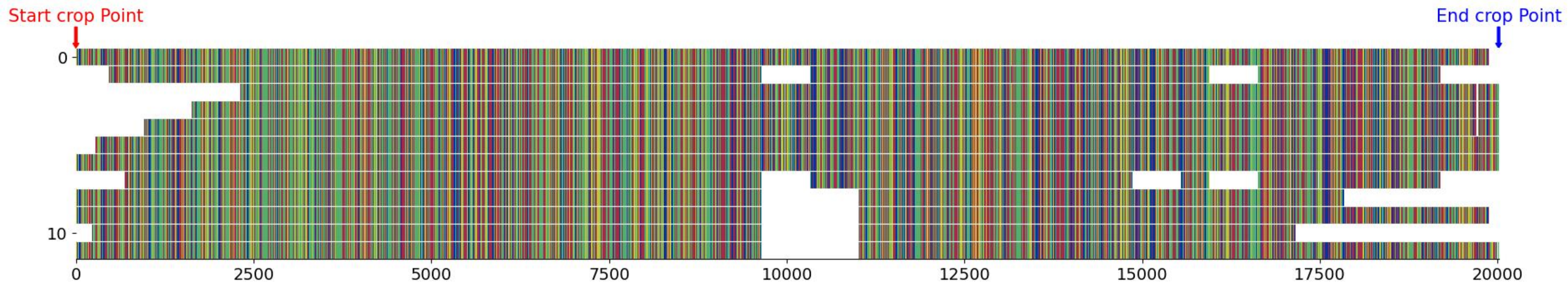




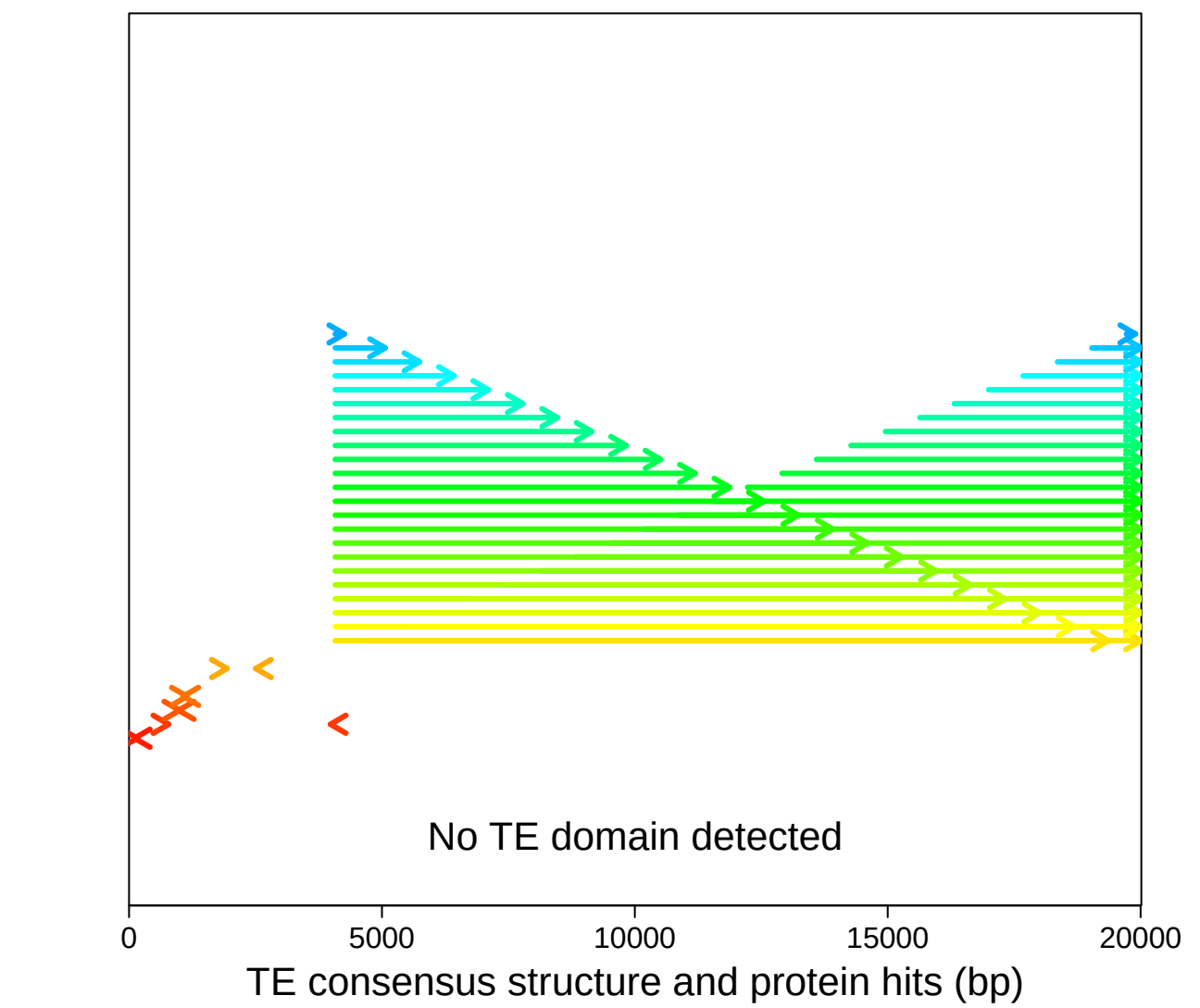
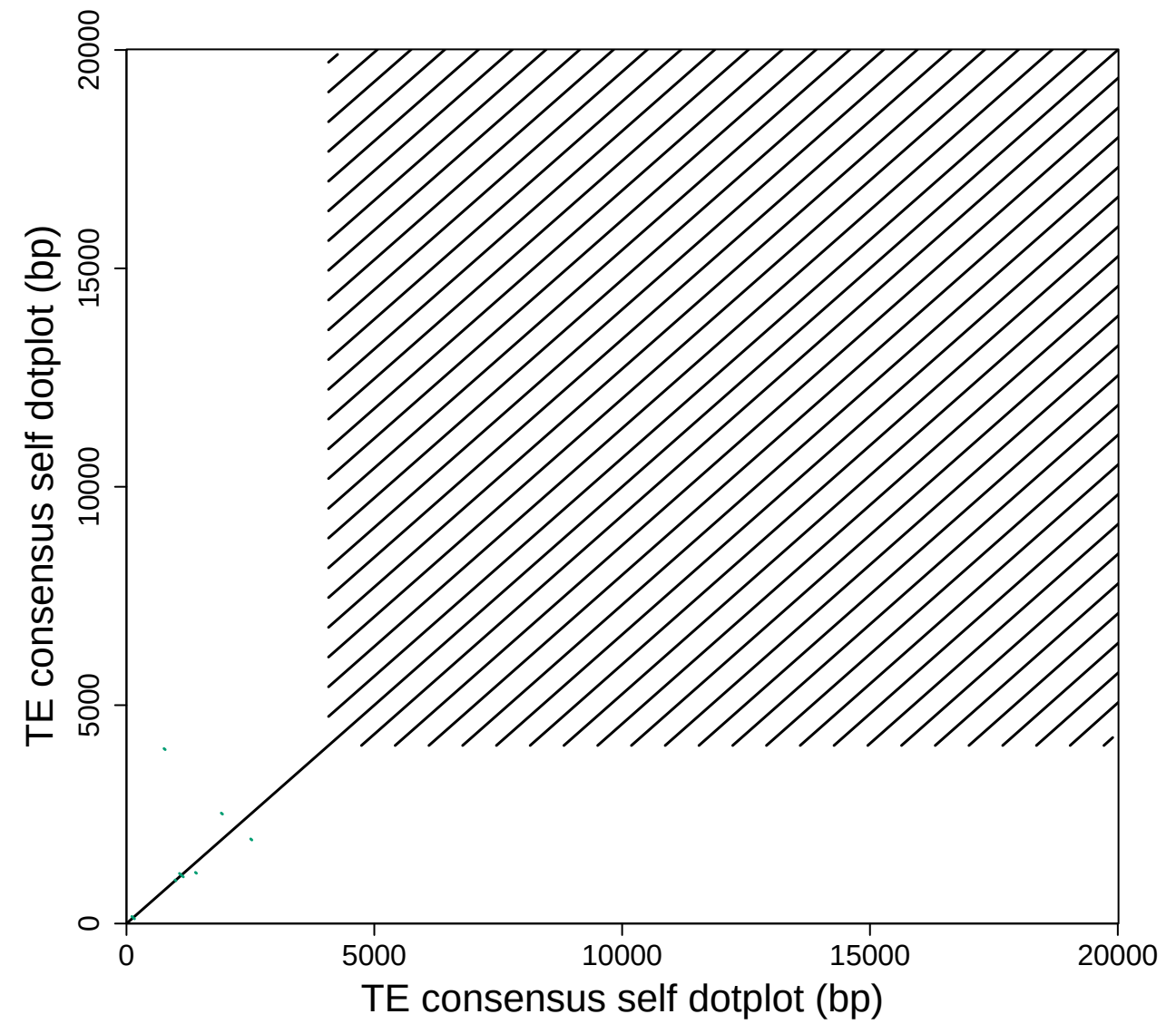
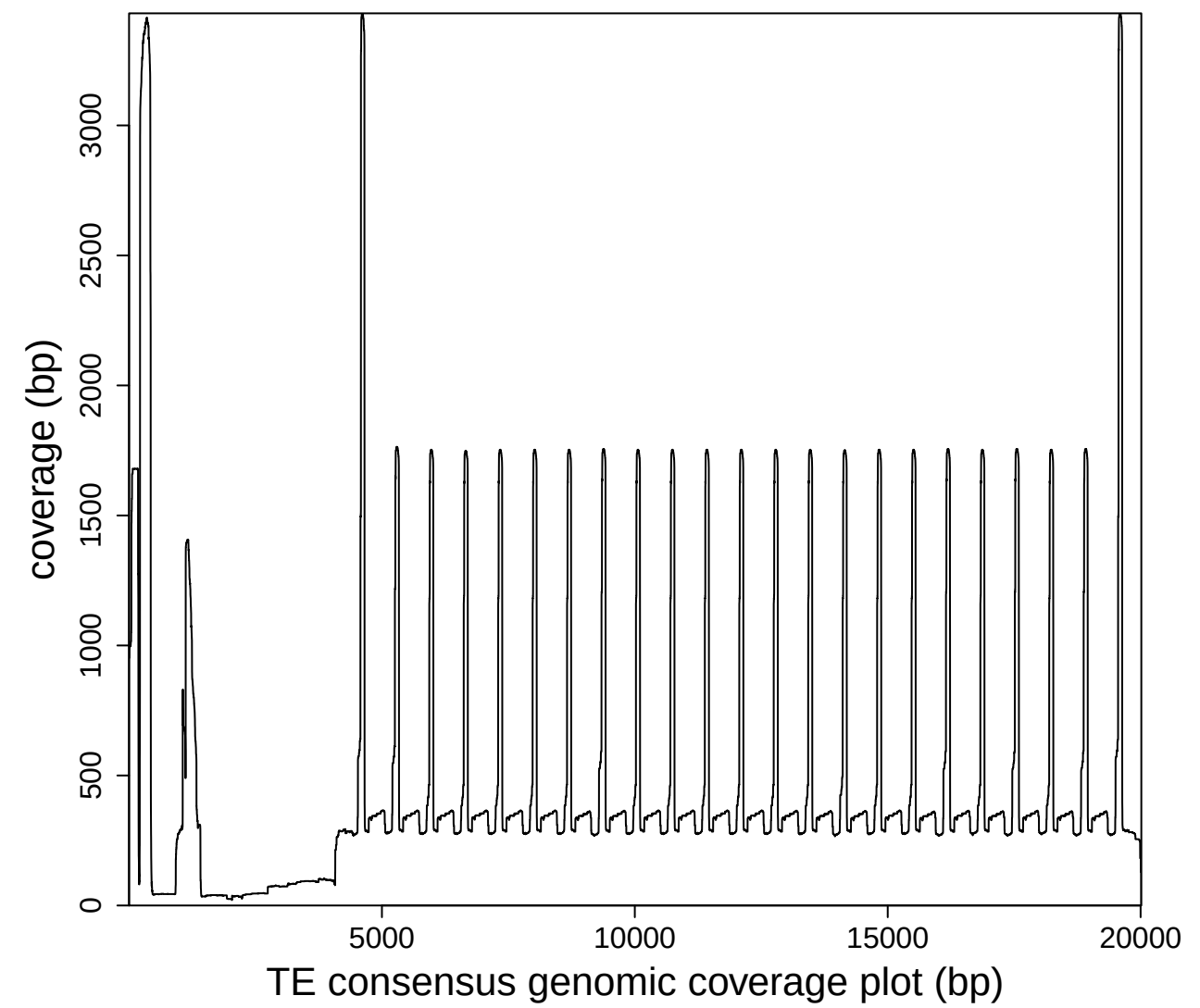
