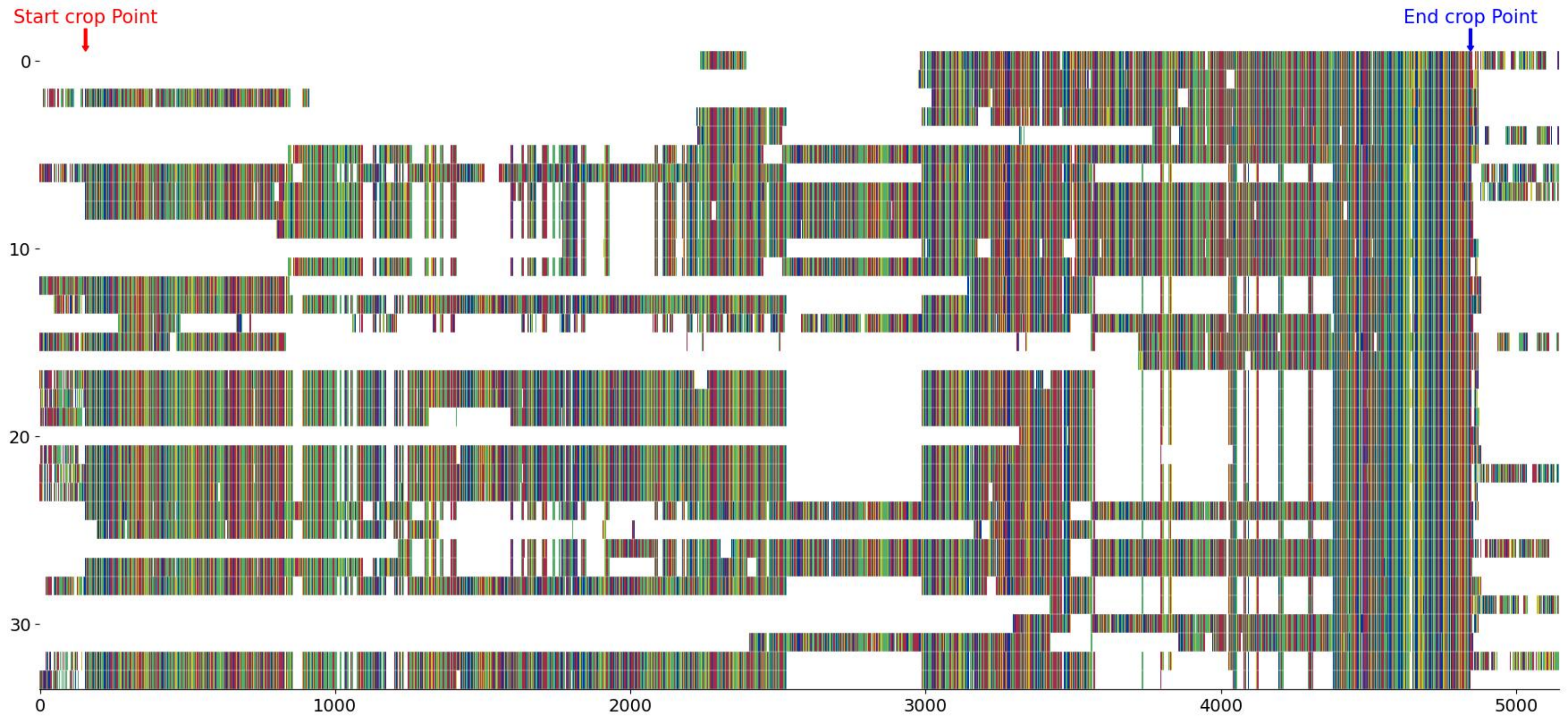


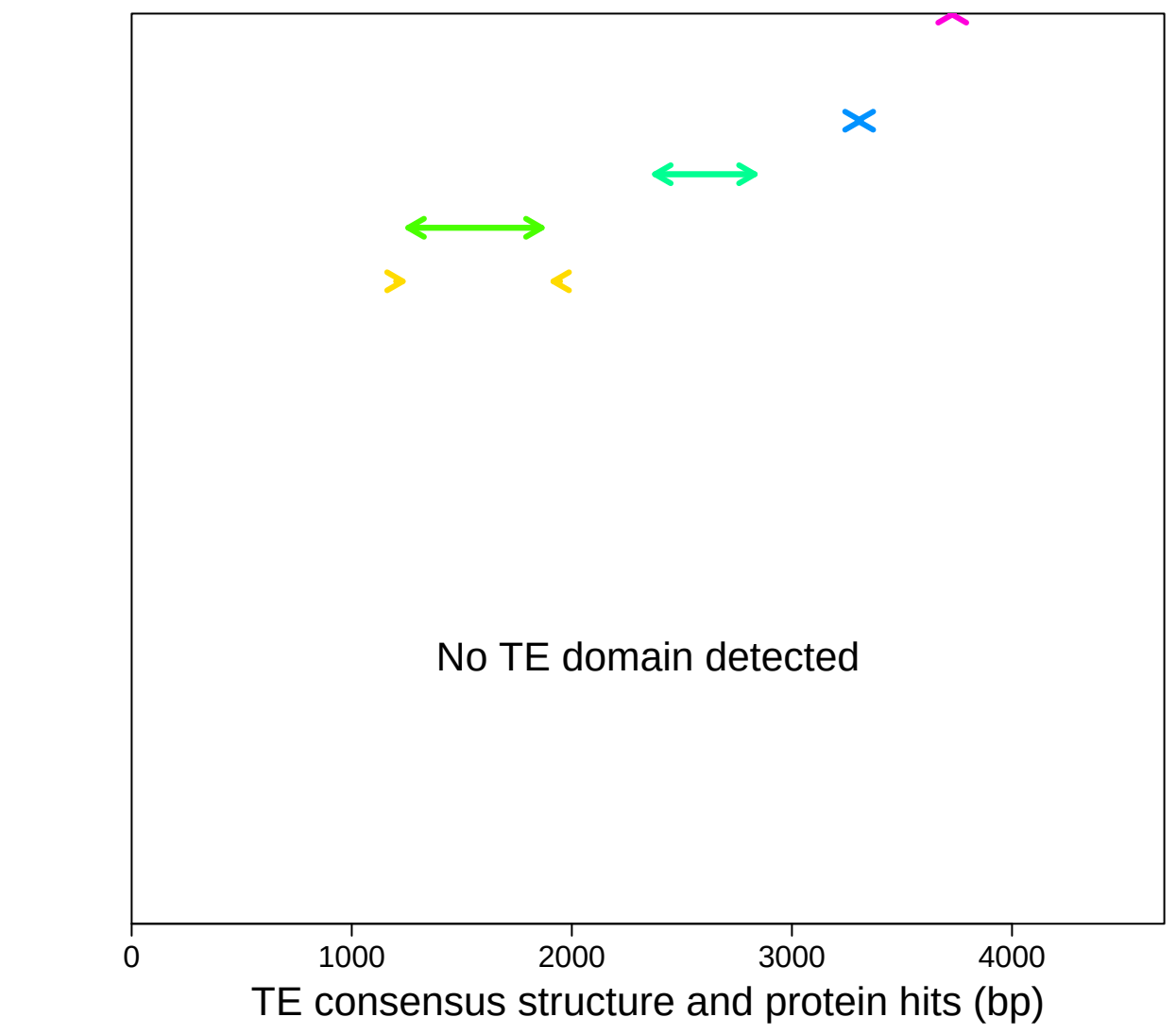
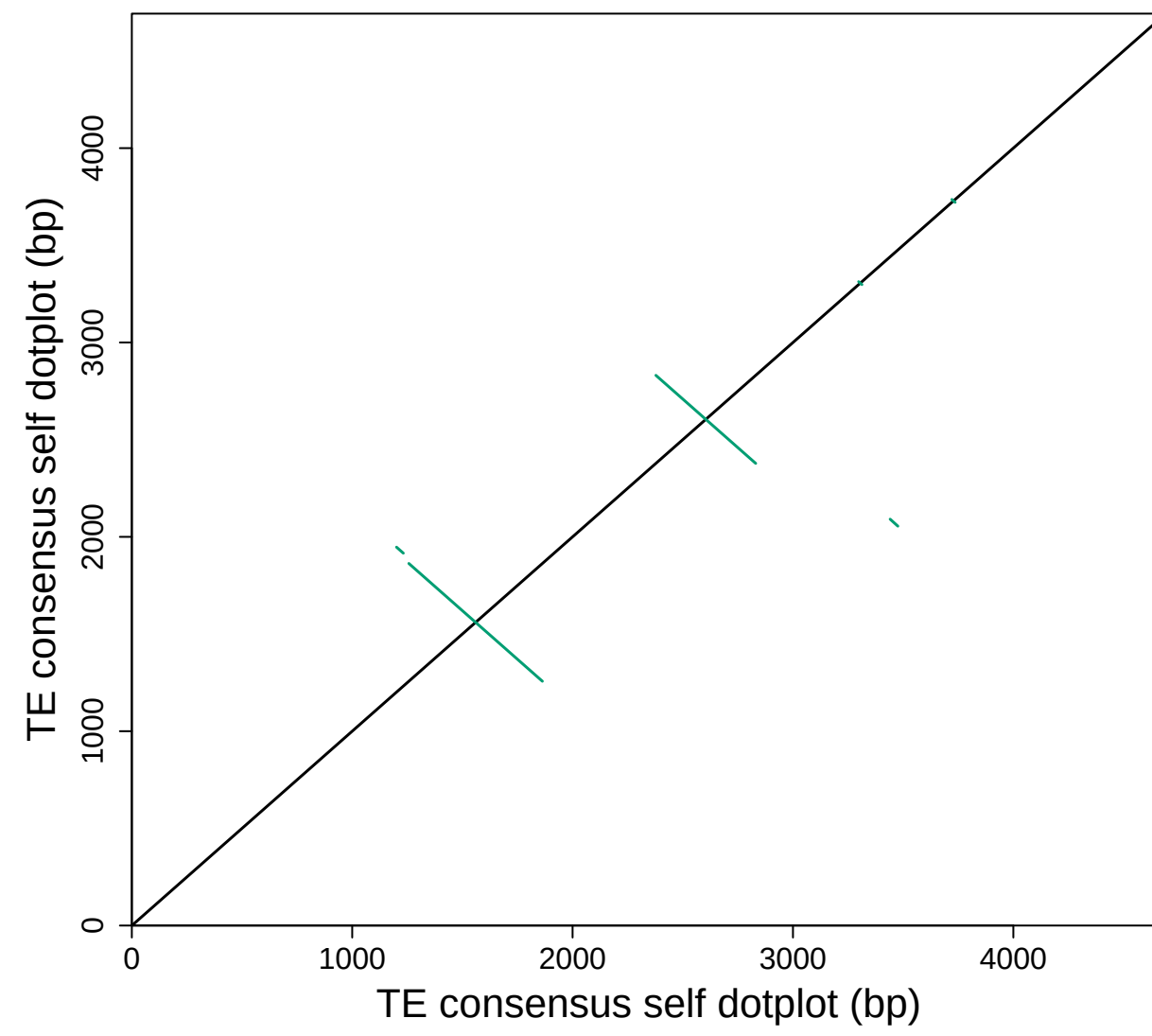
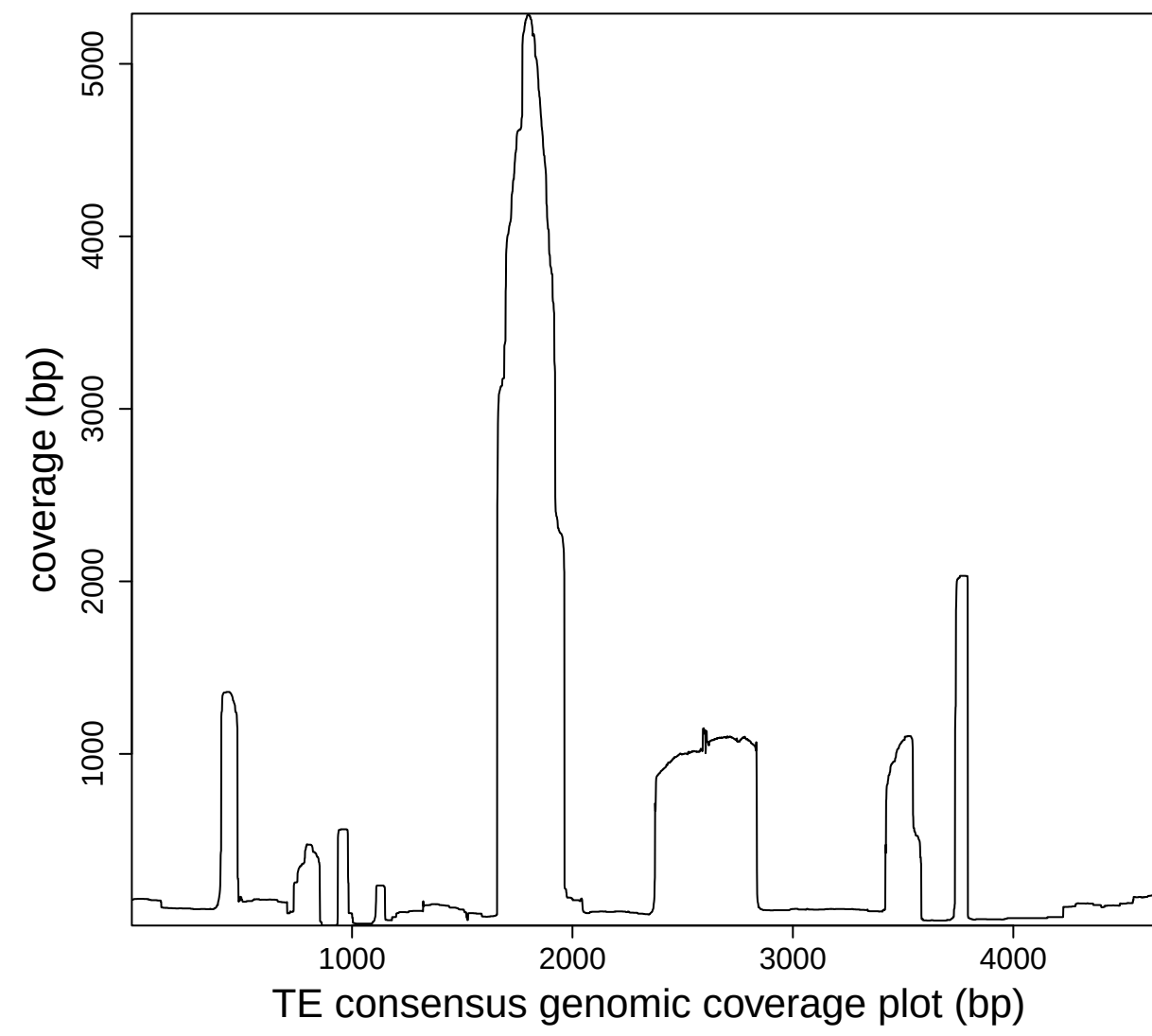
