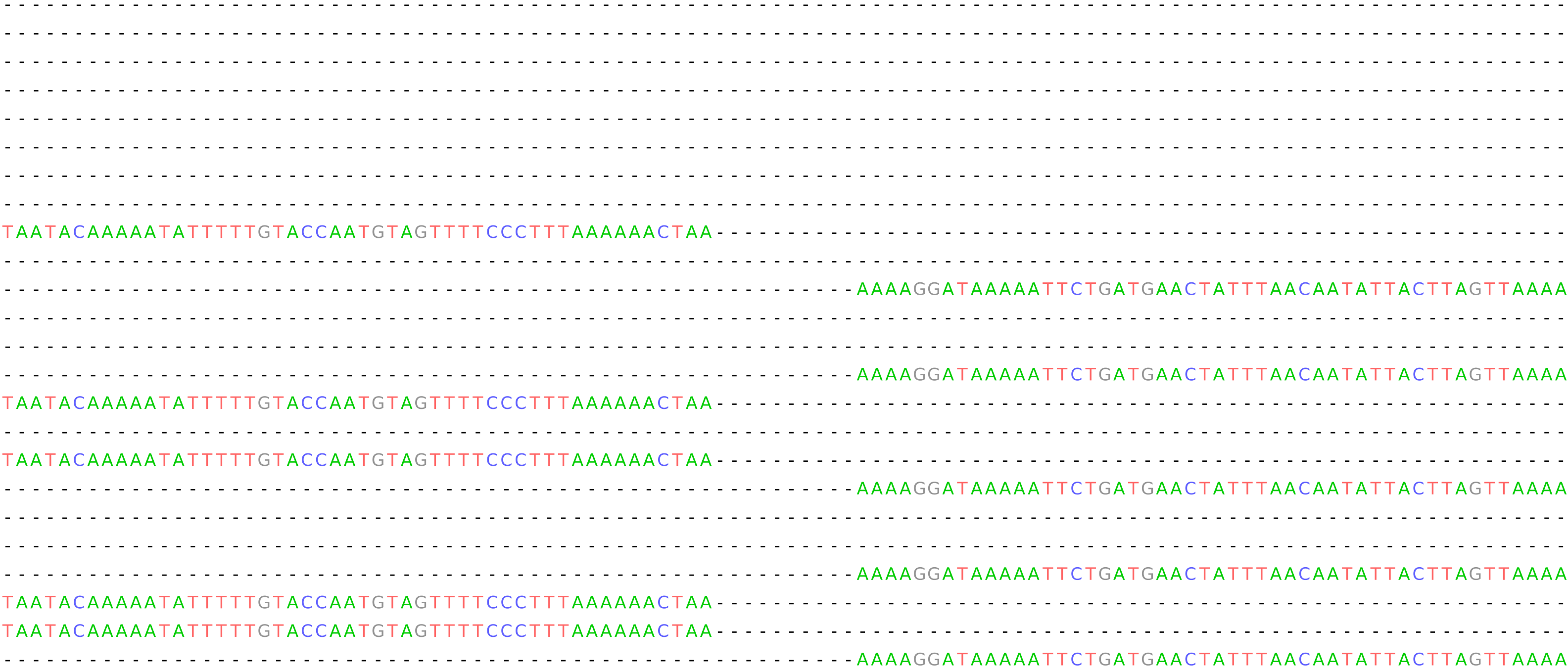
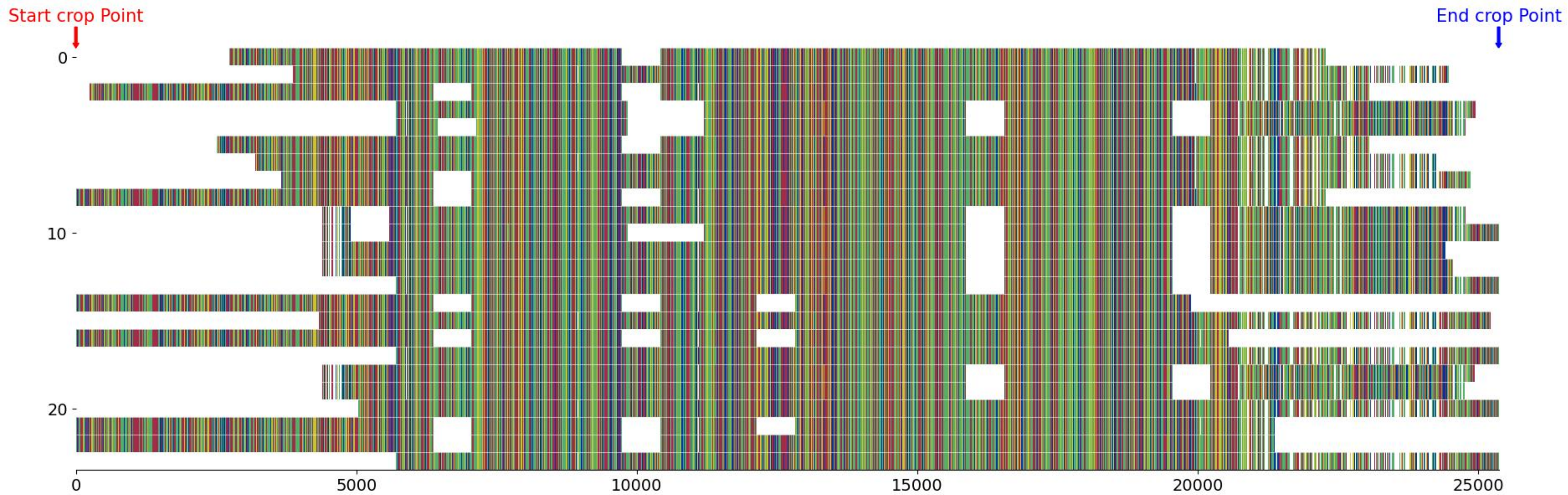