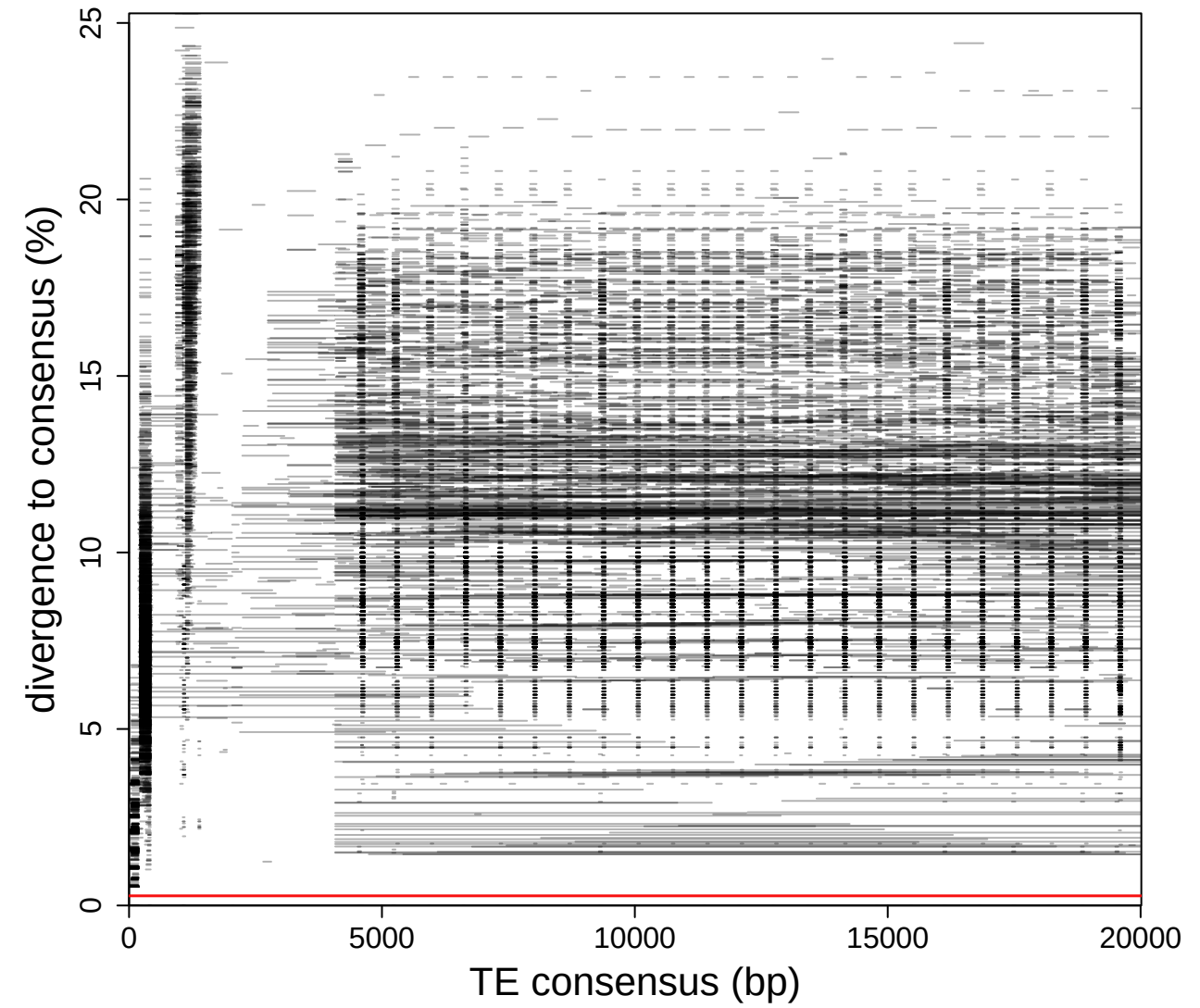
MSA length = 20014

A G GA G GACAAA CGAA GACAAAAA AAAACAACCAAAA - - - - -



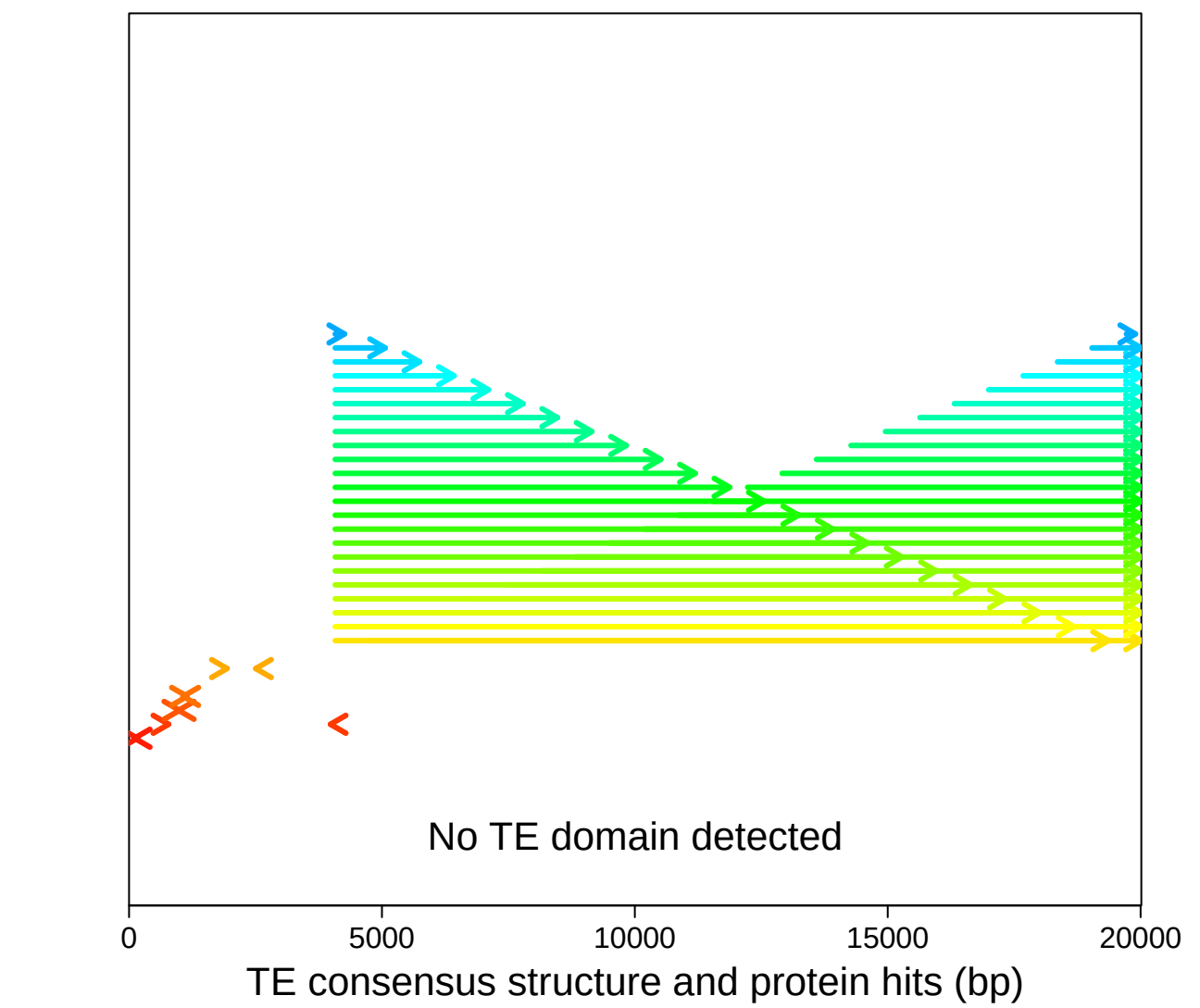
