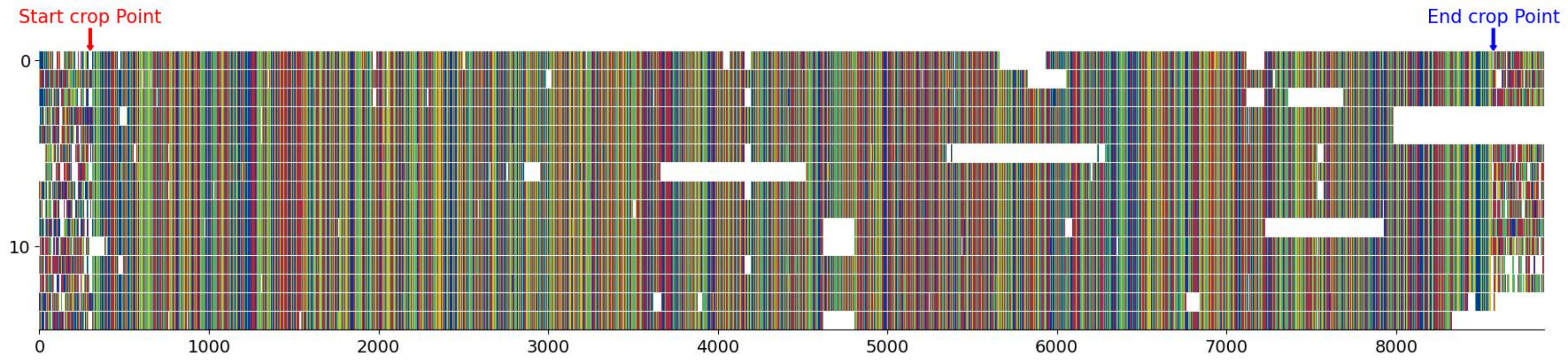
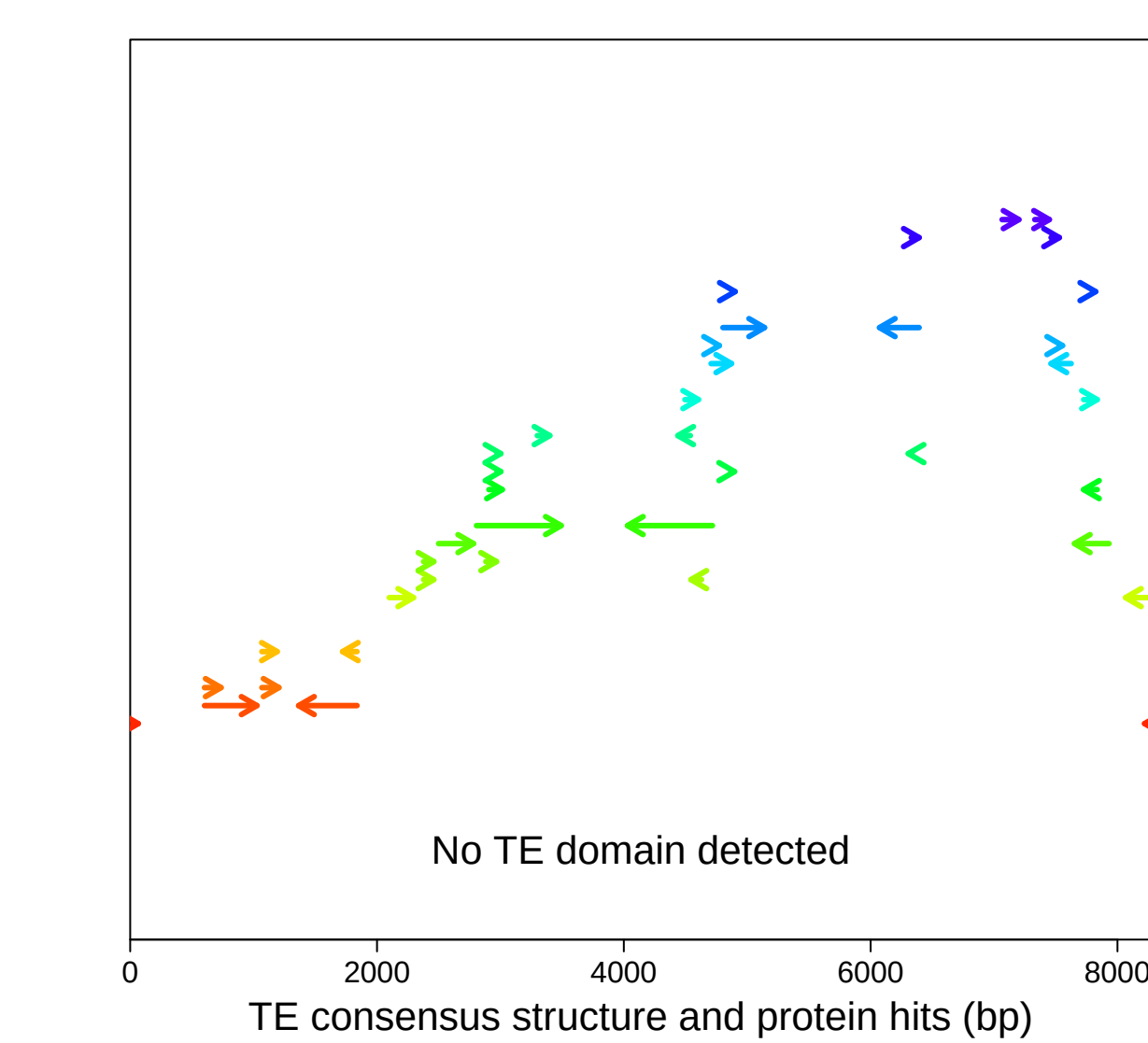
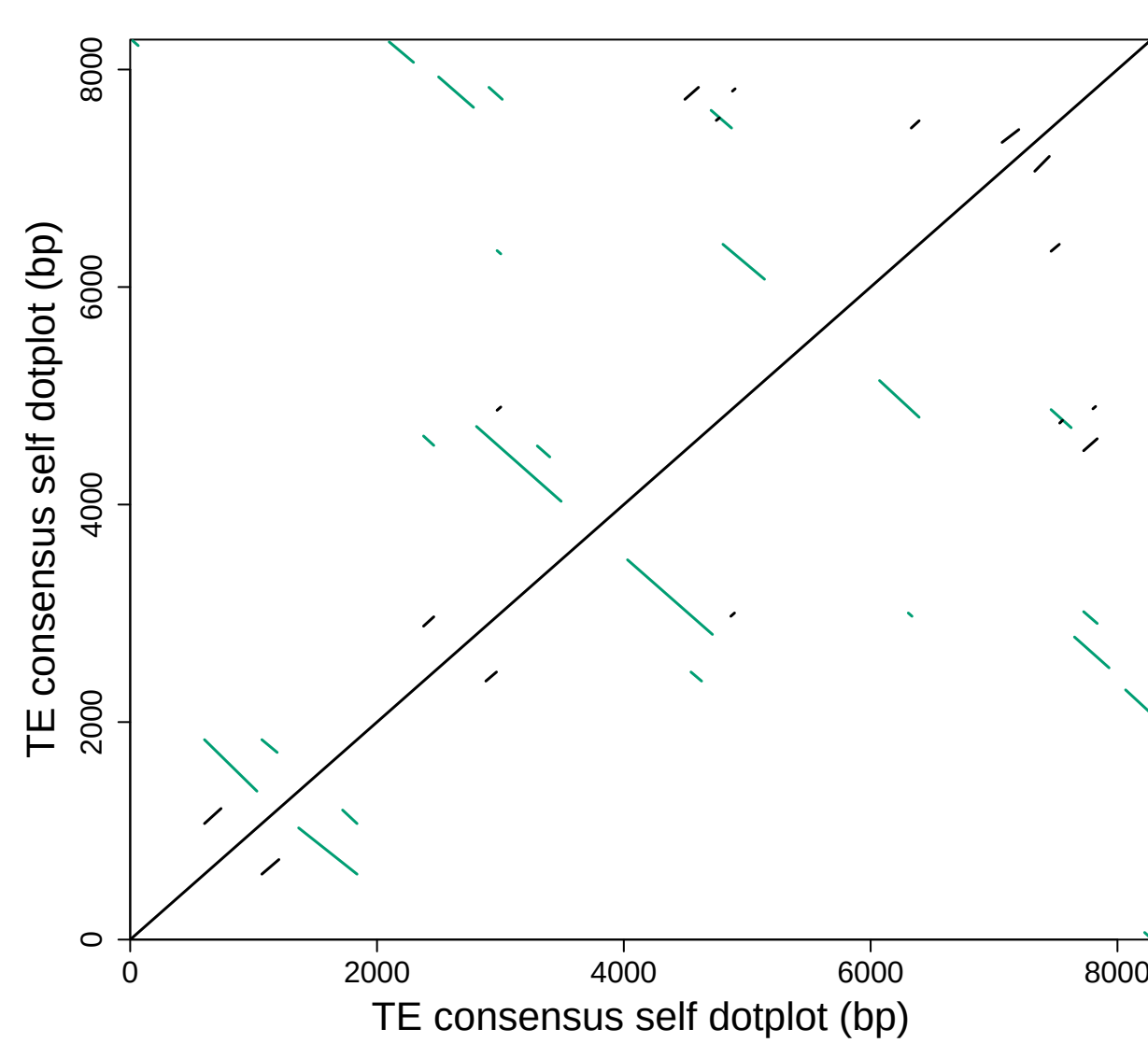
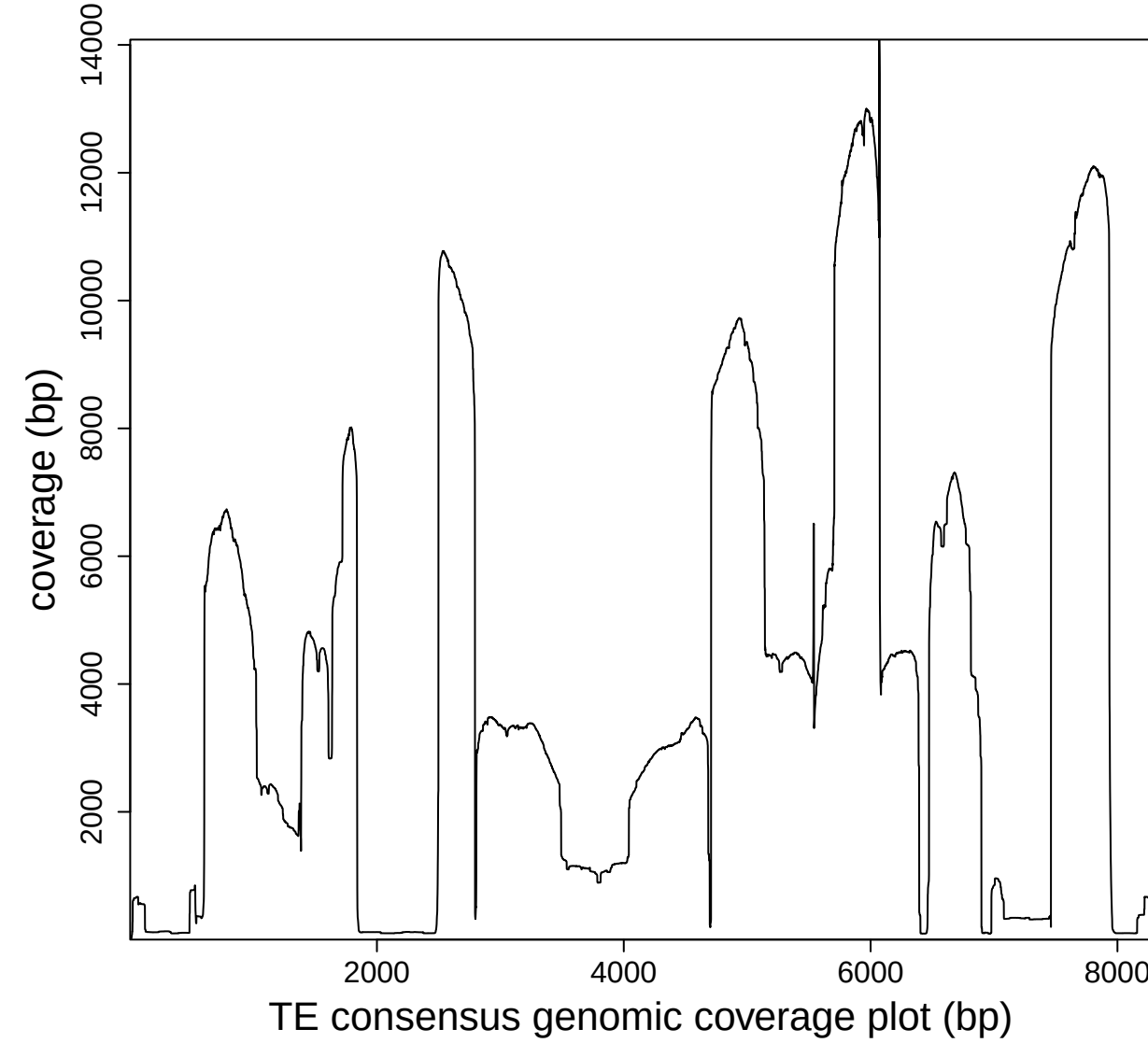
