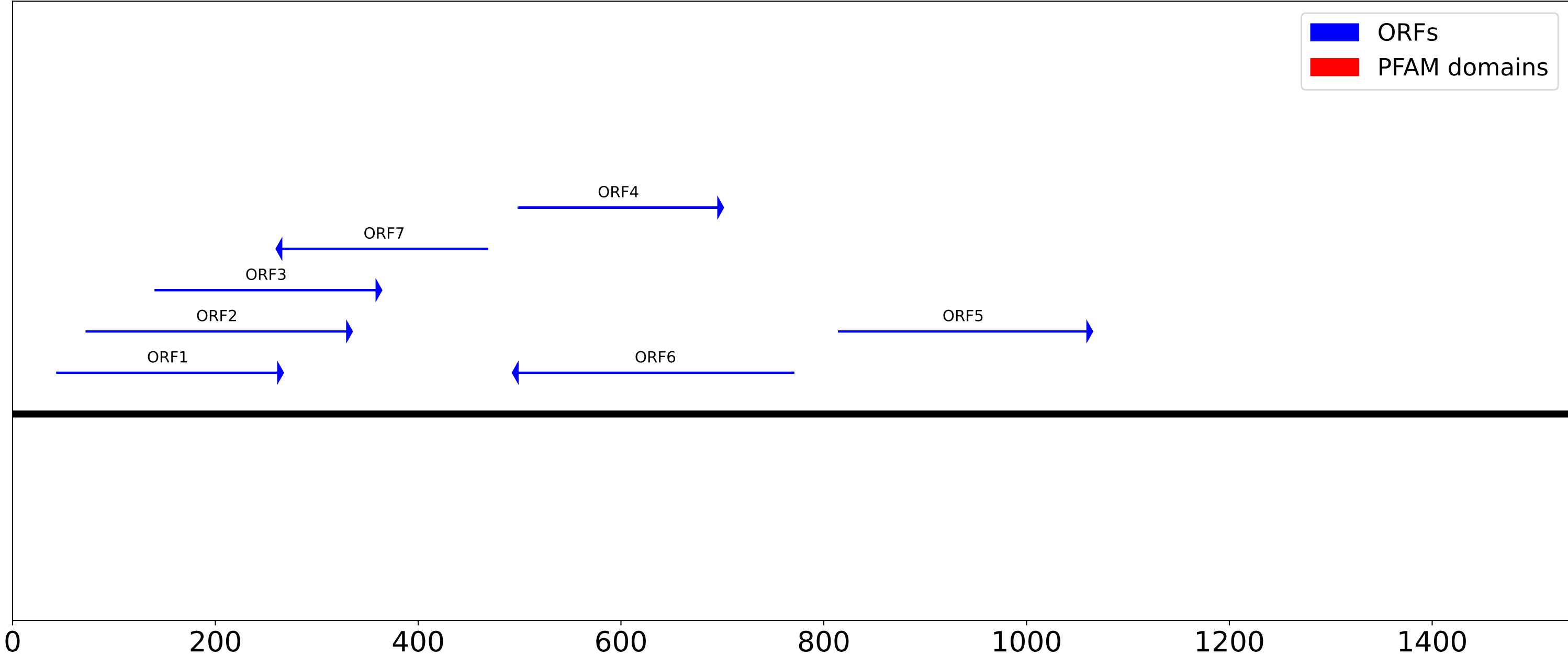
Before TEtrimmer 1536 bp



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)