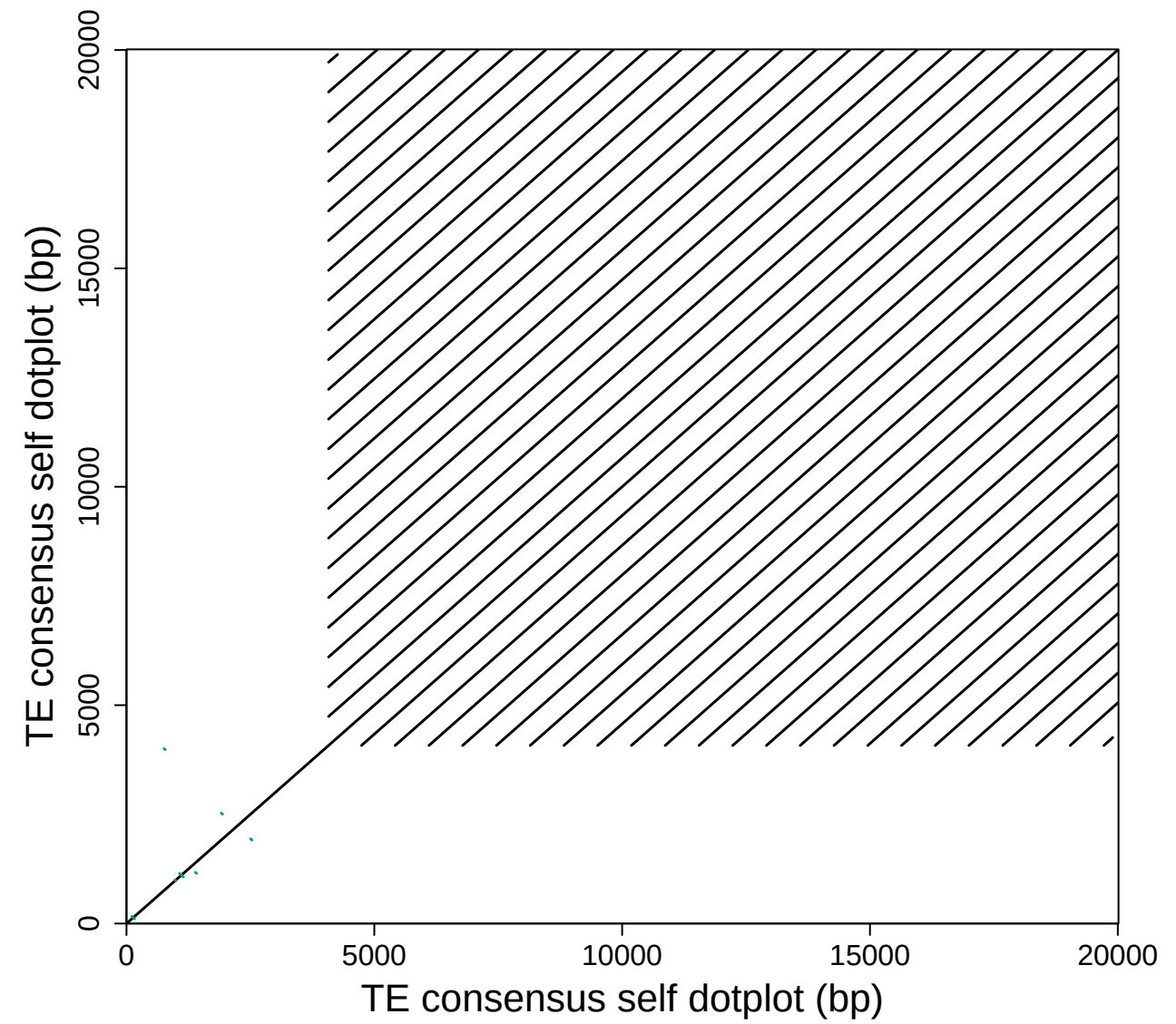
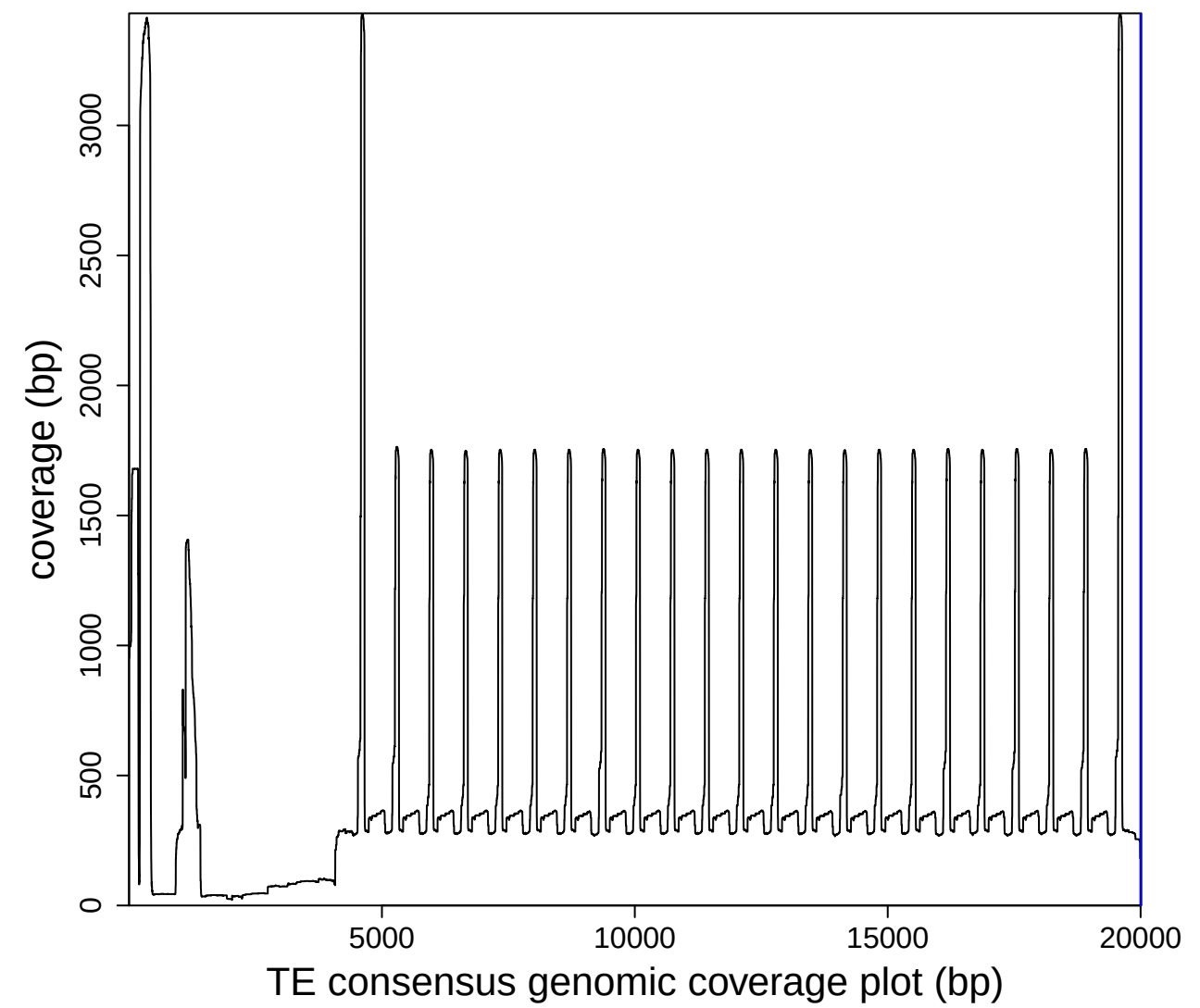
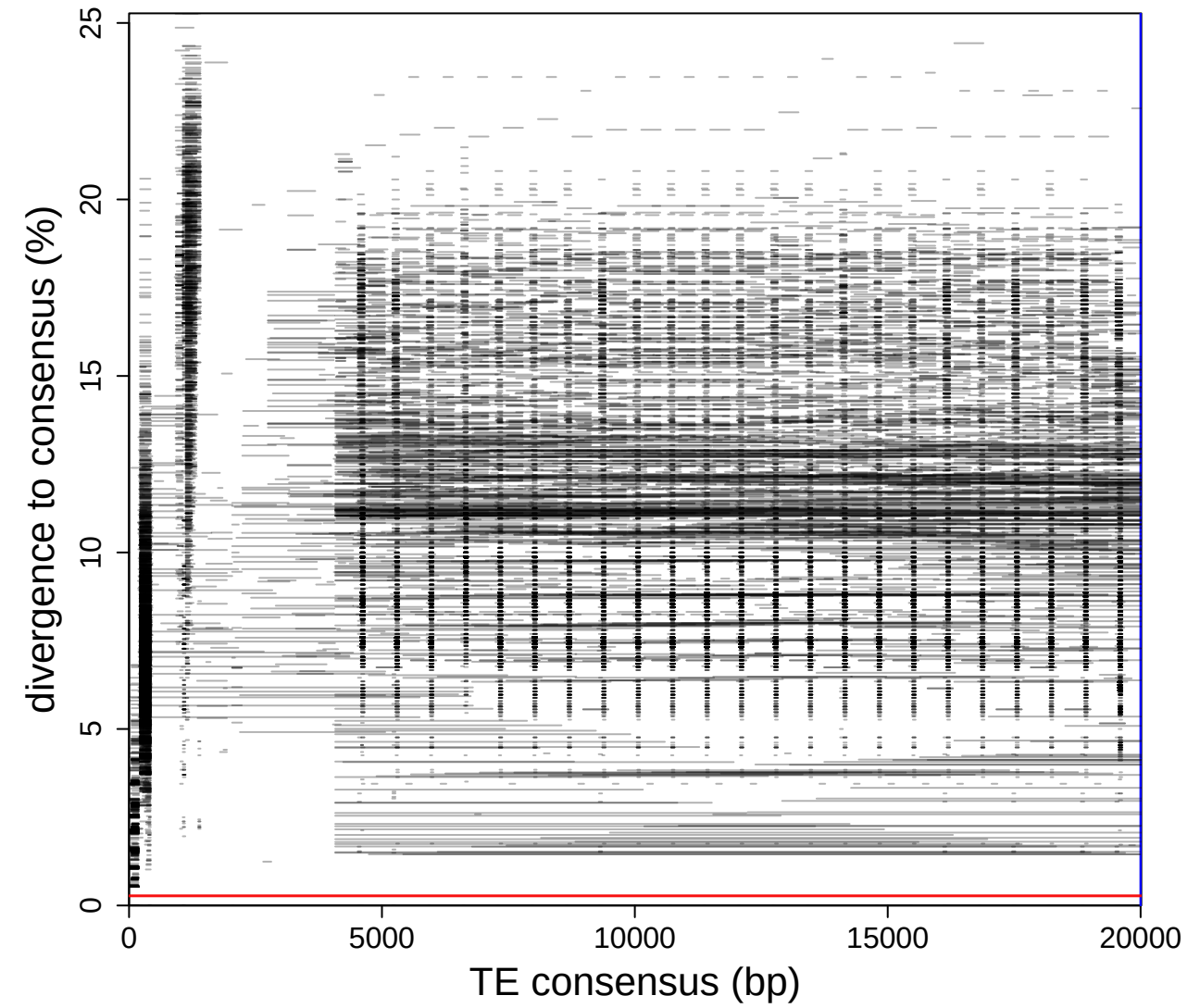
g_1.bed fm_1.bed 0 0 n.bed