Start crop Point

End crop Point

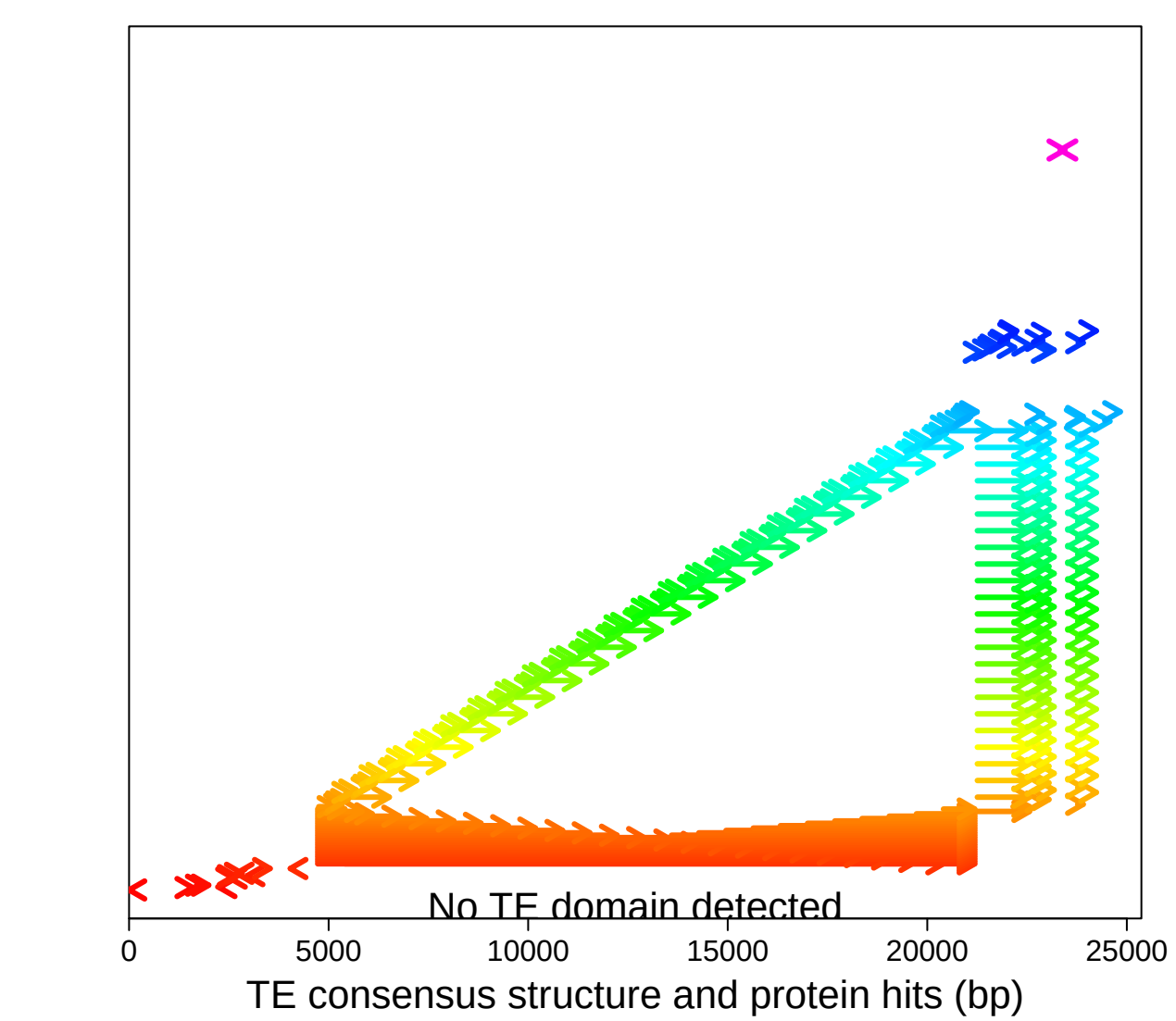
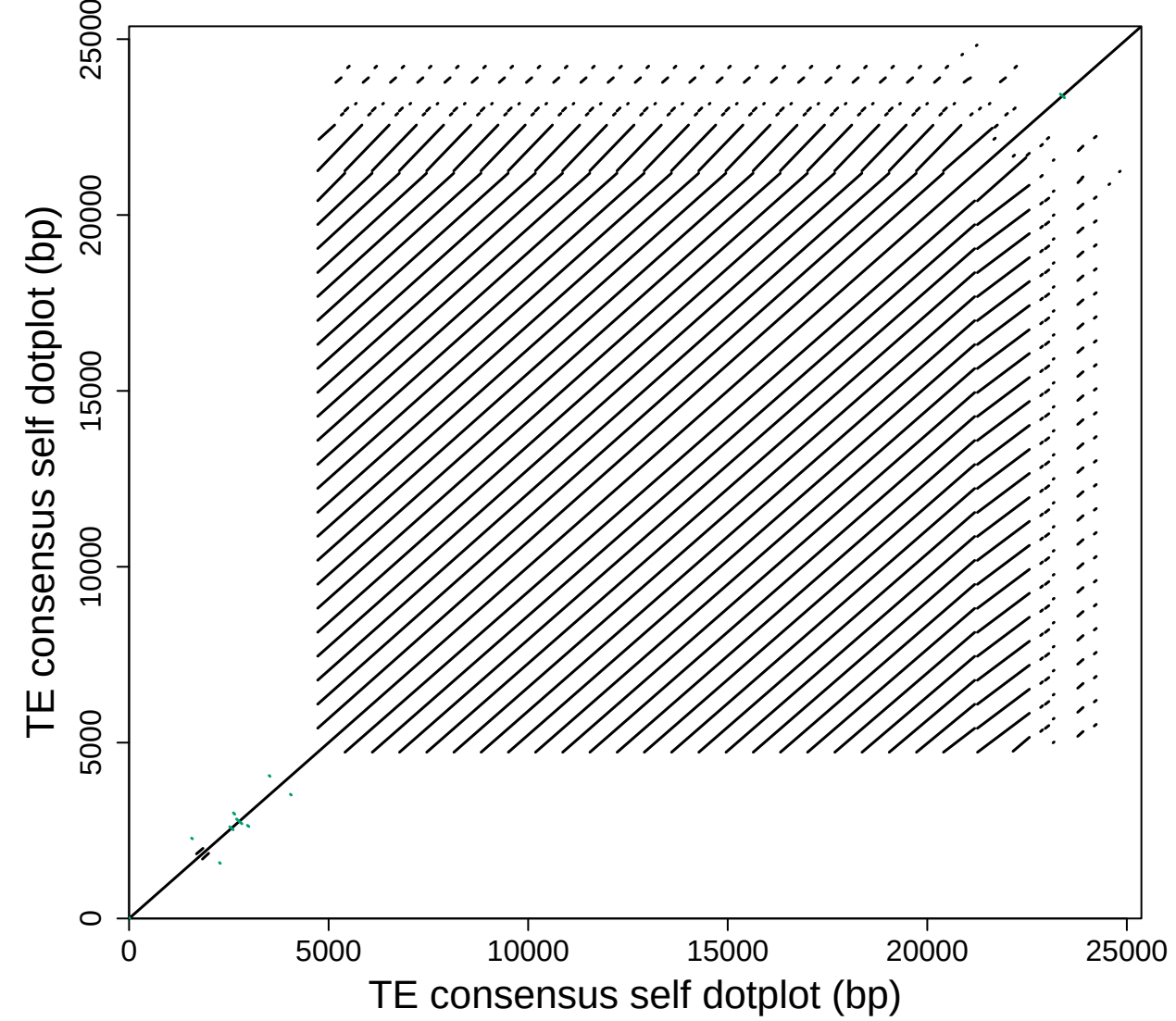