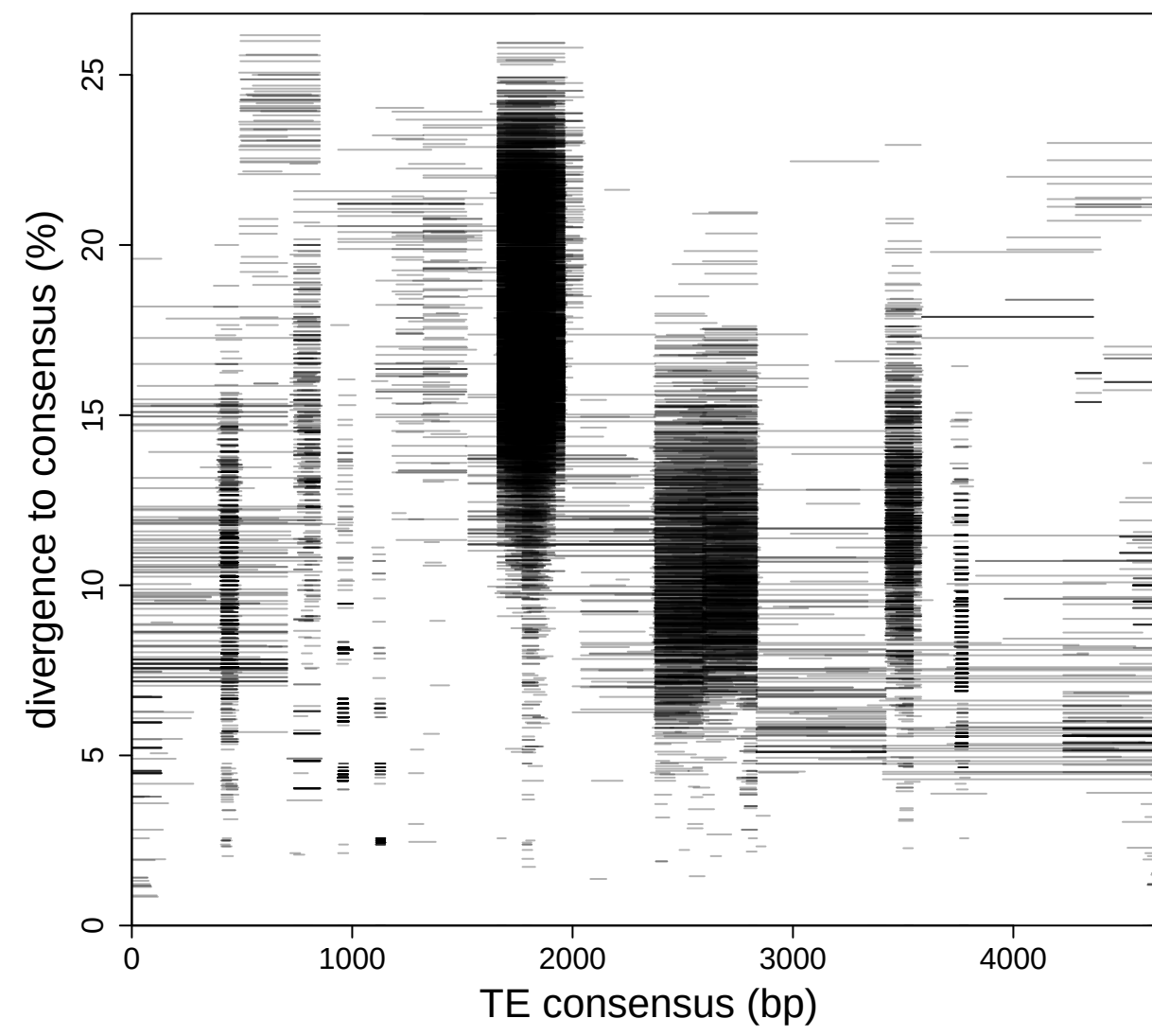
MSA length = 4692





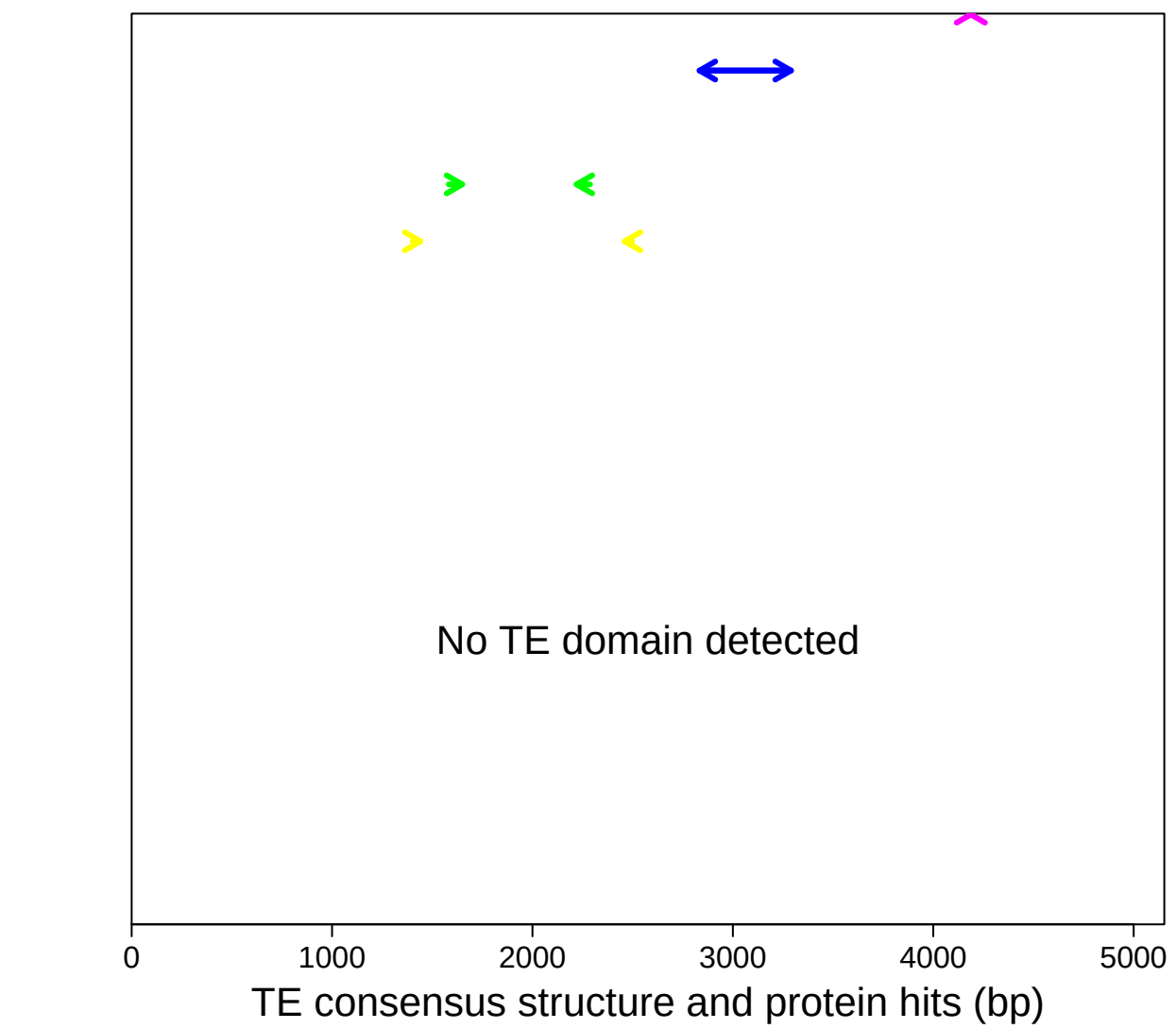
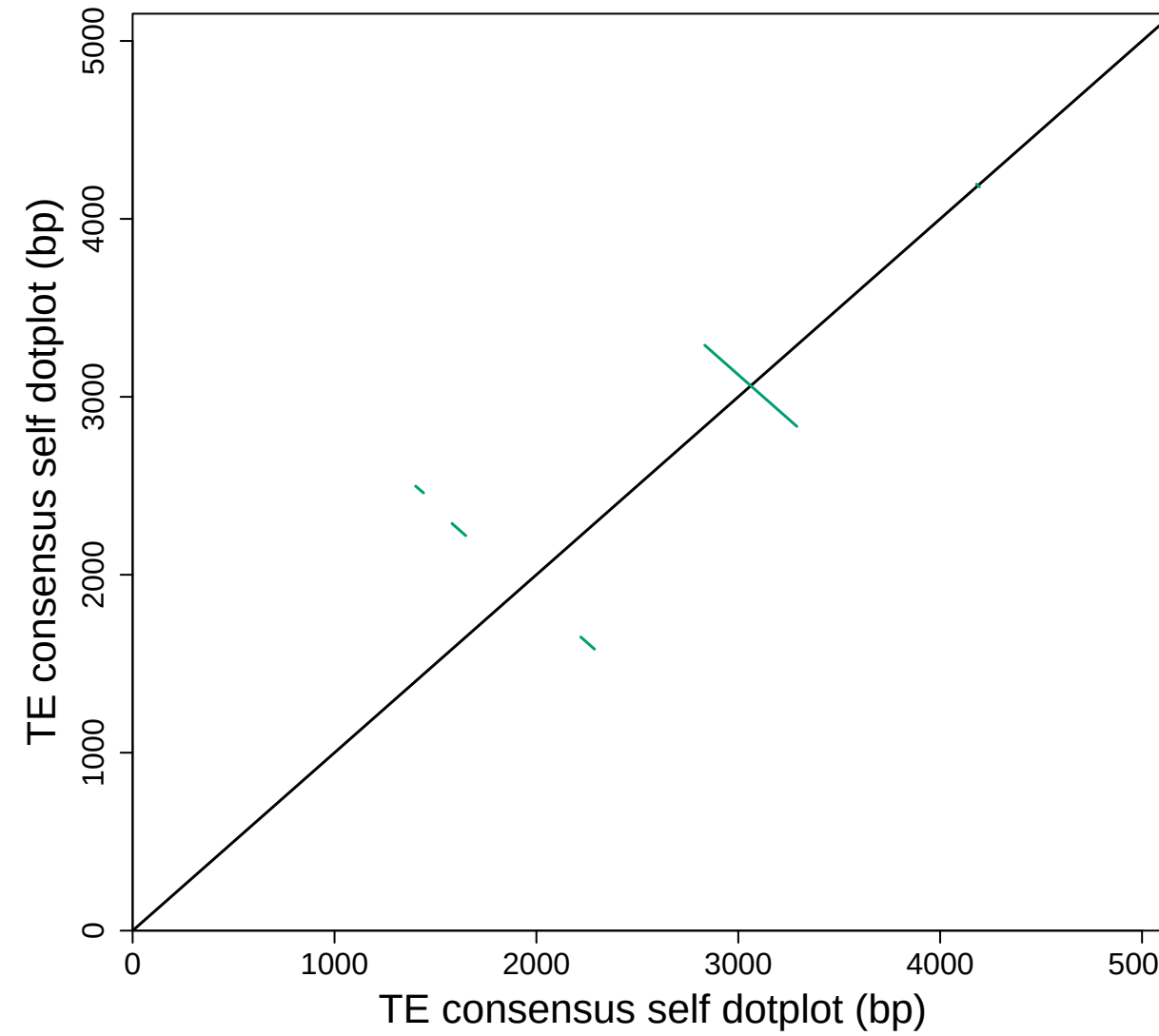