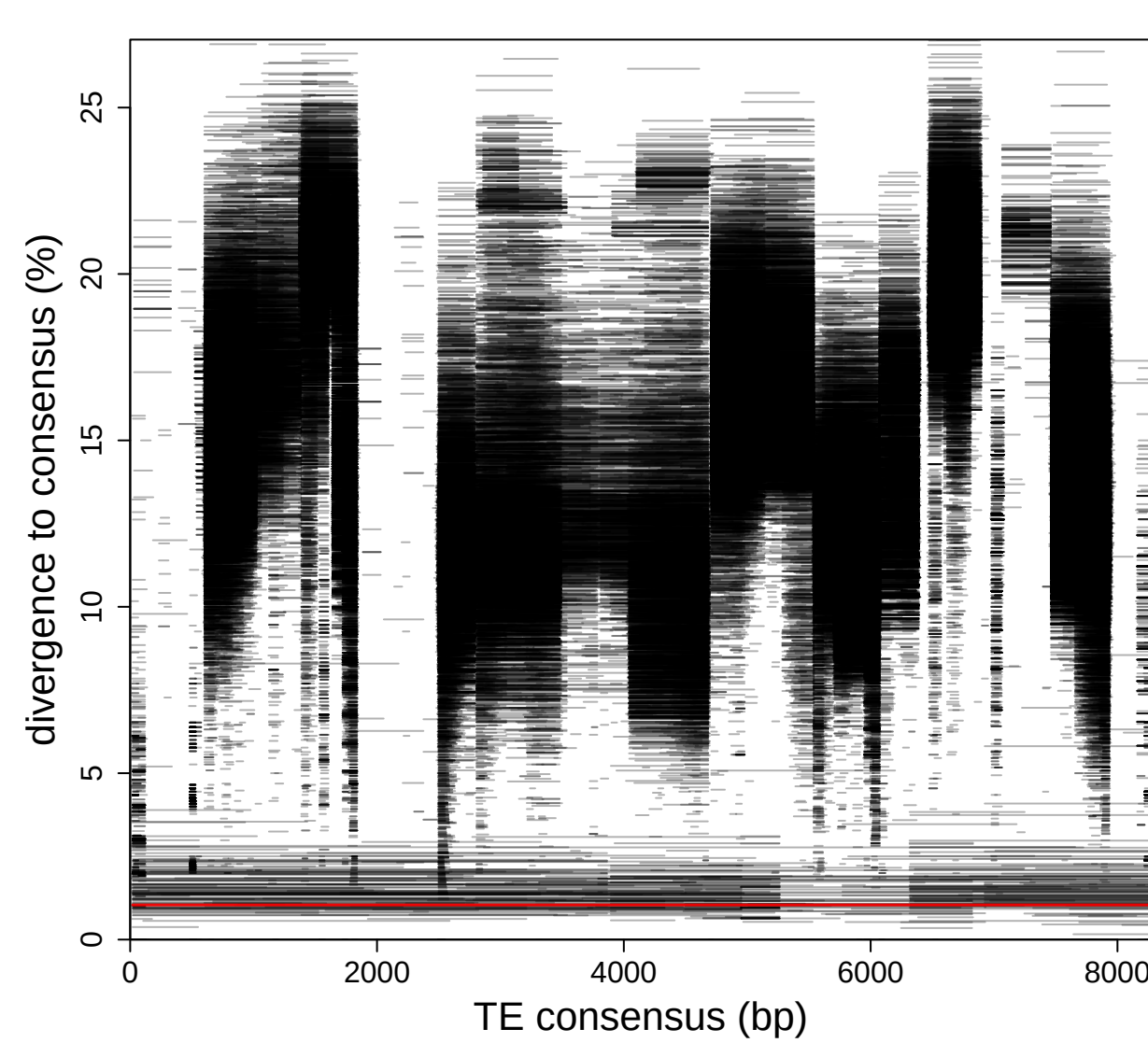






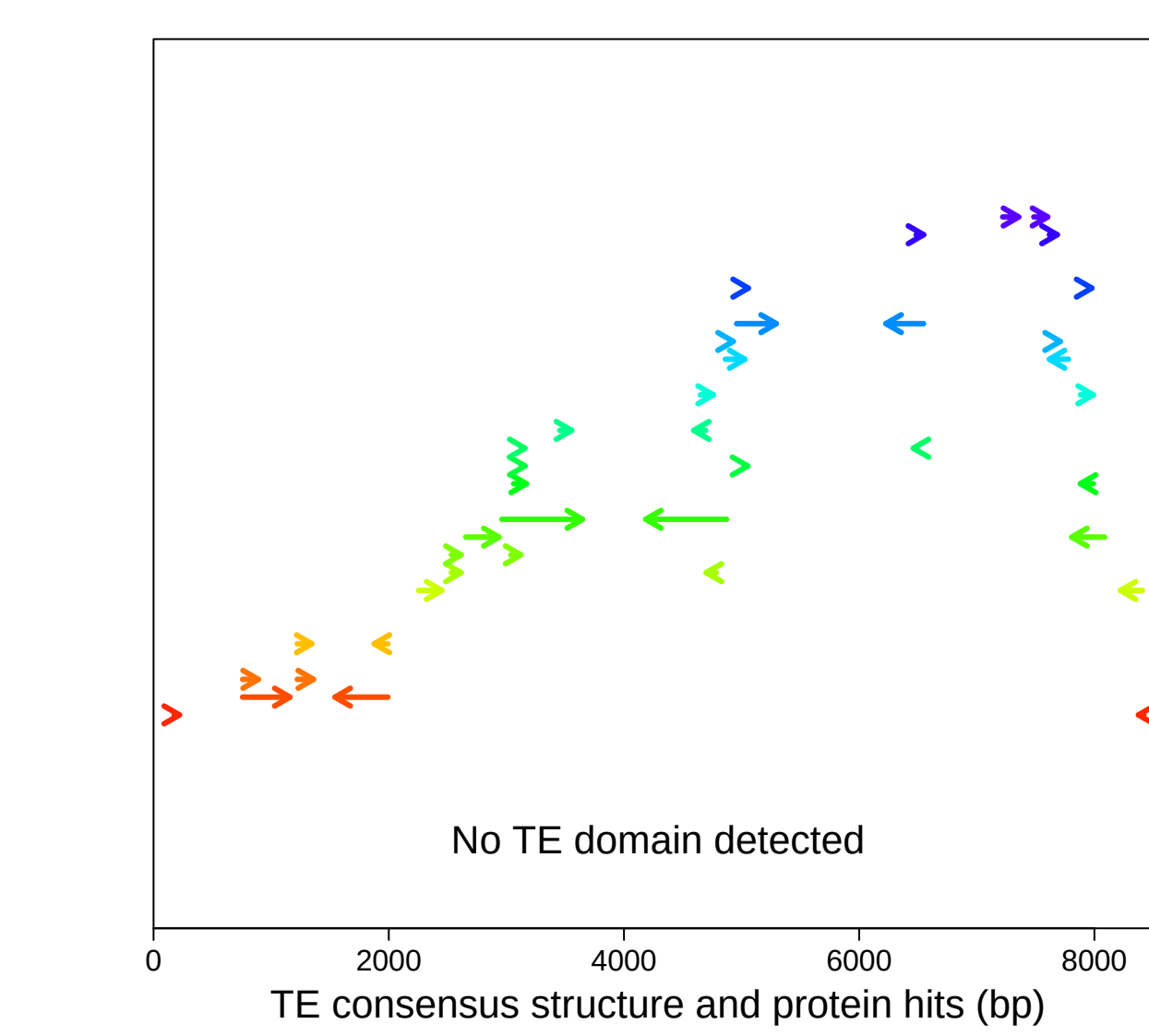
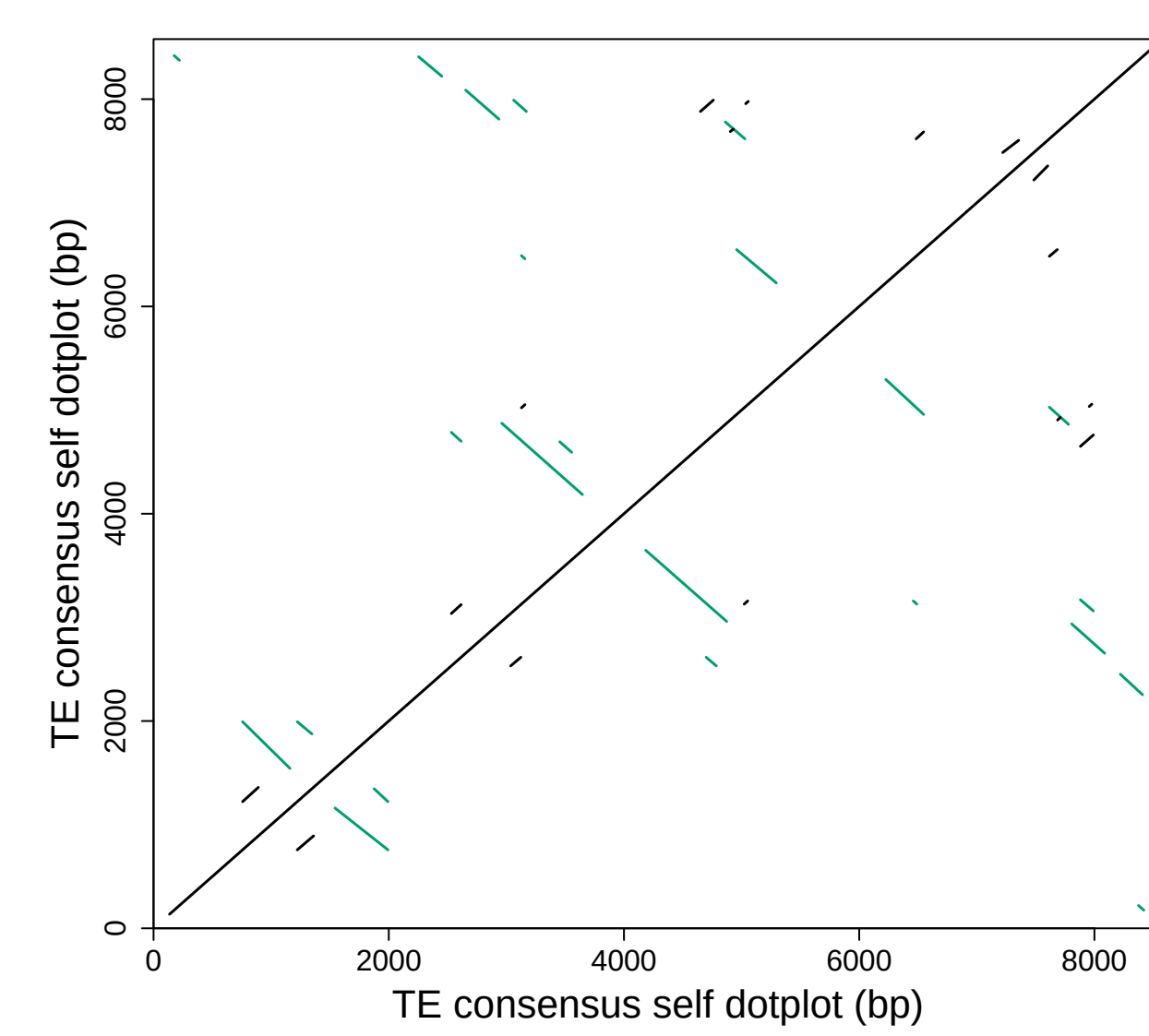
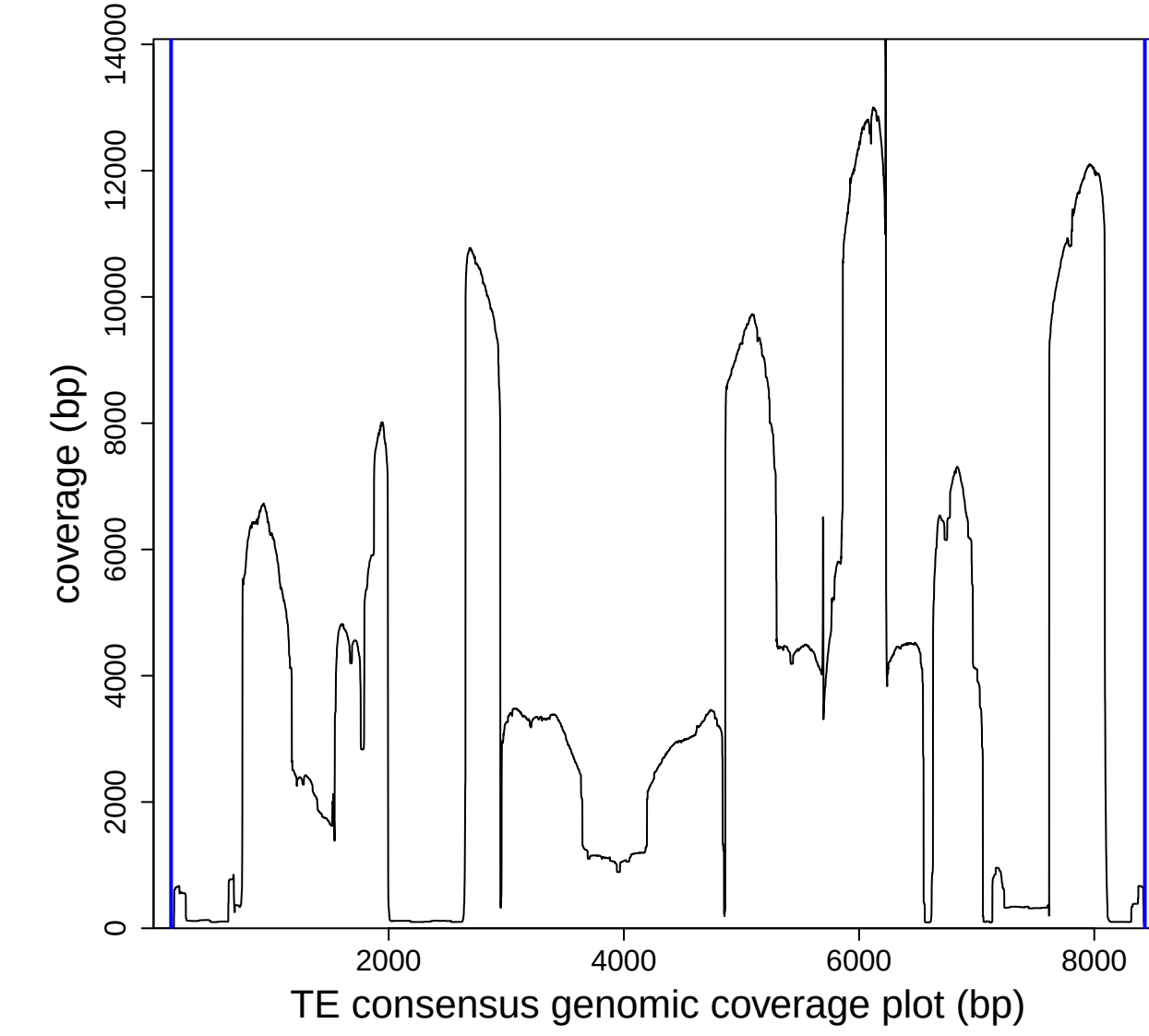