g_1.bed fm_2.bed 0 0 bcln.fa &
size: 20014bp; fragments: 48291; full length: 1 (>=18012.6bp)

After TEtrimmer 20014 bp



f.bed_g_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bc
size: 20014bp; fragments: 48291; full length: 1 (>=18012.6bp)

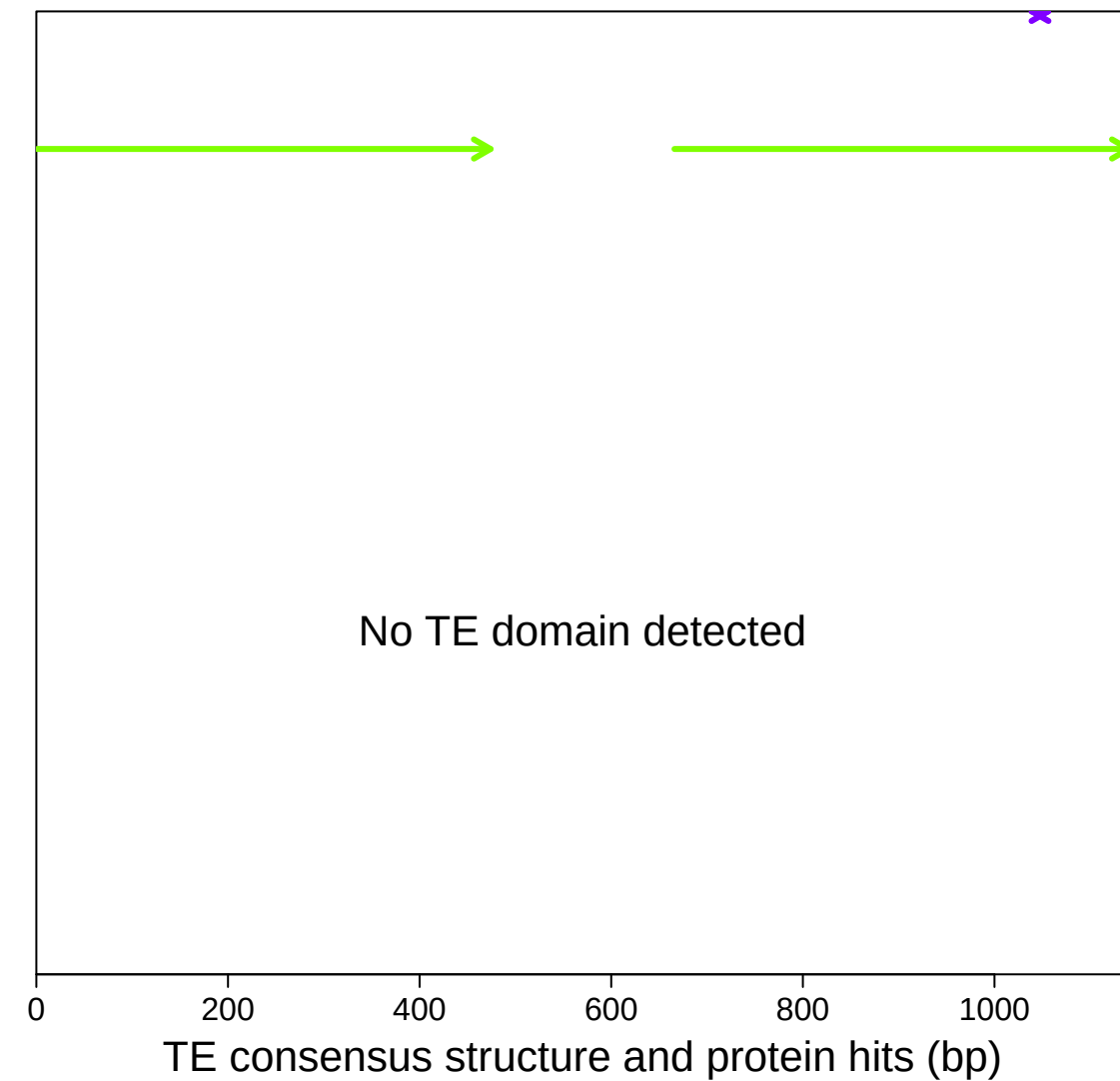
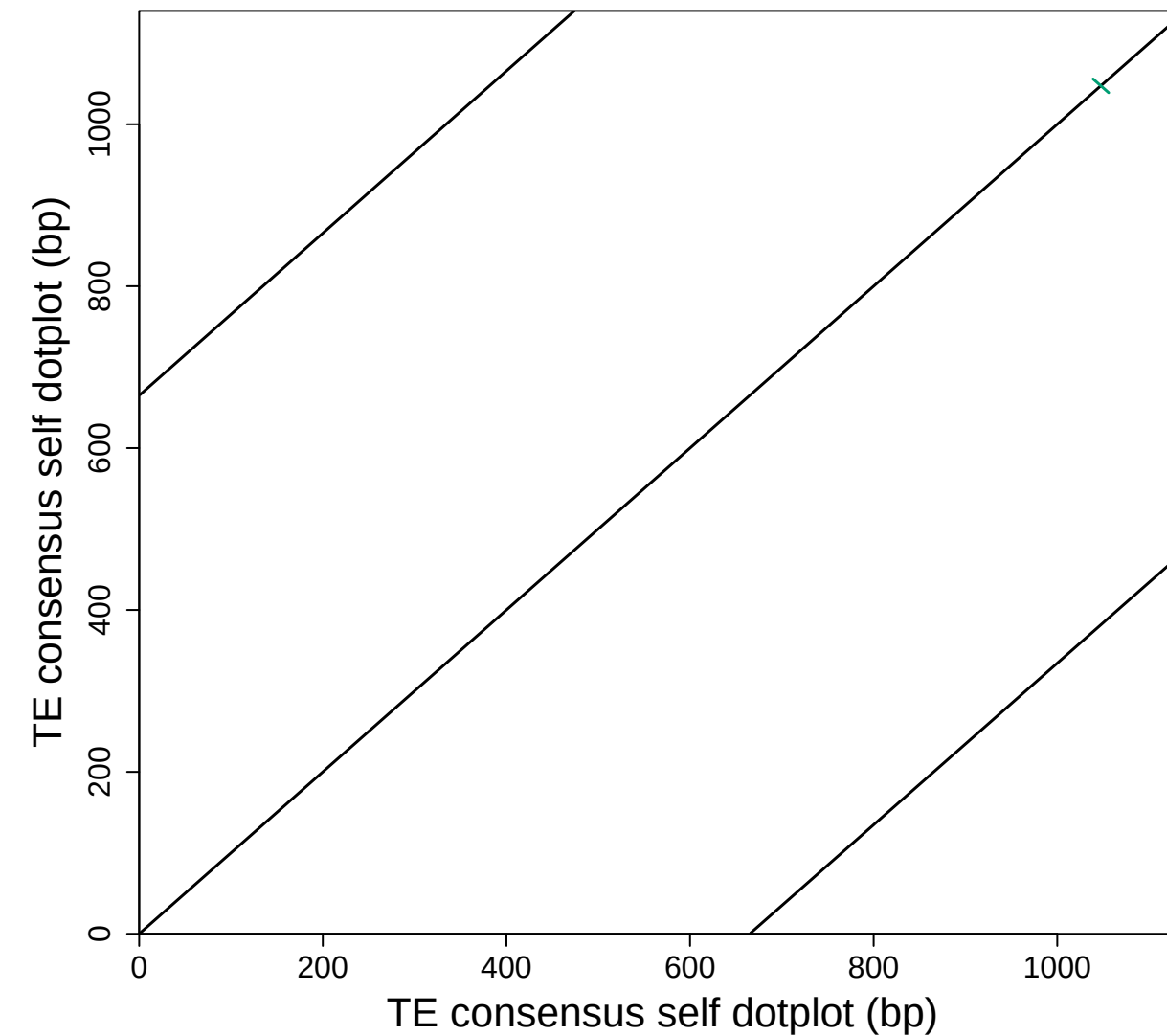