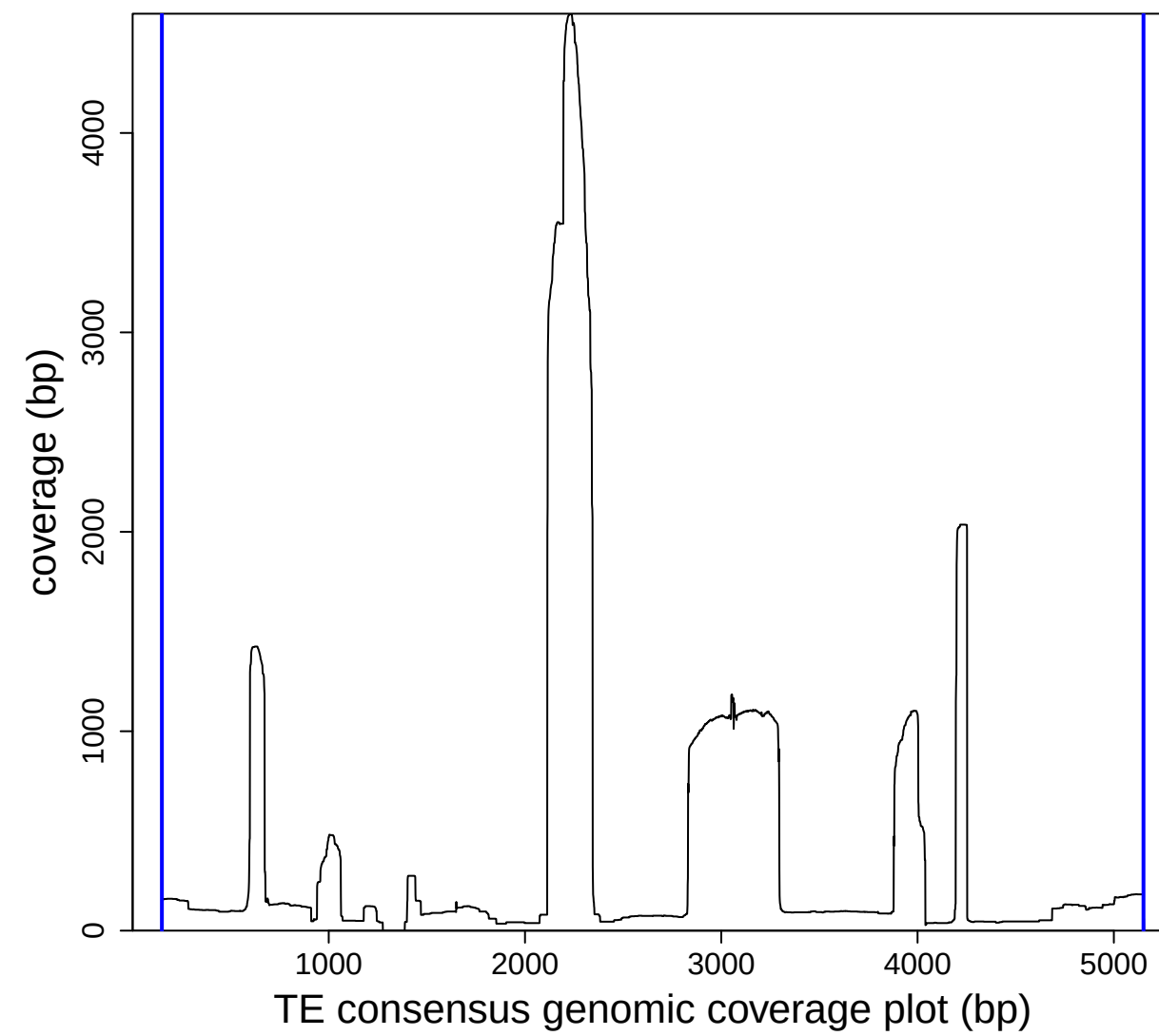
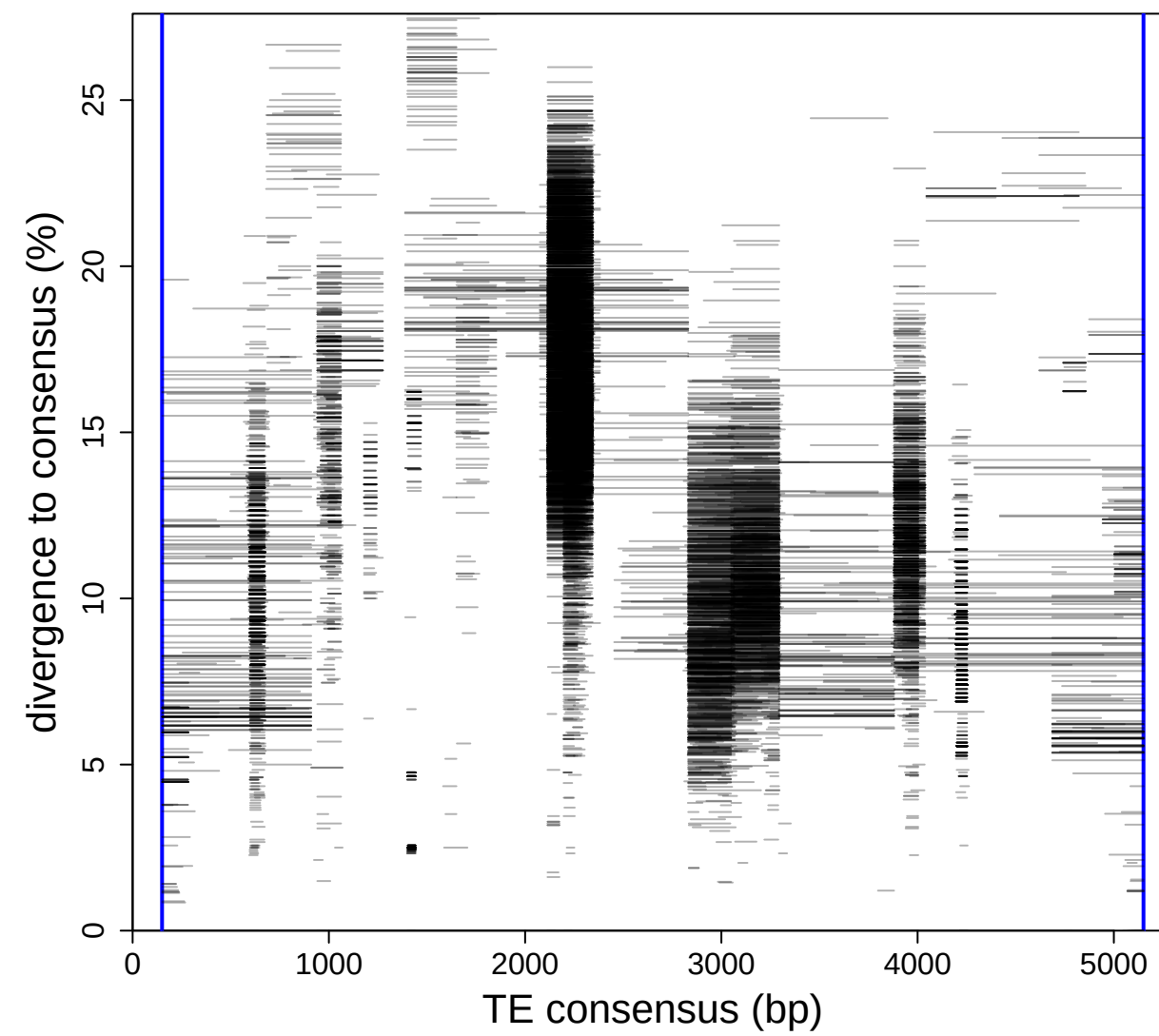
g_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcln.fa