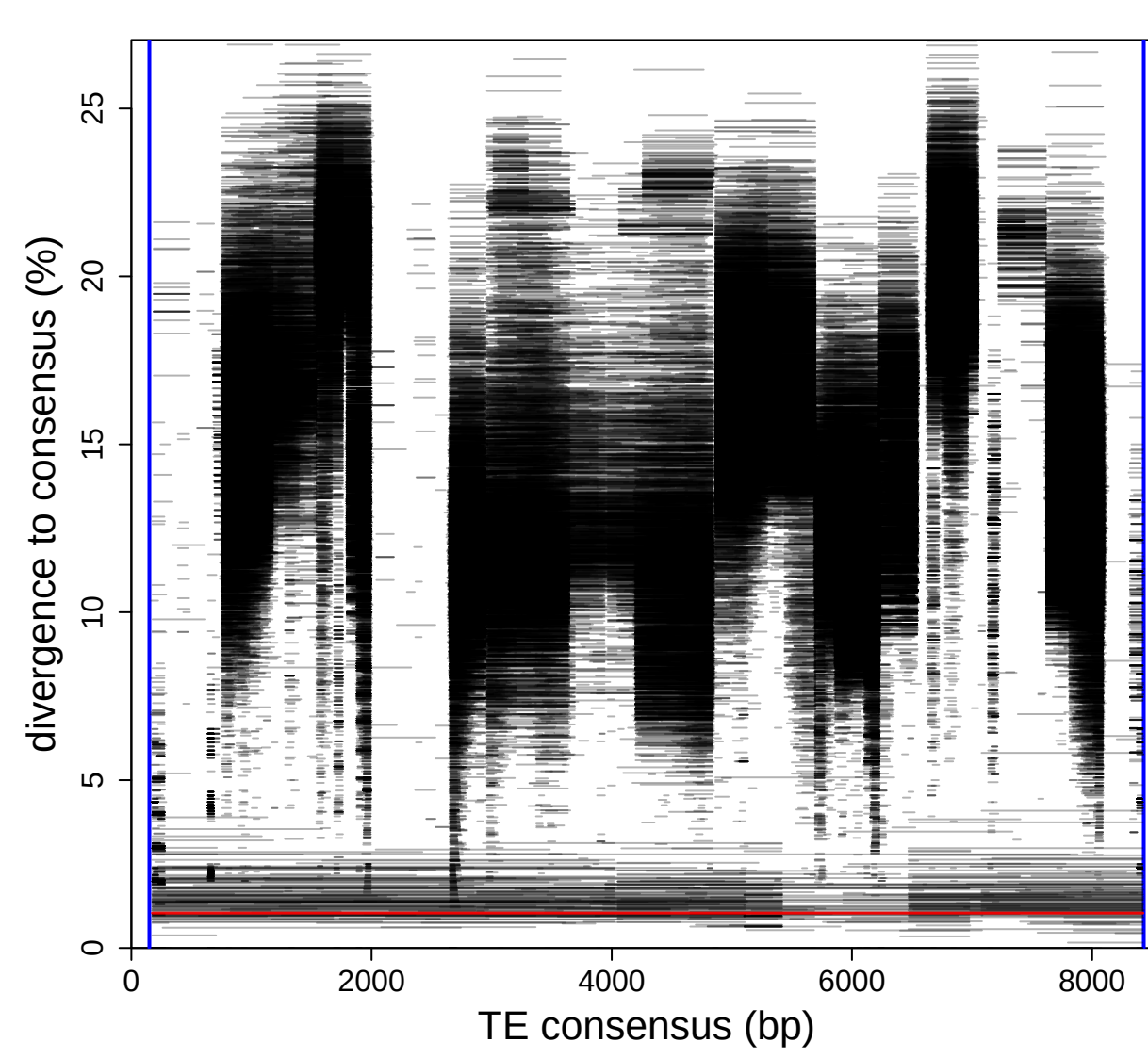
g_1.bed fm_1.bed 0 0 n.bed g_1.bed fm_2.bed 0 0 bcln.fa a
size: 8275bp; fragments: 94695; full length: 1 (>=7447.5bp)