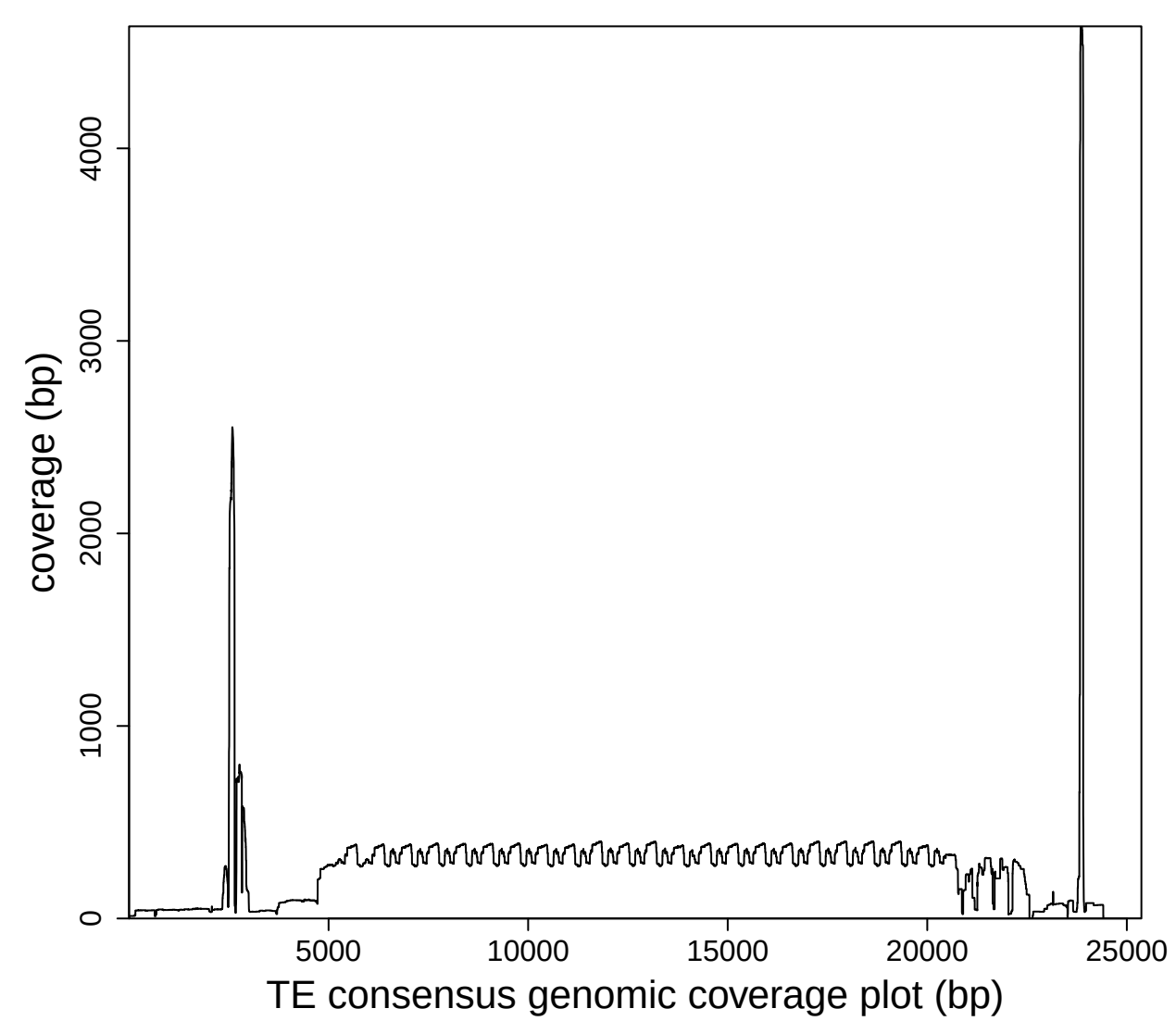
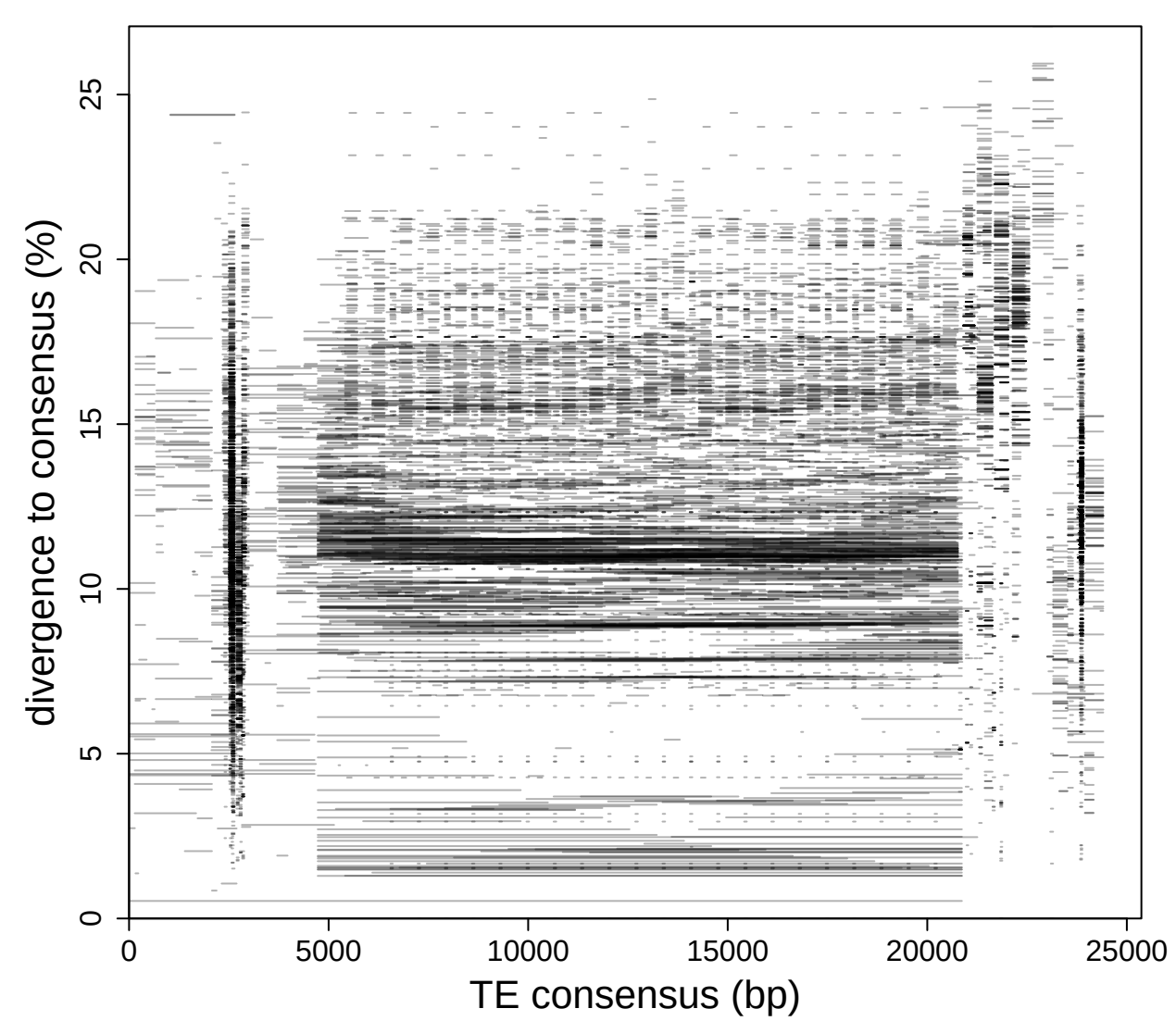
MSA length = 25365