size: 4692bp; fragments: 13347; full length: 0 (>=4222.8bp)

After TEtrimmer 4692 bp

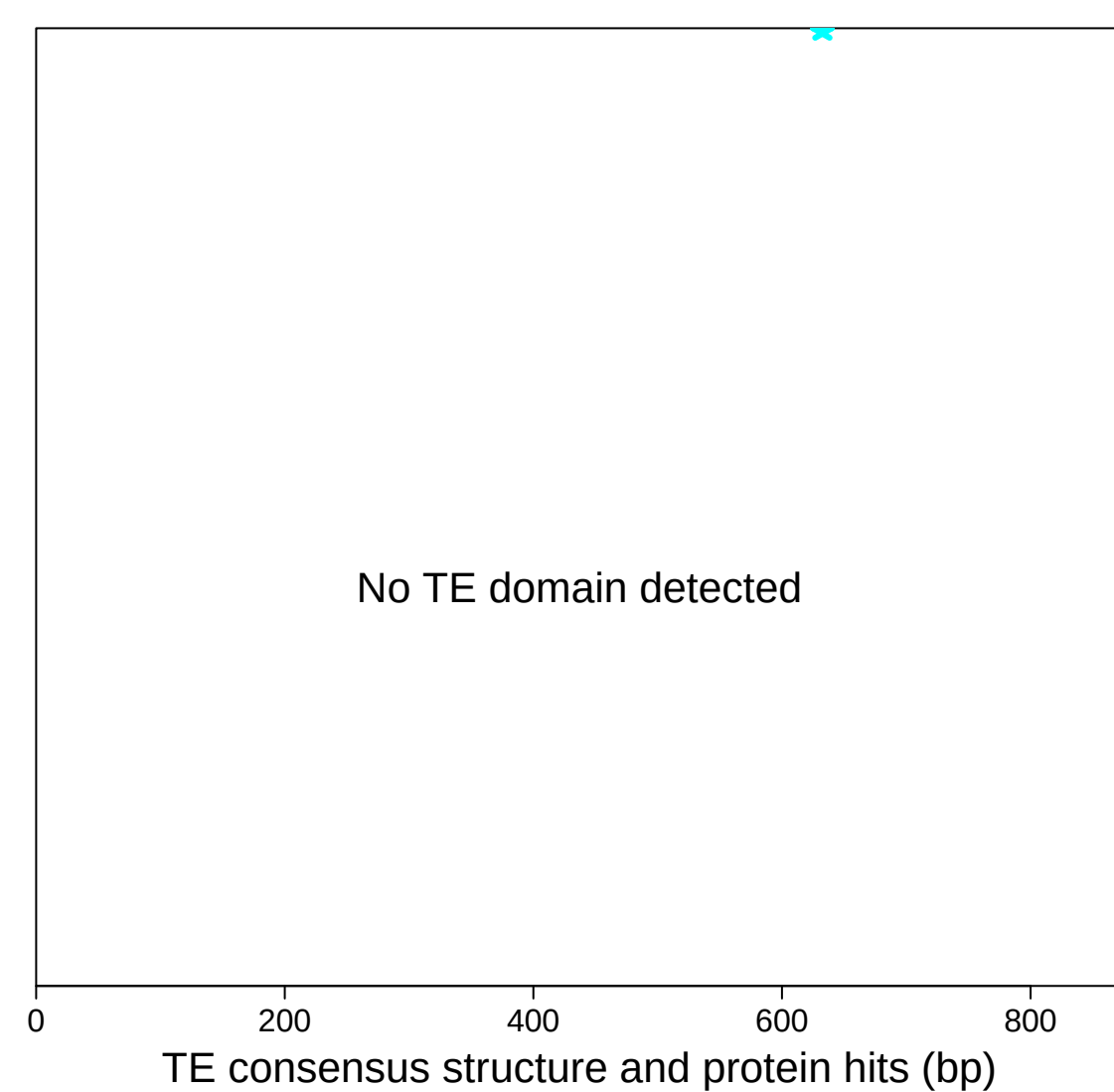
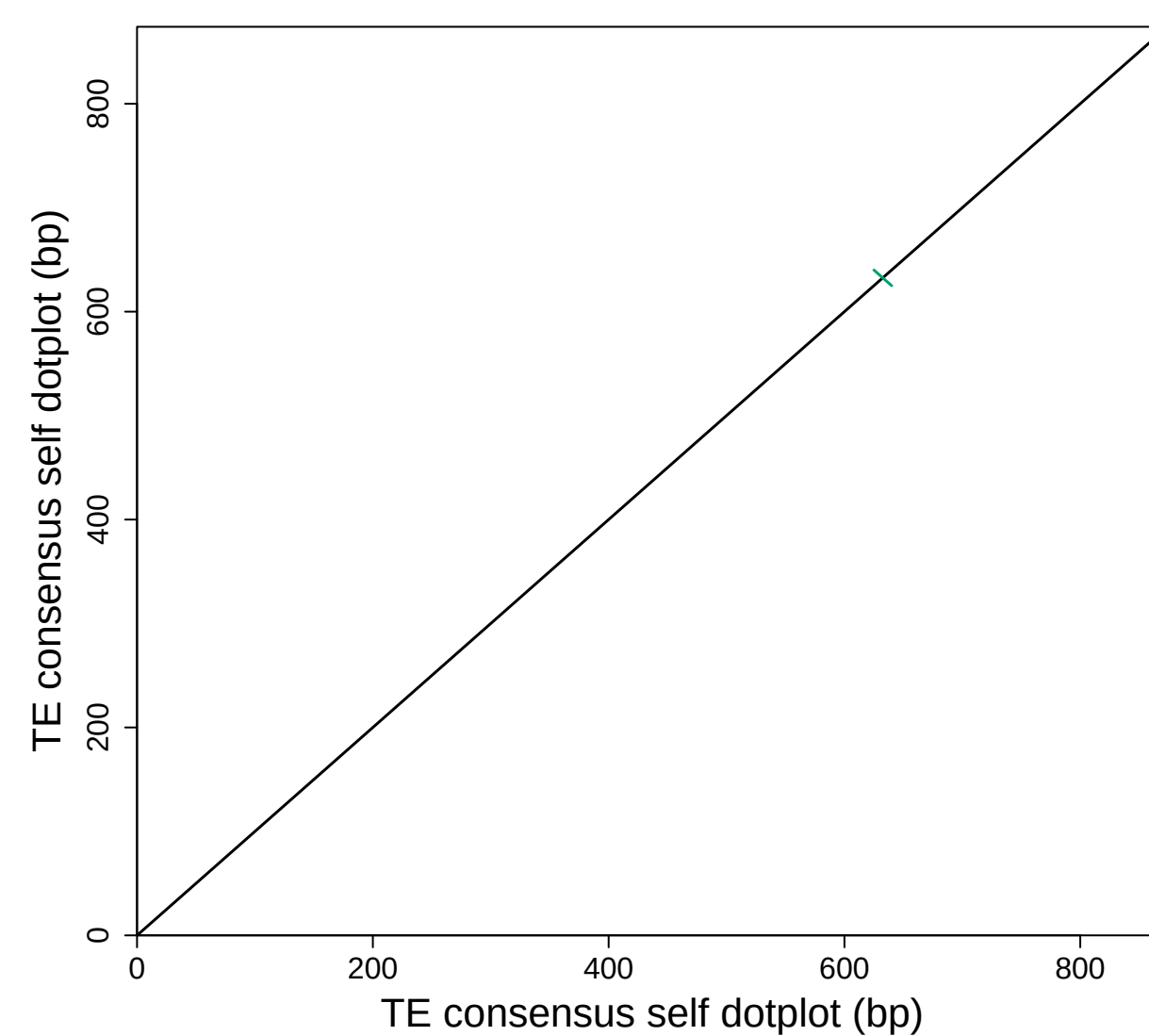