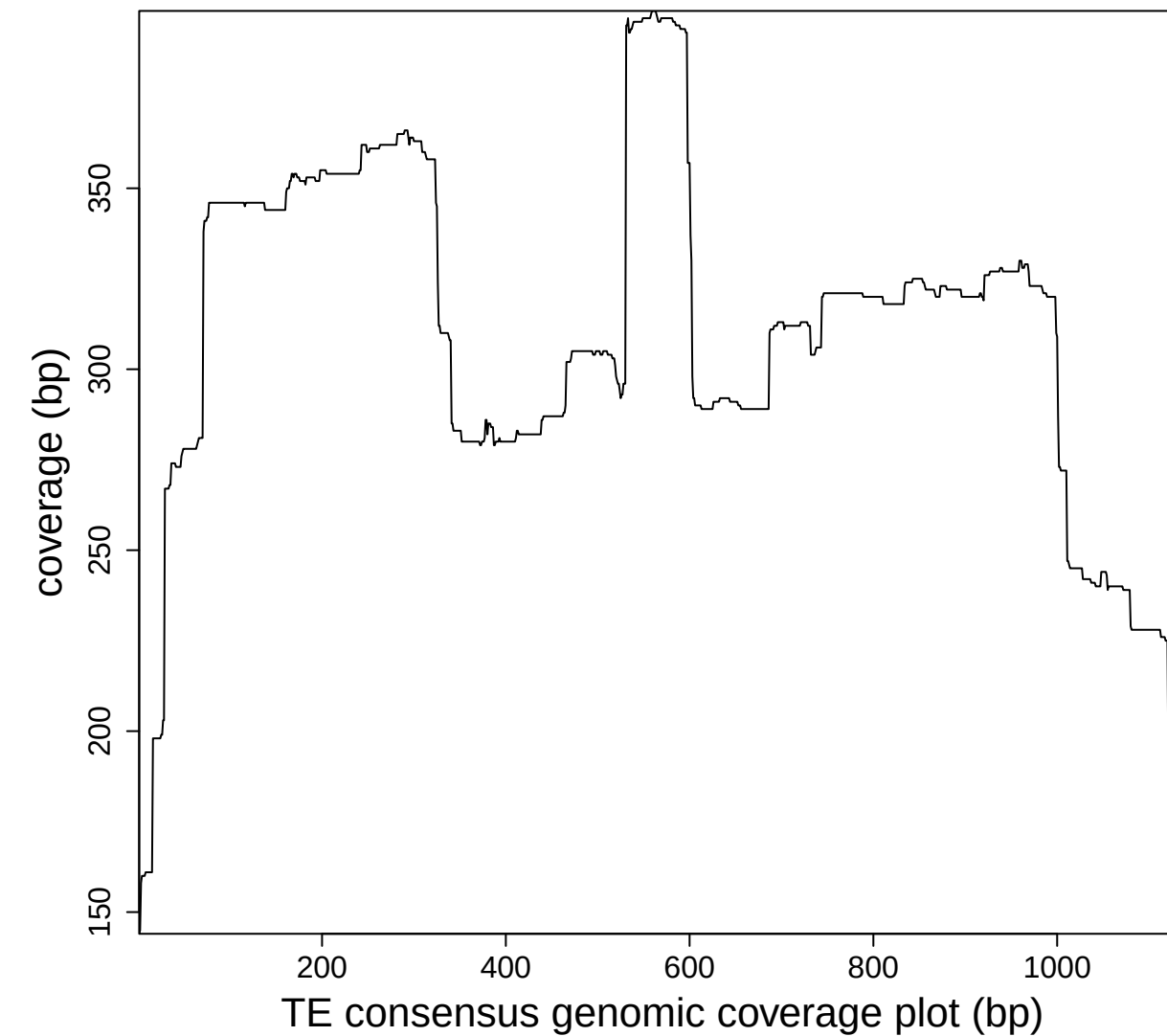
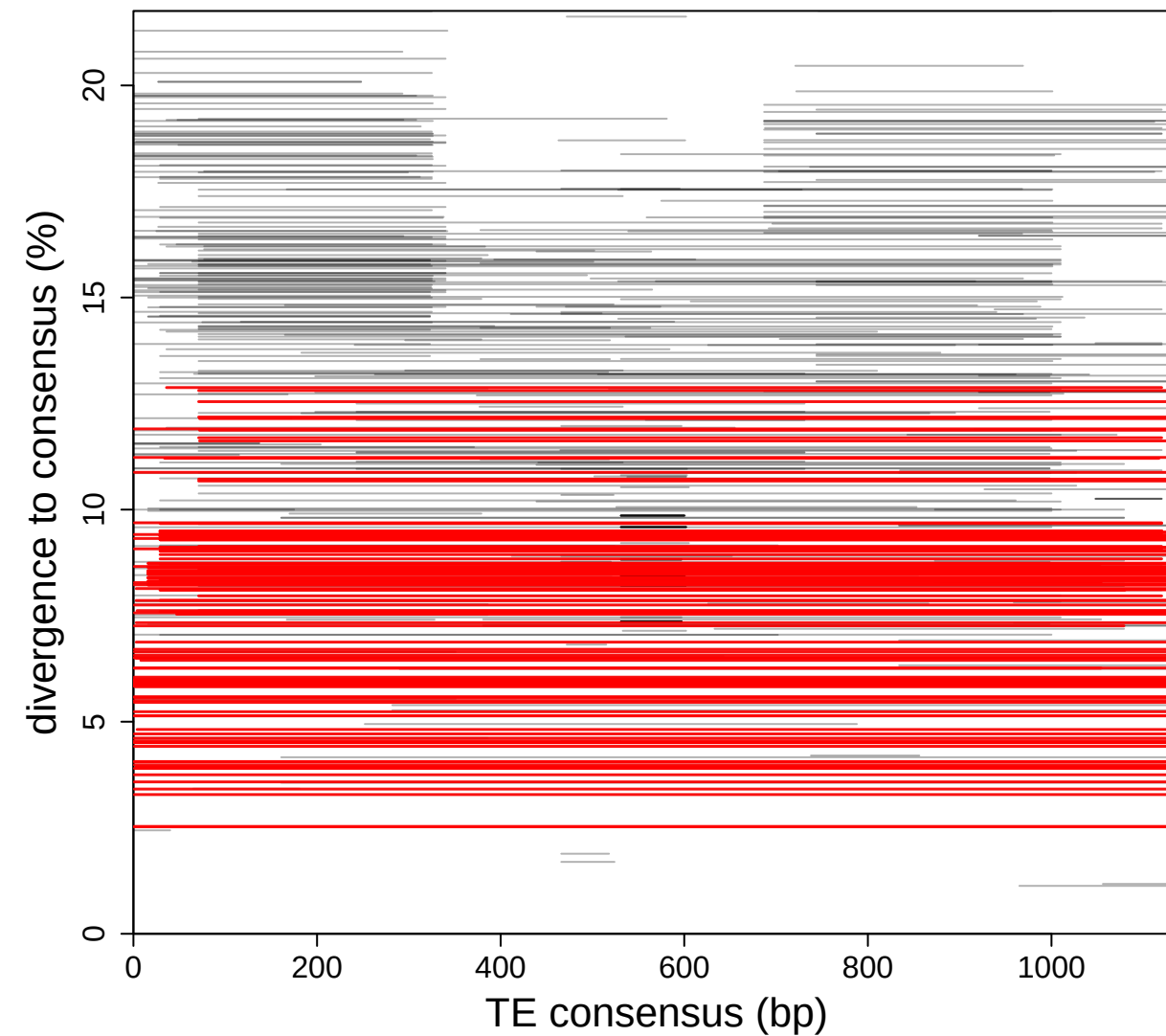
After TEtrimmer Extended plot Blue lines are boundaries