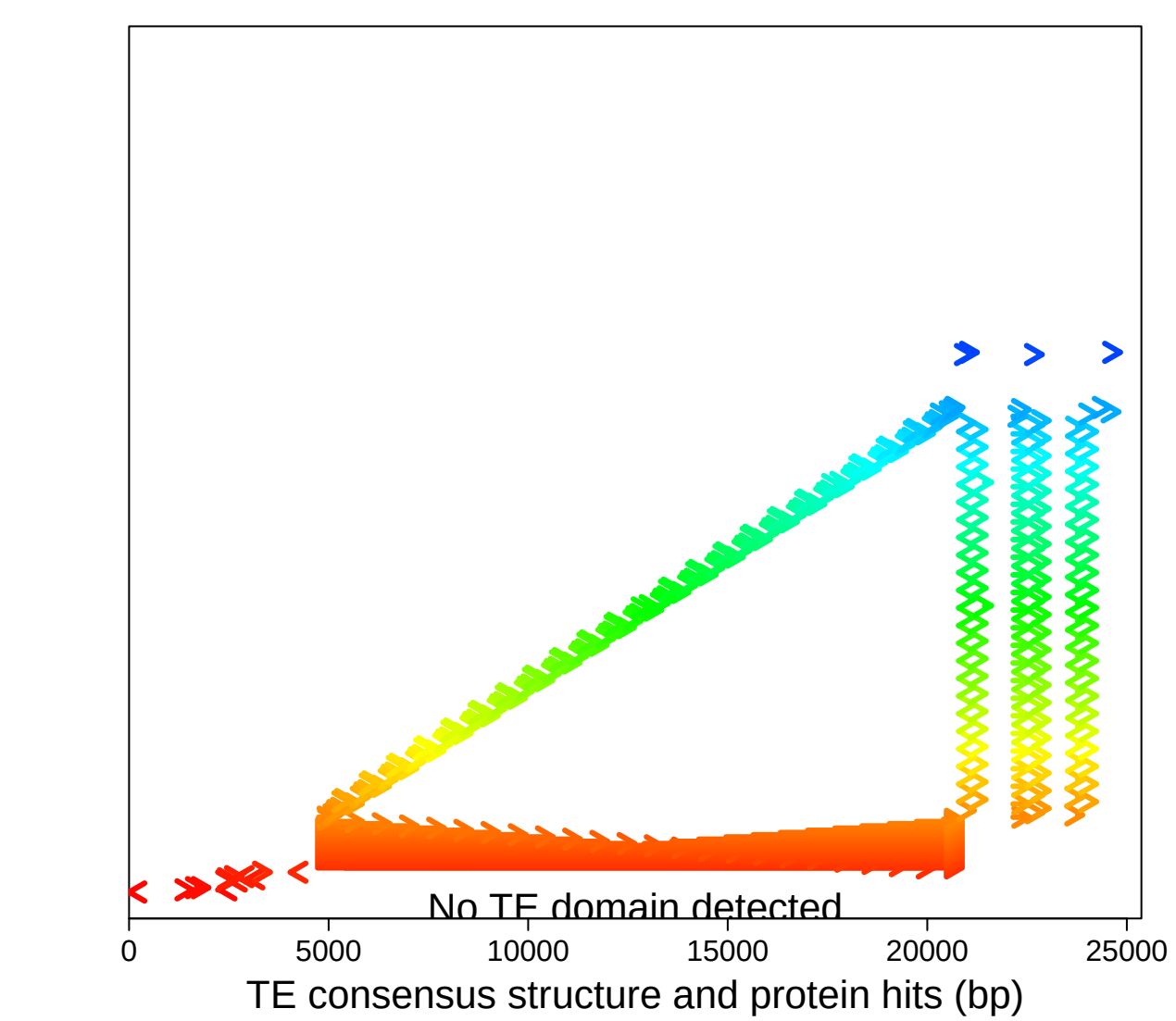
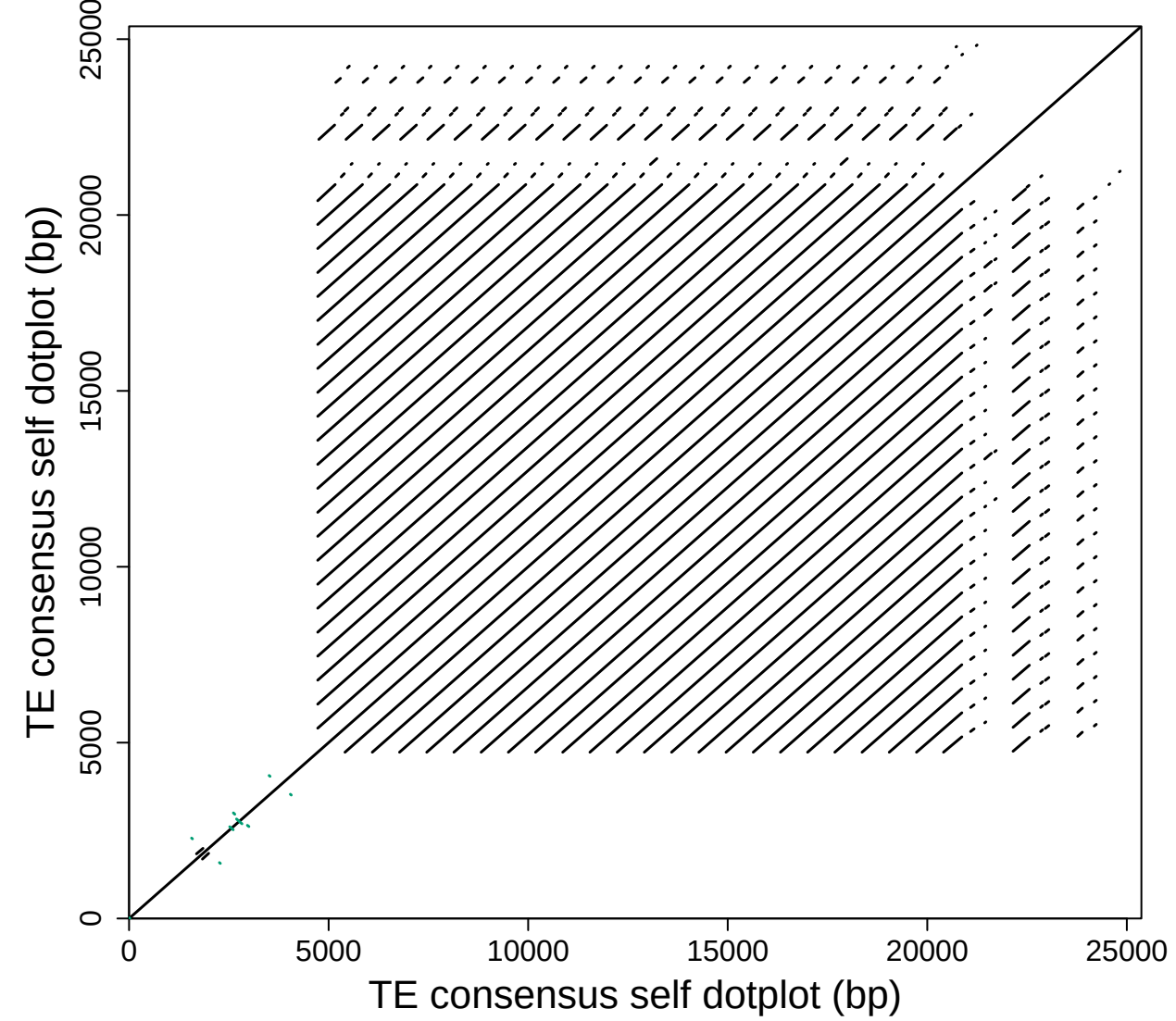
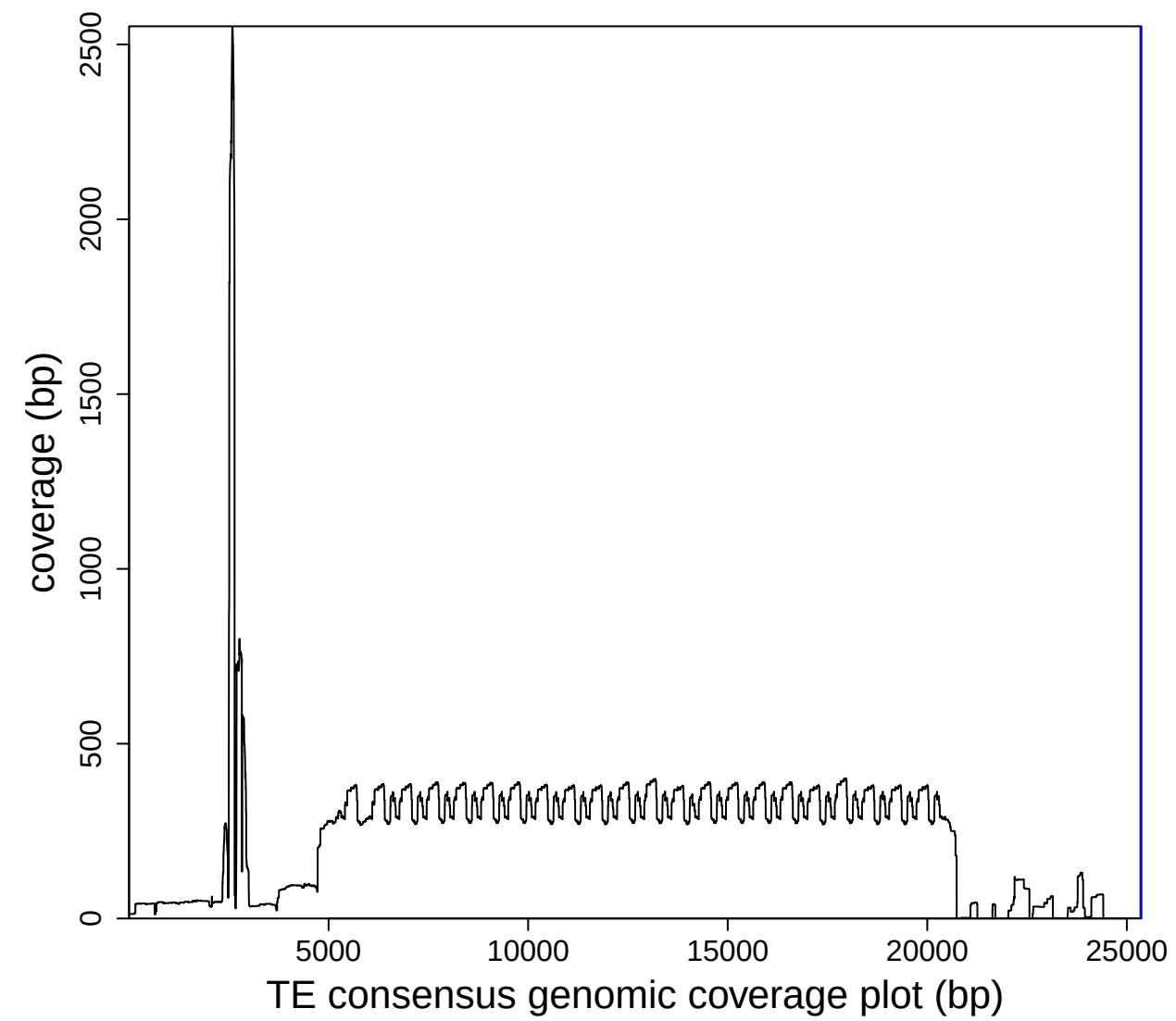