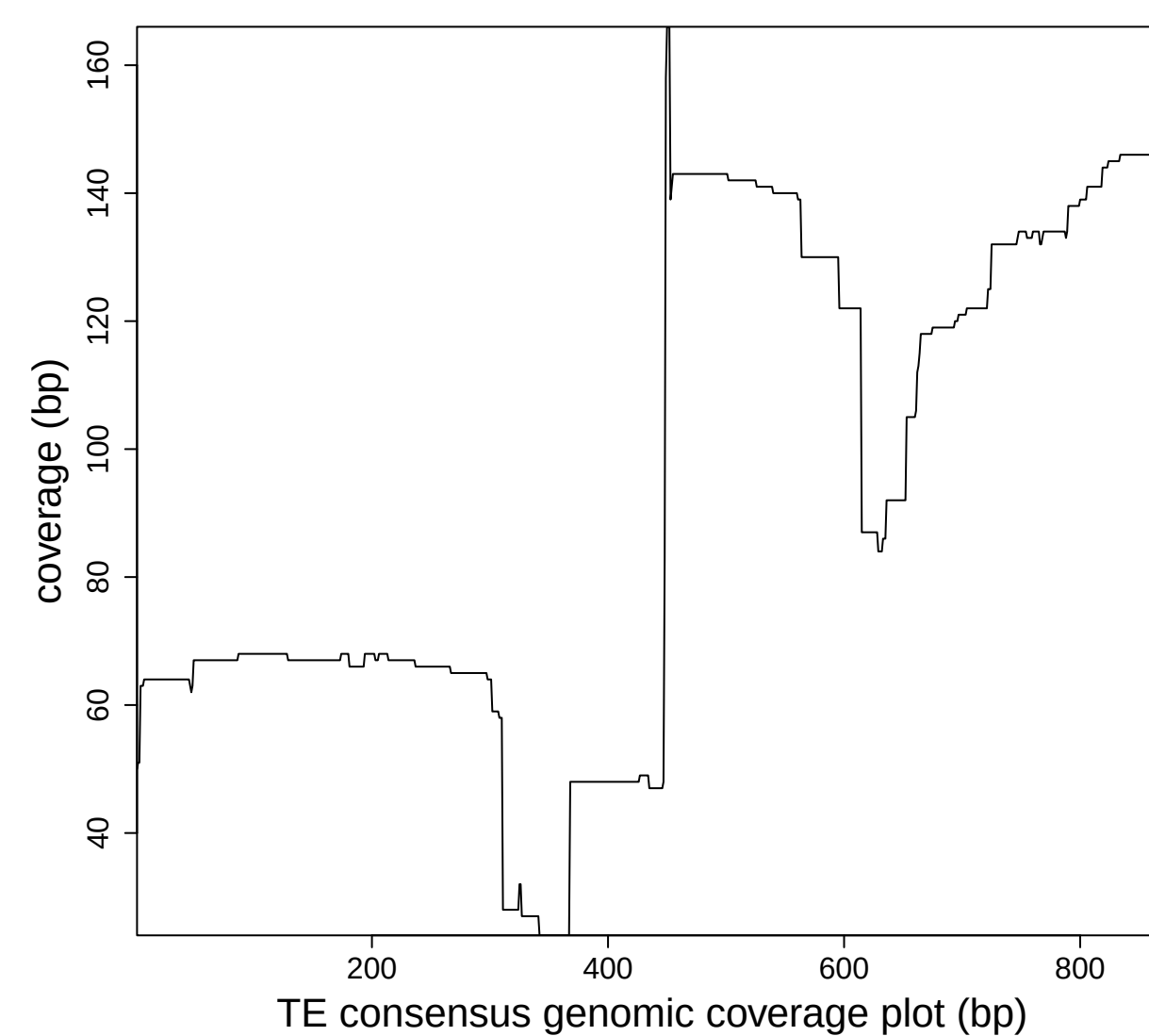
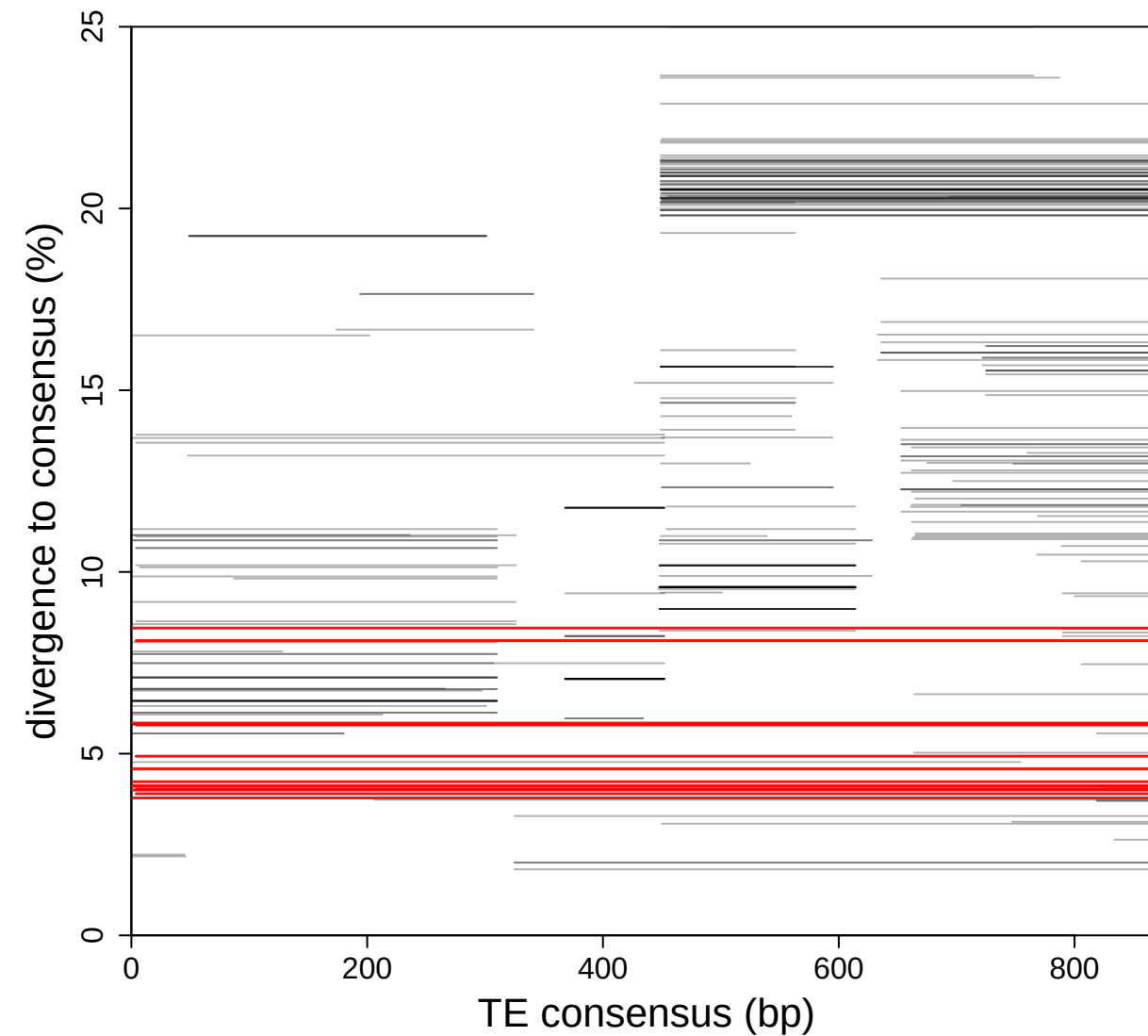


uf.bed g 1.bed fm 1.bed 0 0 n.bed g 1.bed fm 2.bed 0 0 bc
size: 5301bp; fragments: 12287; full