After TETrimmer 8275 bp



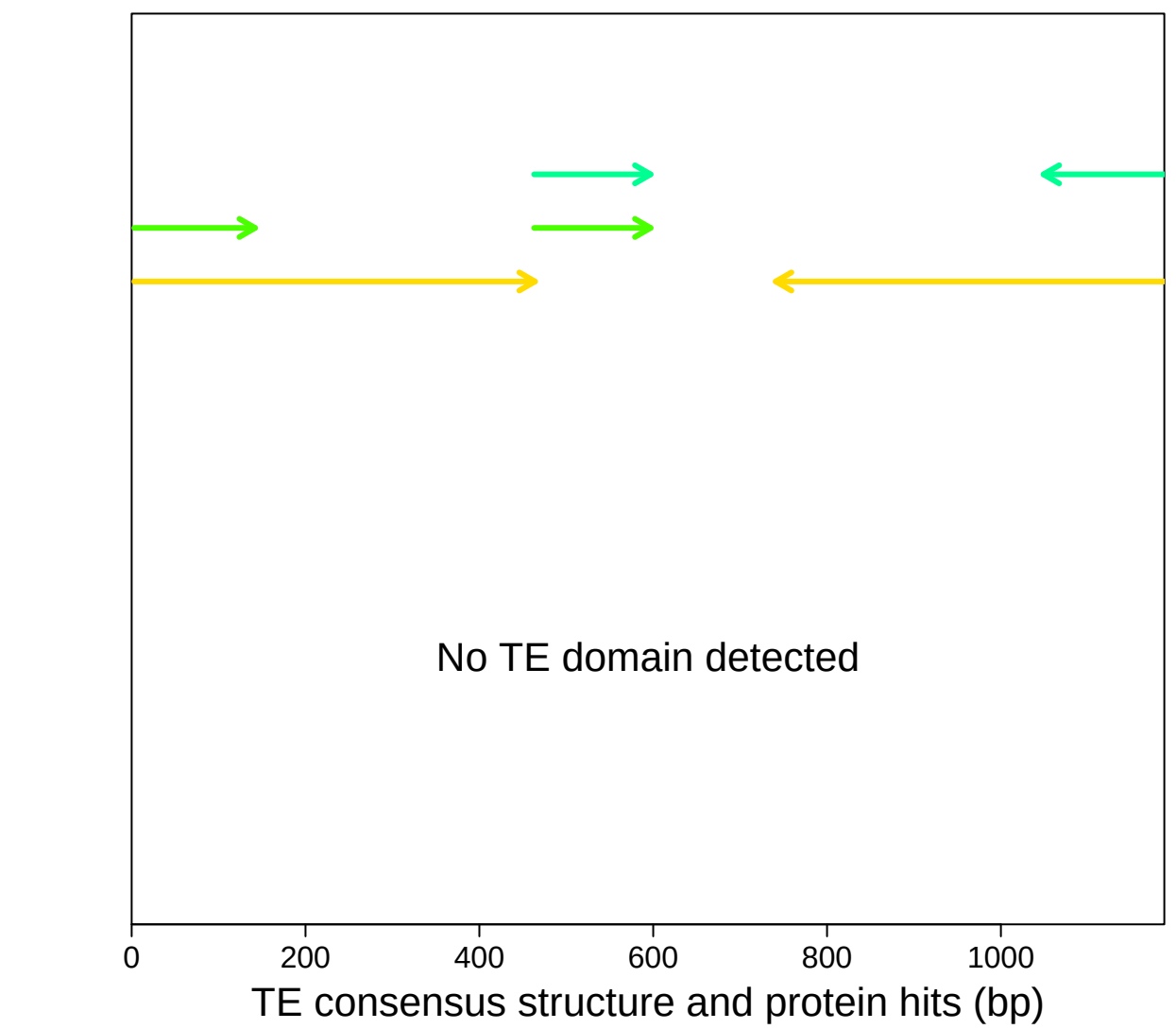
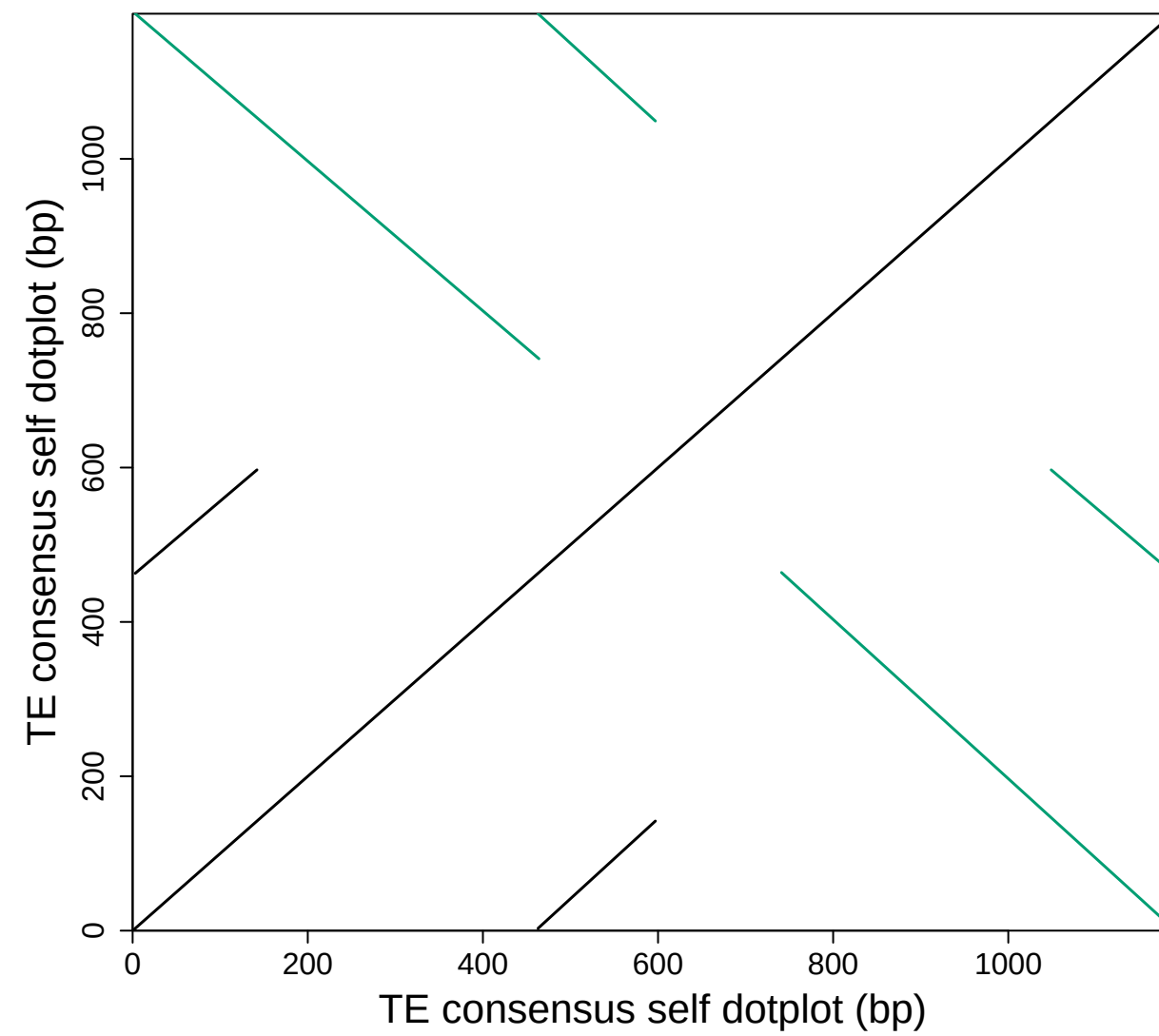
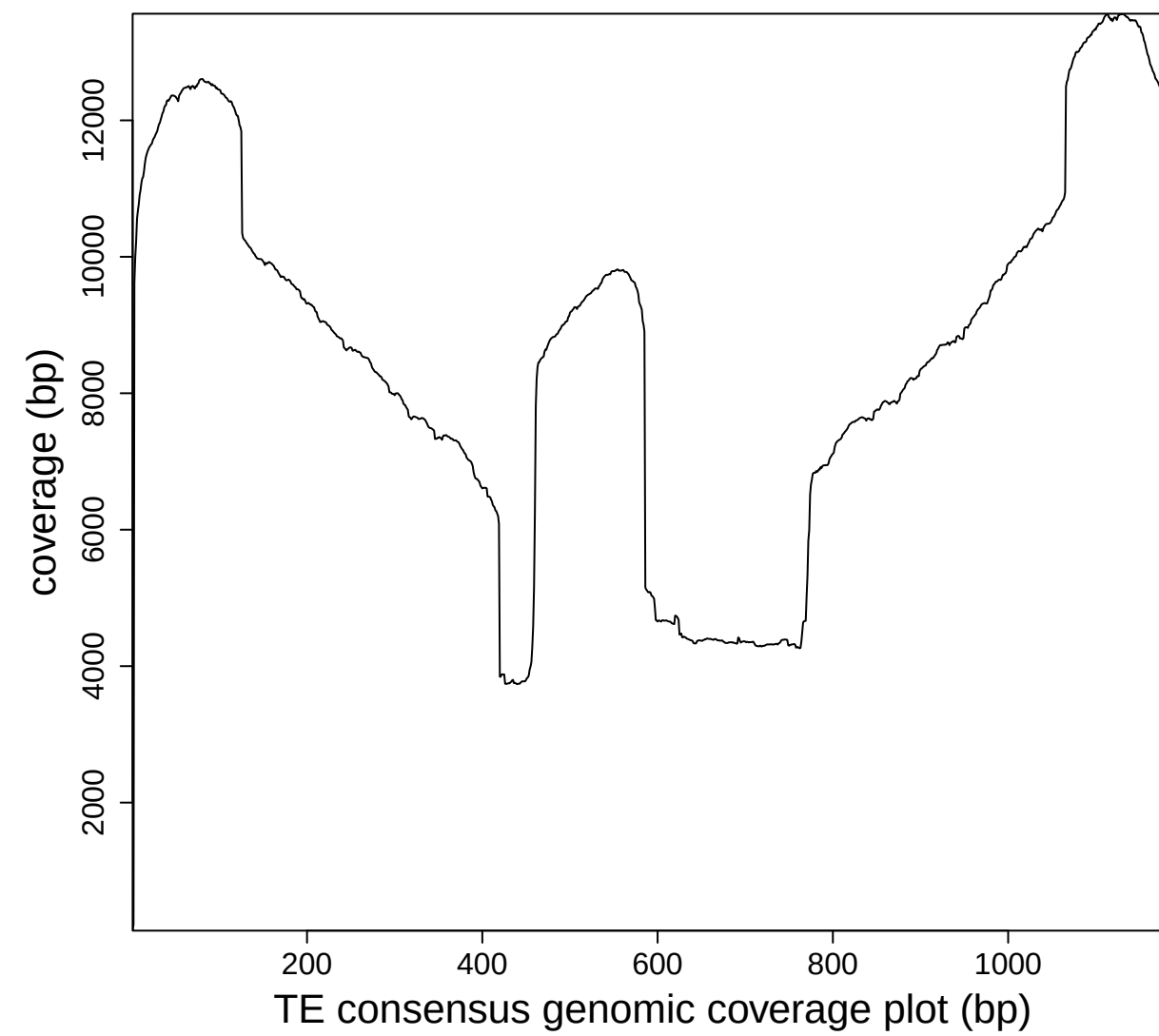