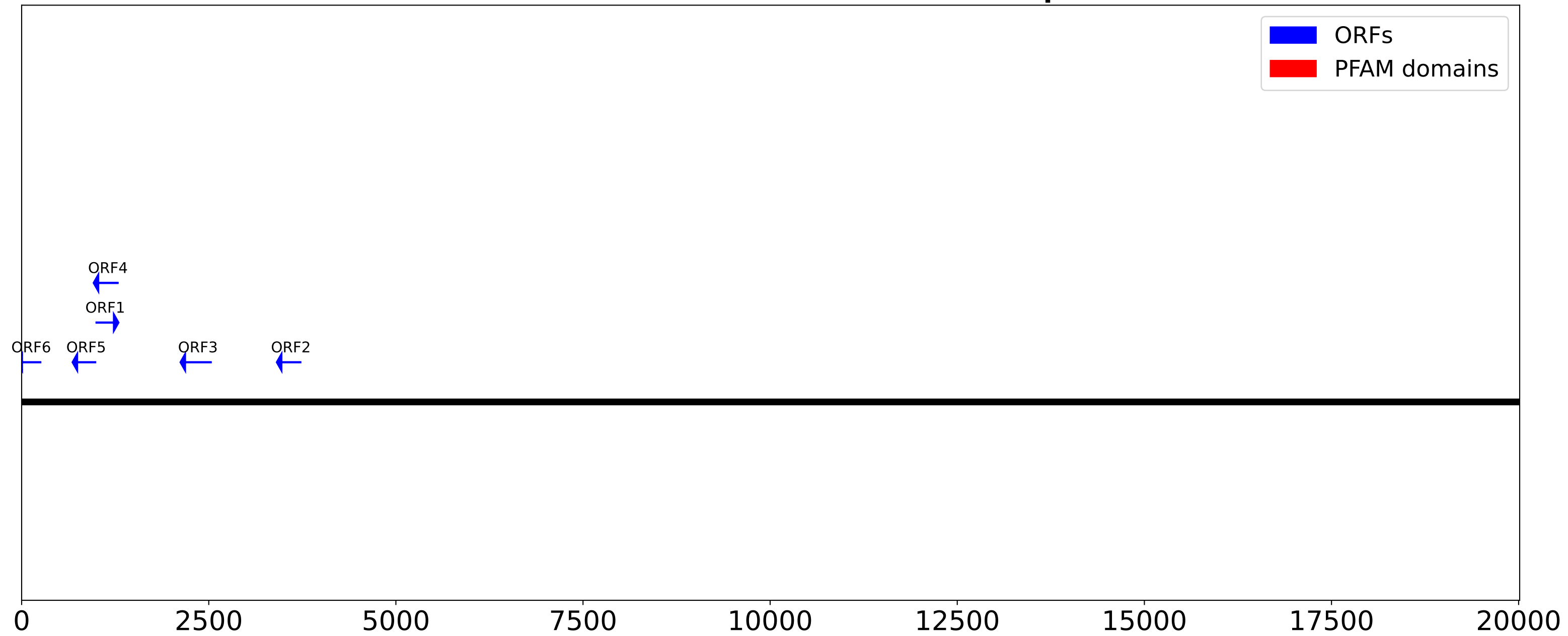
Before TEtrimmer 1140 bp

TE: rnd_4_family_7213

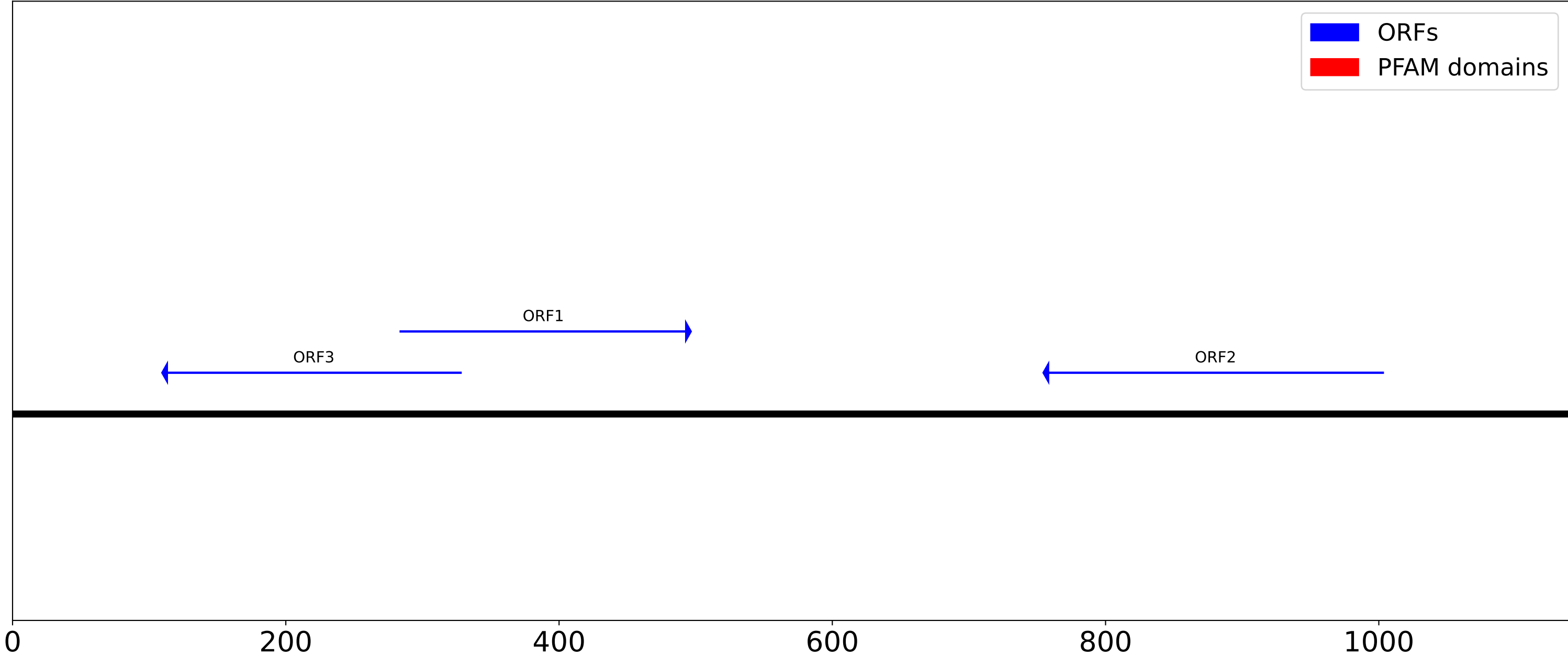
size: 1140bp; fragments: 614; full length: 157 (≥ 1026 bp)



After TEtrimmer ORF and PFAM domain plot



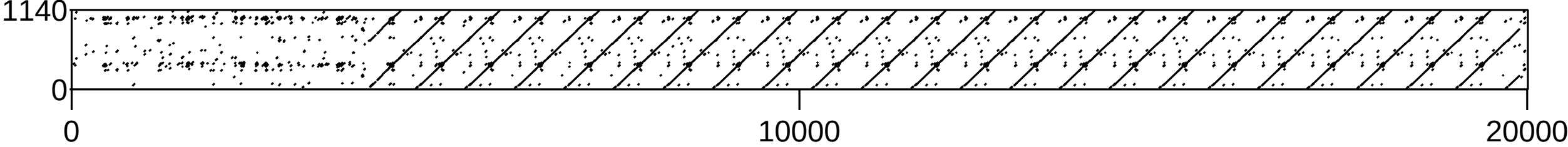
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)