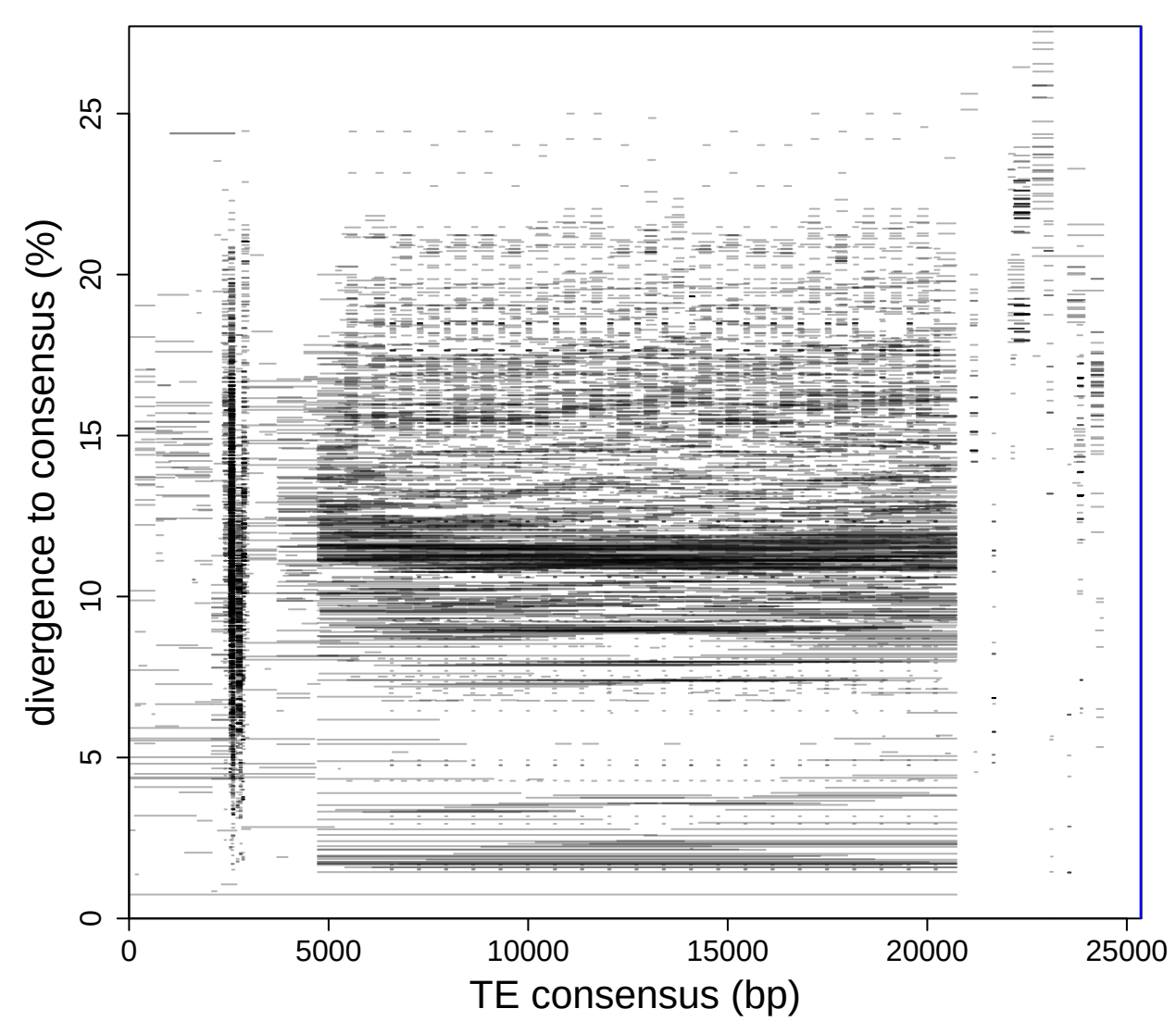
g_2.bed fm_1.bed 0 0 n.bed g_1.bed fm_2.bed 0 0 bcln.fa &
size: 25365bp; fragments: 16864; full length: 0 (>=22828.5bp)