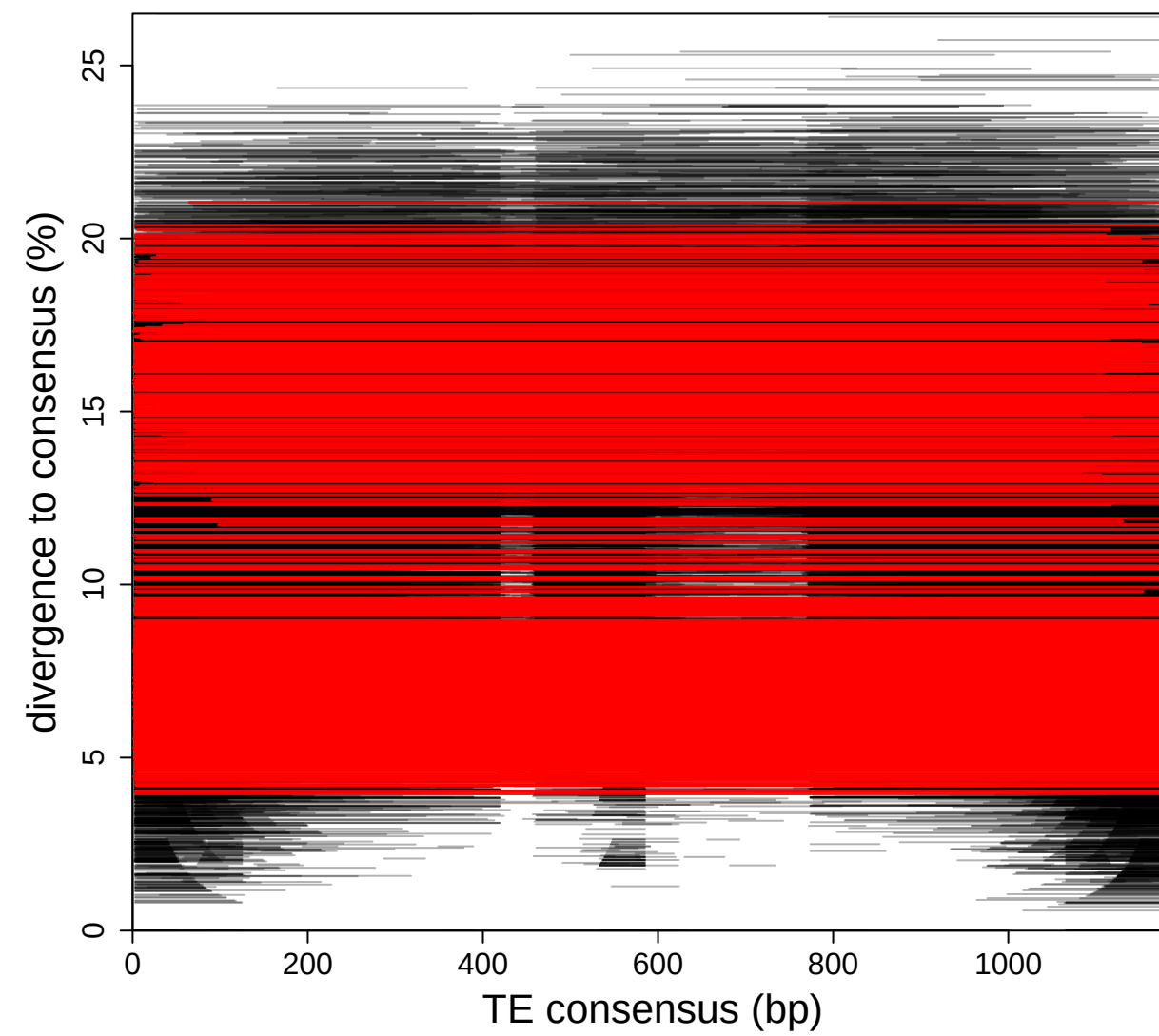
f.bed_g_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcl
size: 8580bp; fragments: 94642; full length: 1 (>=7722bp)

