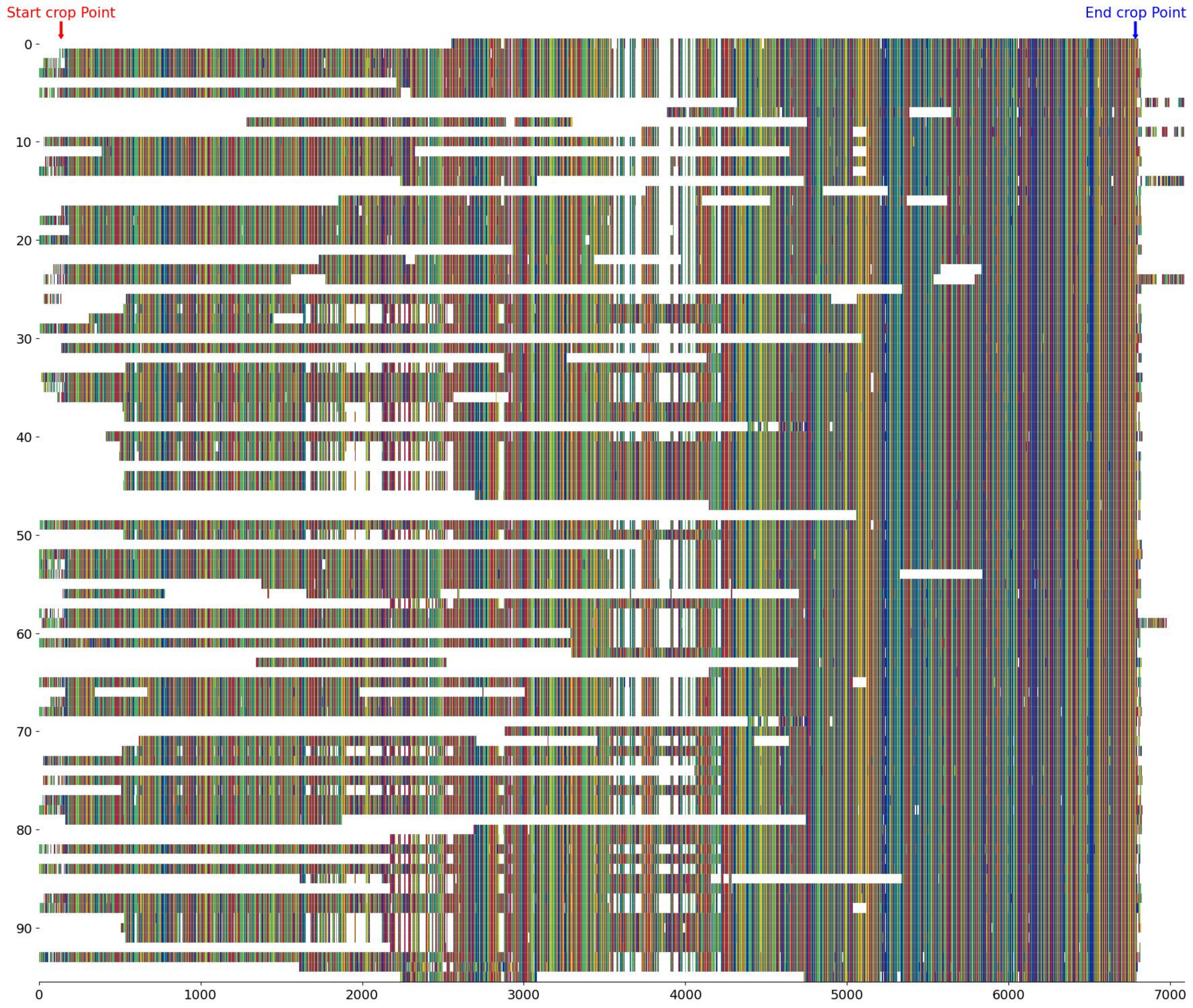