After TEtrimmer 25365 bp

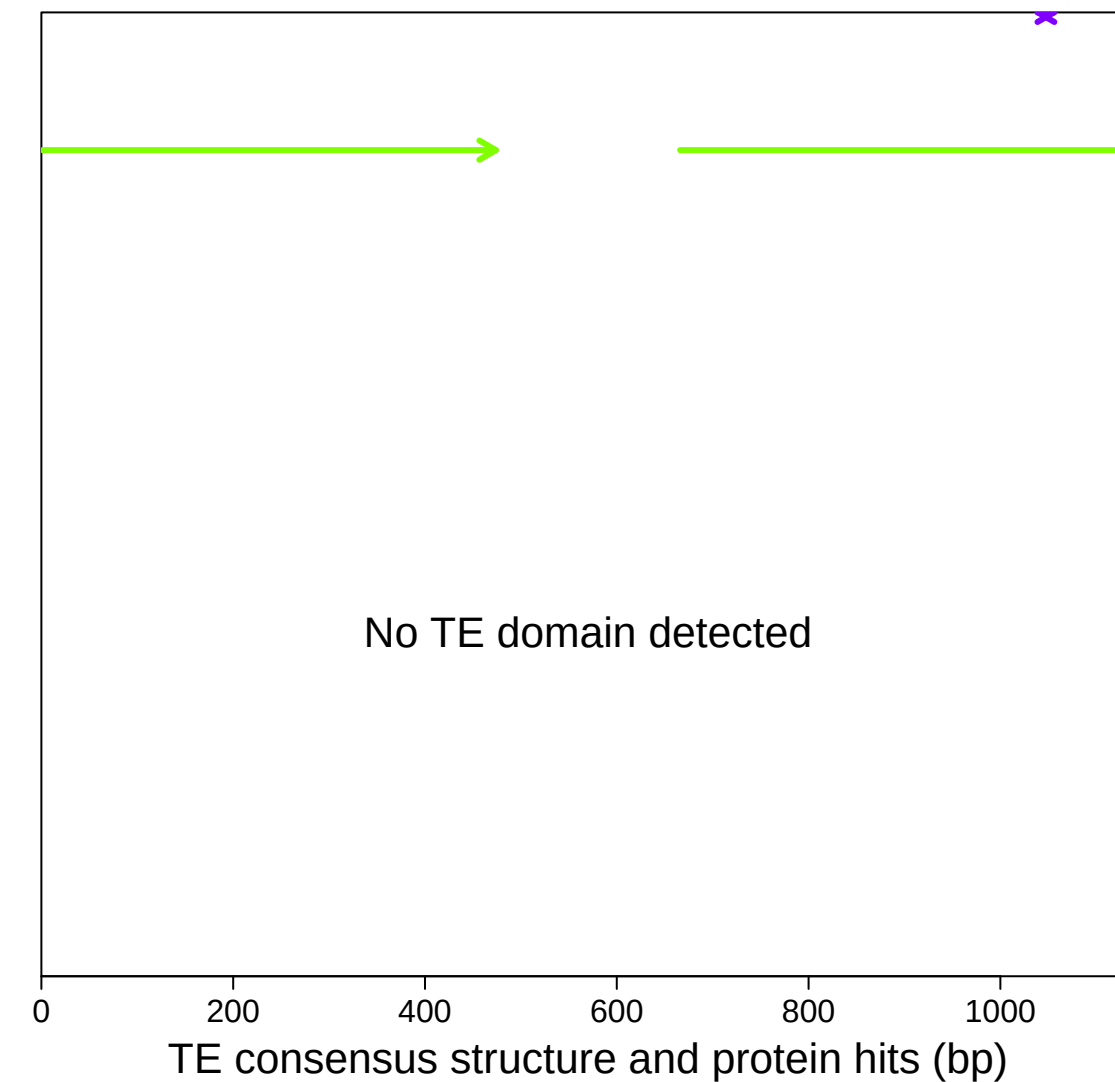
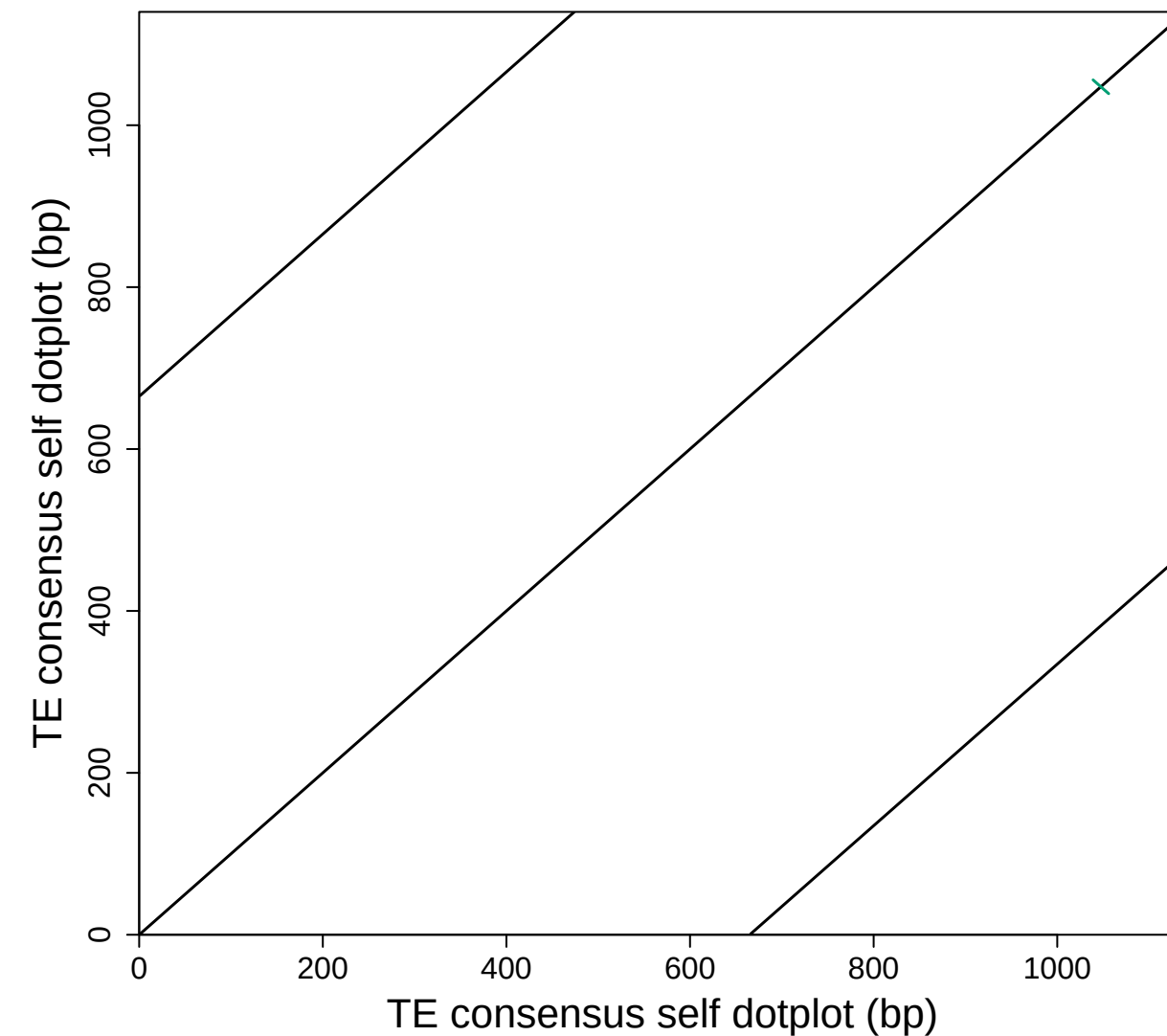