After TEtrimmer Extended plot Blue lines are boundaries

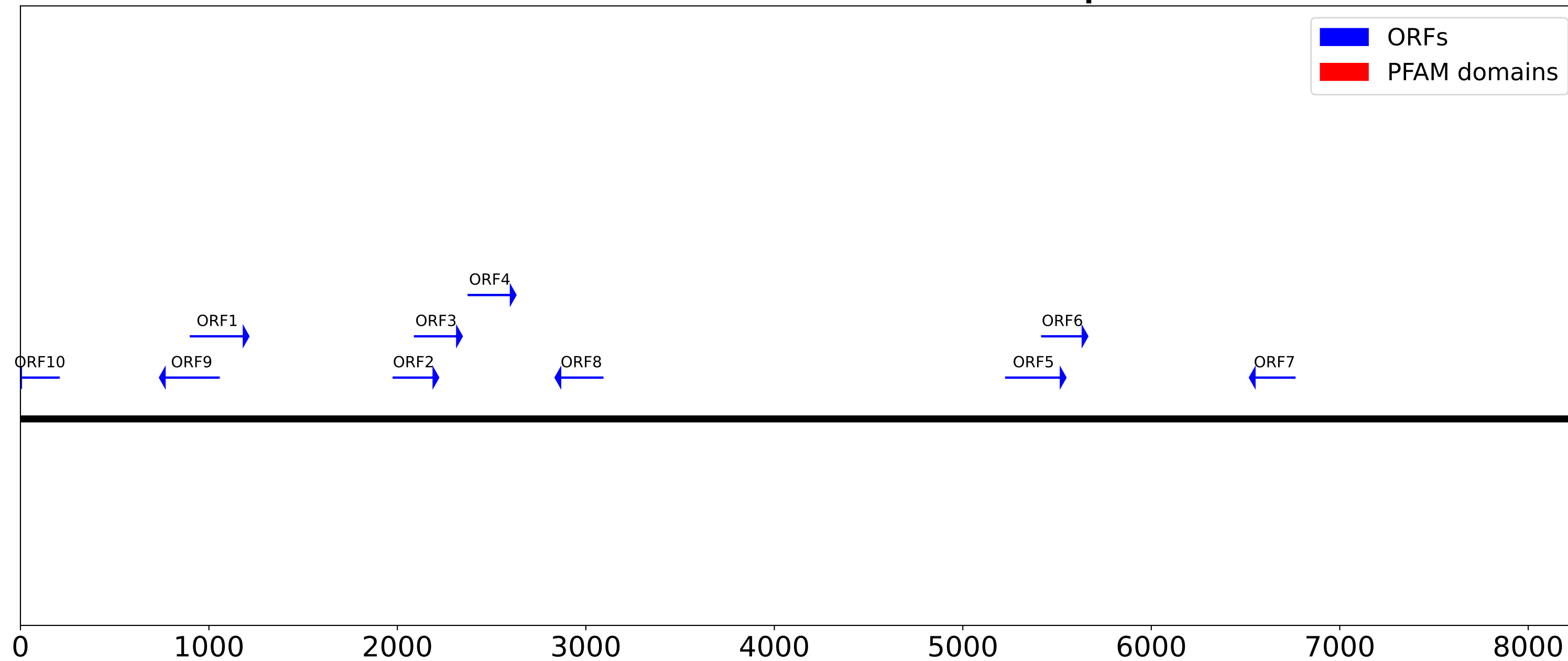


TE: rnd_1_family_137
size: 1188bp; fragments: 36317; full length: 1047 (≥ 1069.2 bp)