length: 0 (>=4770.9bp)

After TEtrimmer Extended plot Blue lines are boundaries

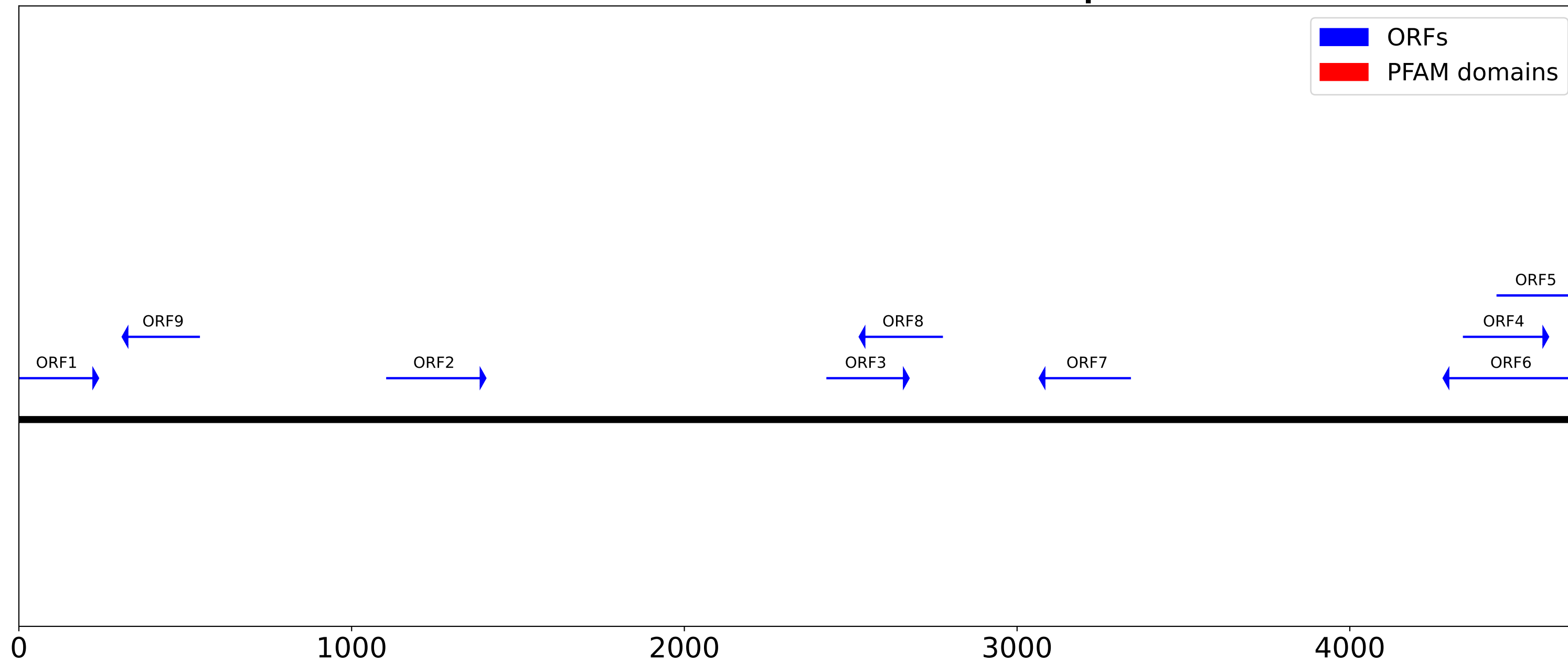


TE: rnd_5_family_11408