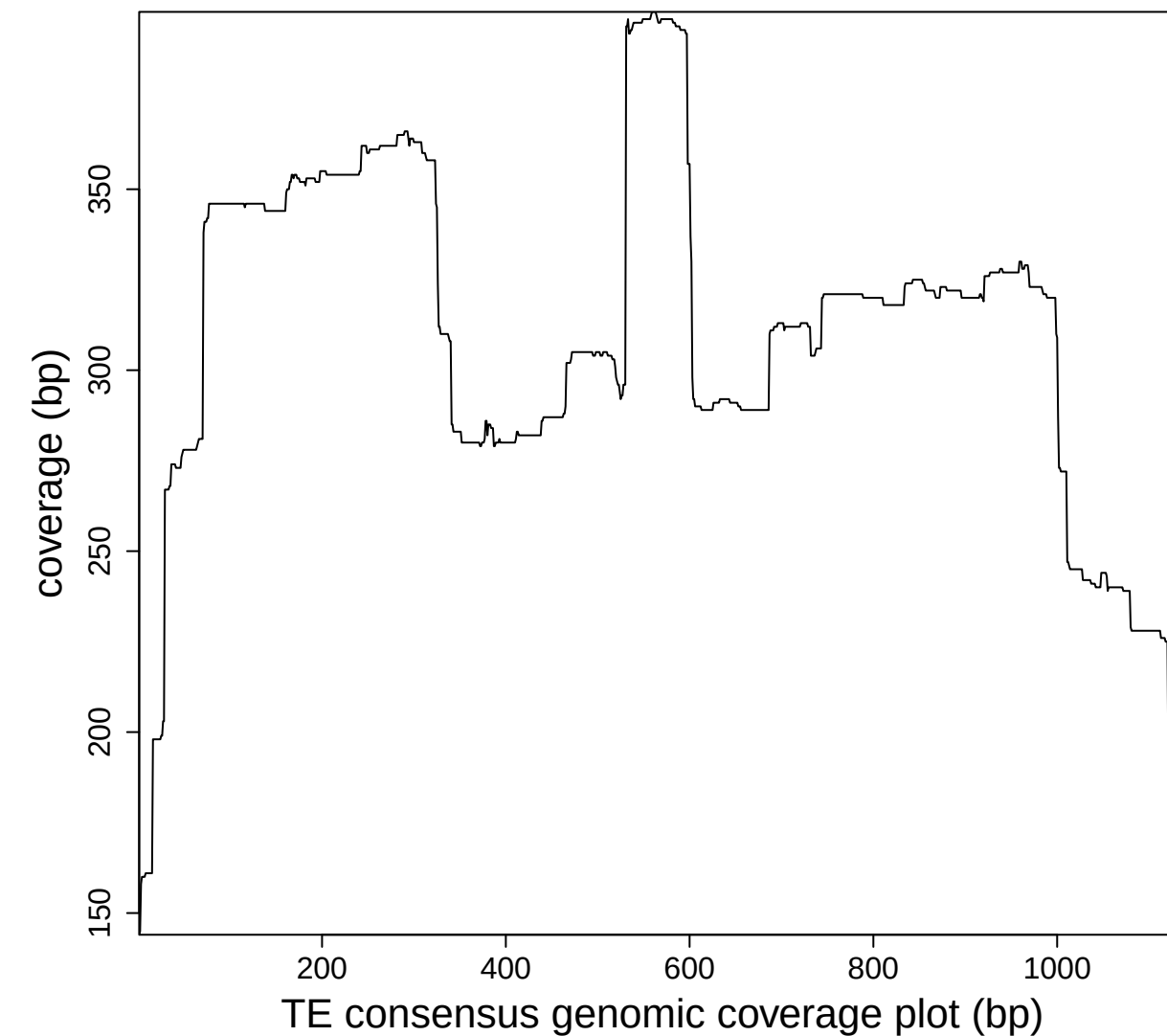
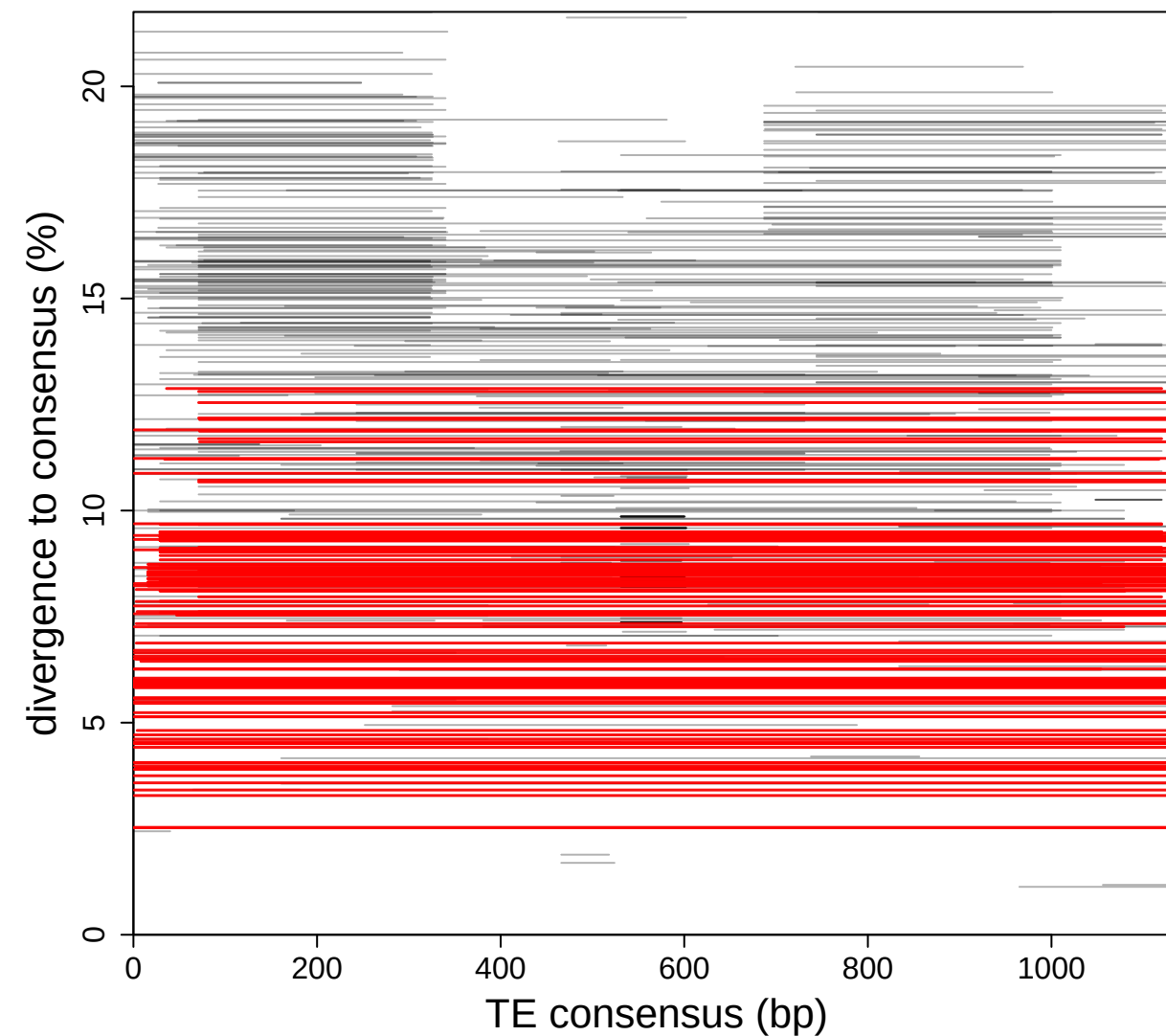