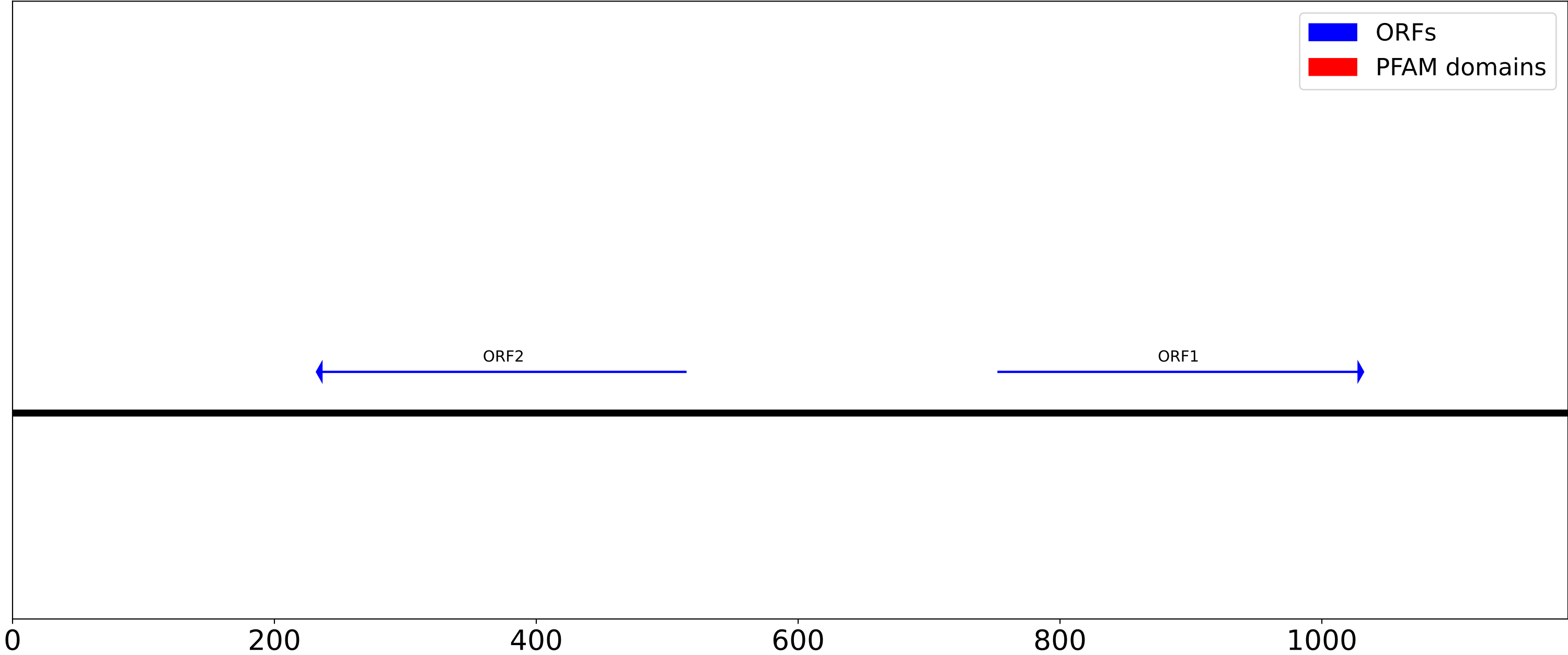
Before TEtrimmer 1188 bp



After TEtrimmer ORF and PFAM domain plot



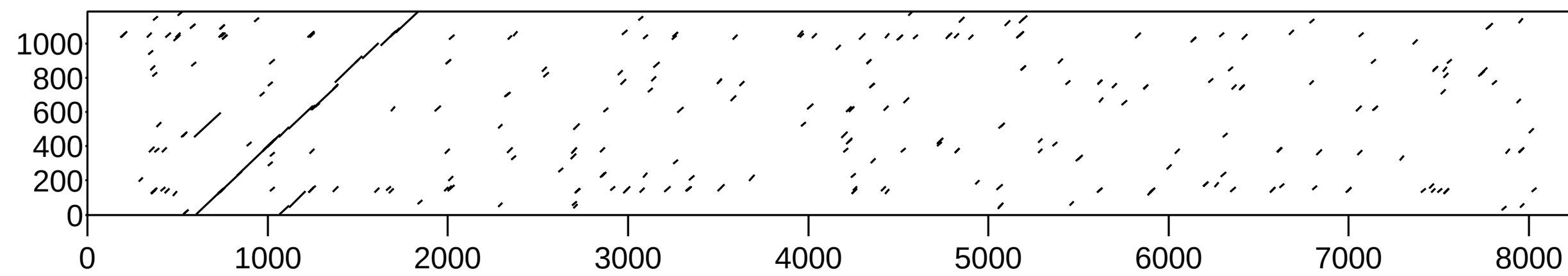
Before TEtrimmer ORF and PFAM domain plot



Dotplot

(windowsize = 25, threshold = 50.00 09/10/25)

isensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)