size: 874bp; fragments: 292; full length: 13 (≥ 786.6 bp)

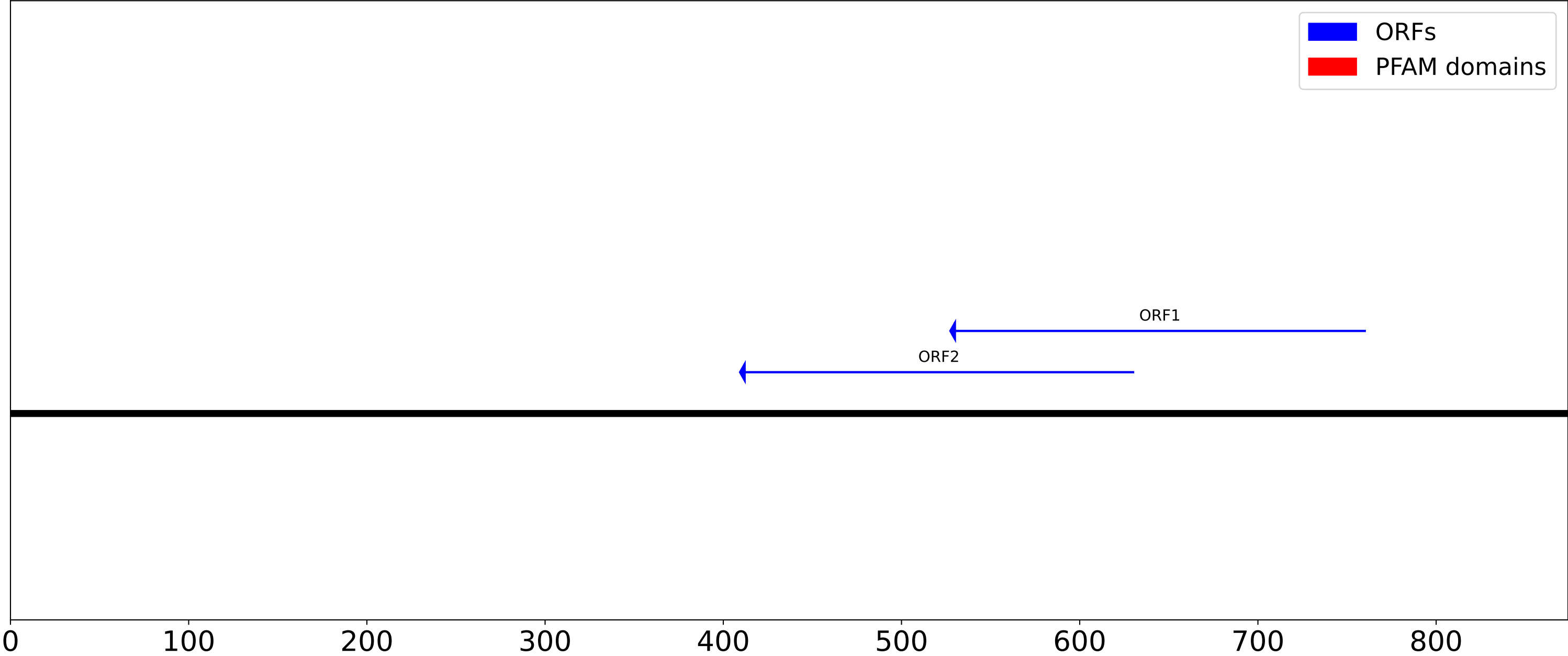


Before TEtrimmer 874 bp

After TEtrimmer ORF and PFAM domain plot