f.bed_g 2.bed_fm 1.bed_0 0 n.bed_g 1.bed_fm 2.bed_0 0 bc
size: 25365bp; fragments: 10946; full length: 0 (>=22828.5bp)

After TEtrimmer Extended plot Blue lines are boundaries

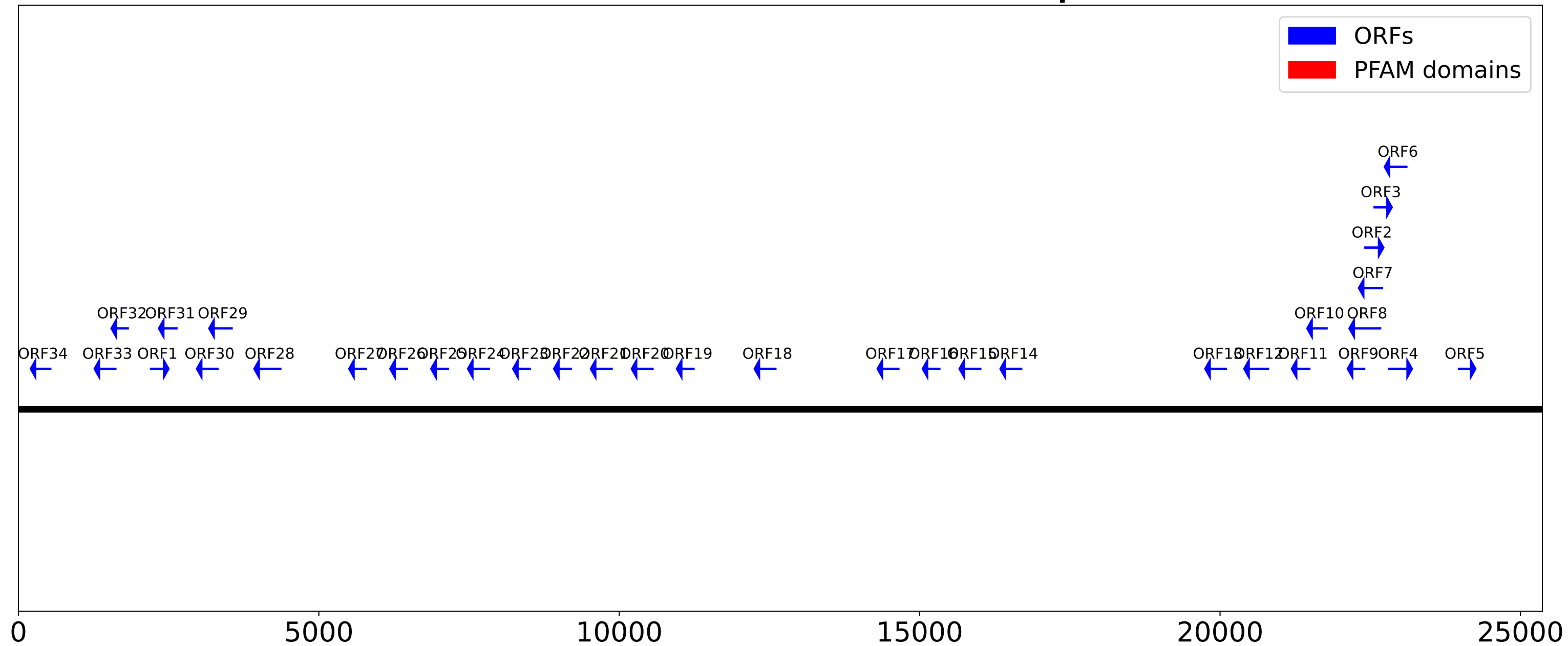


TE: rnd_4_family_7213
size: 1140bp; fragments: 614; full length: 157 (≥ 1026 bp)

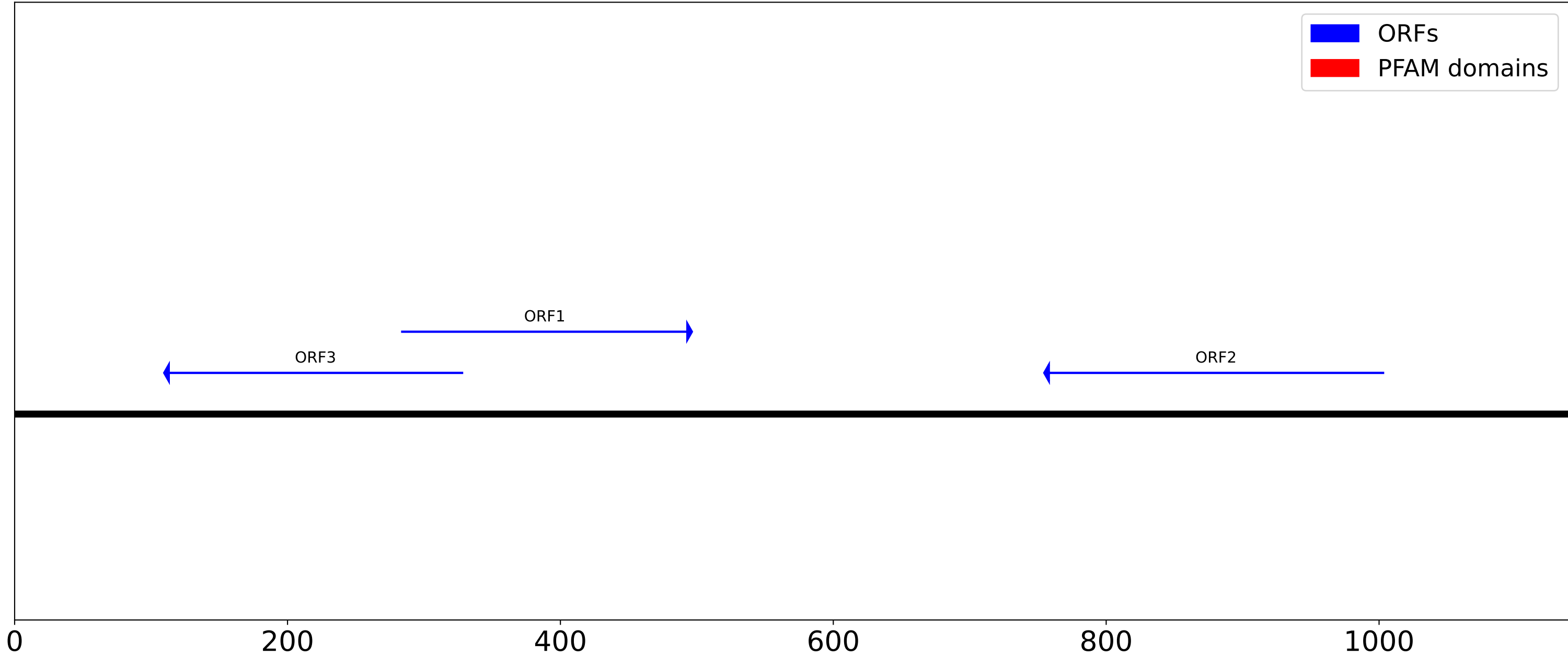


Before TEtrimmer 1140 bp

After TEtrimmer ORF and PFAM domain plot



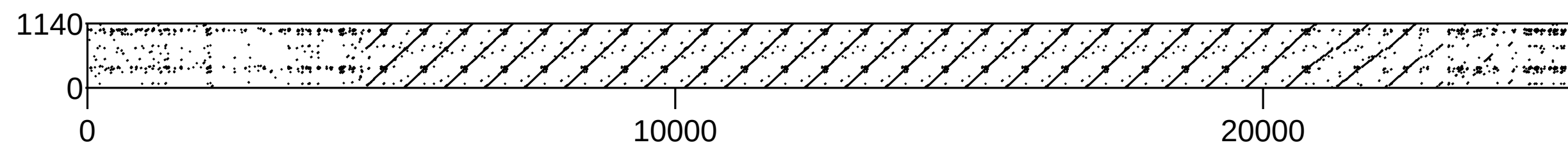
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

nsus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)