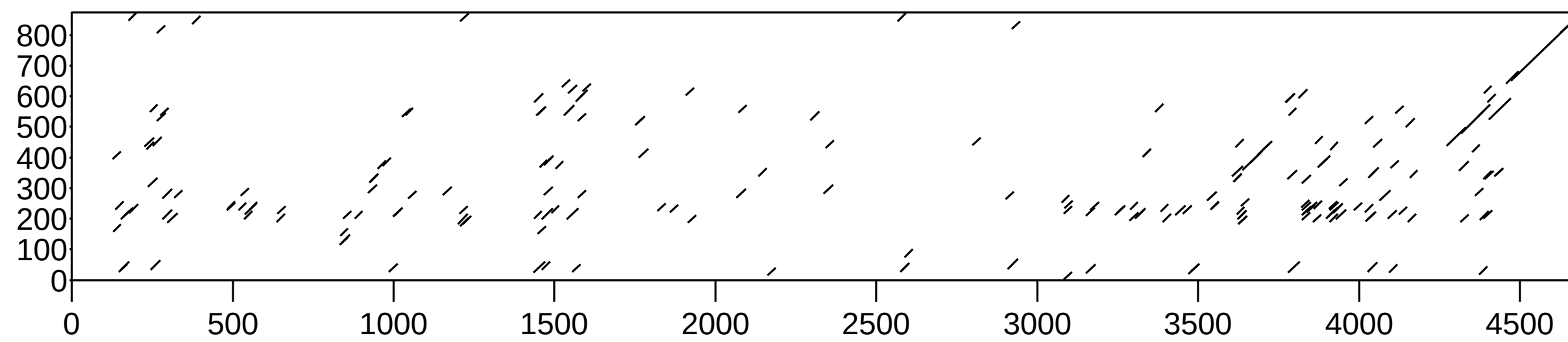
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

nsensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)