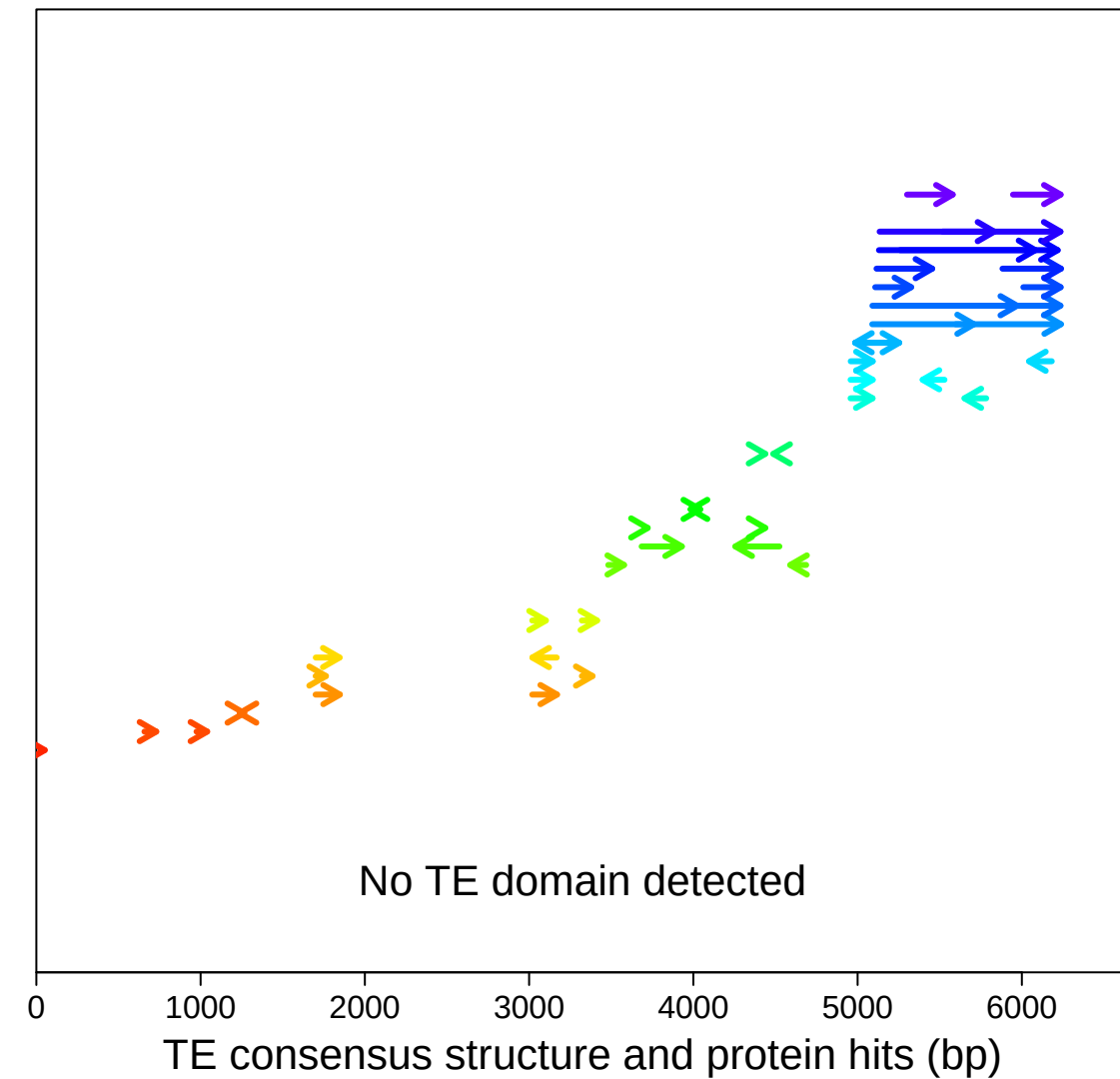
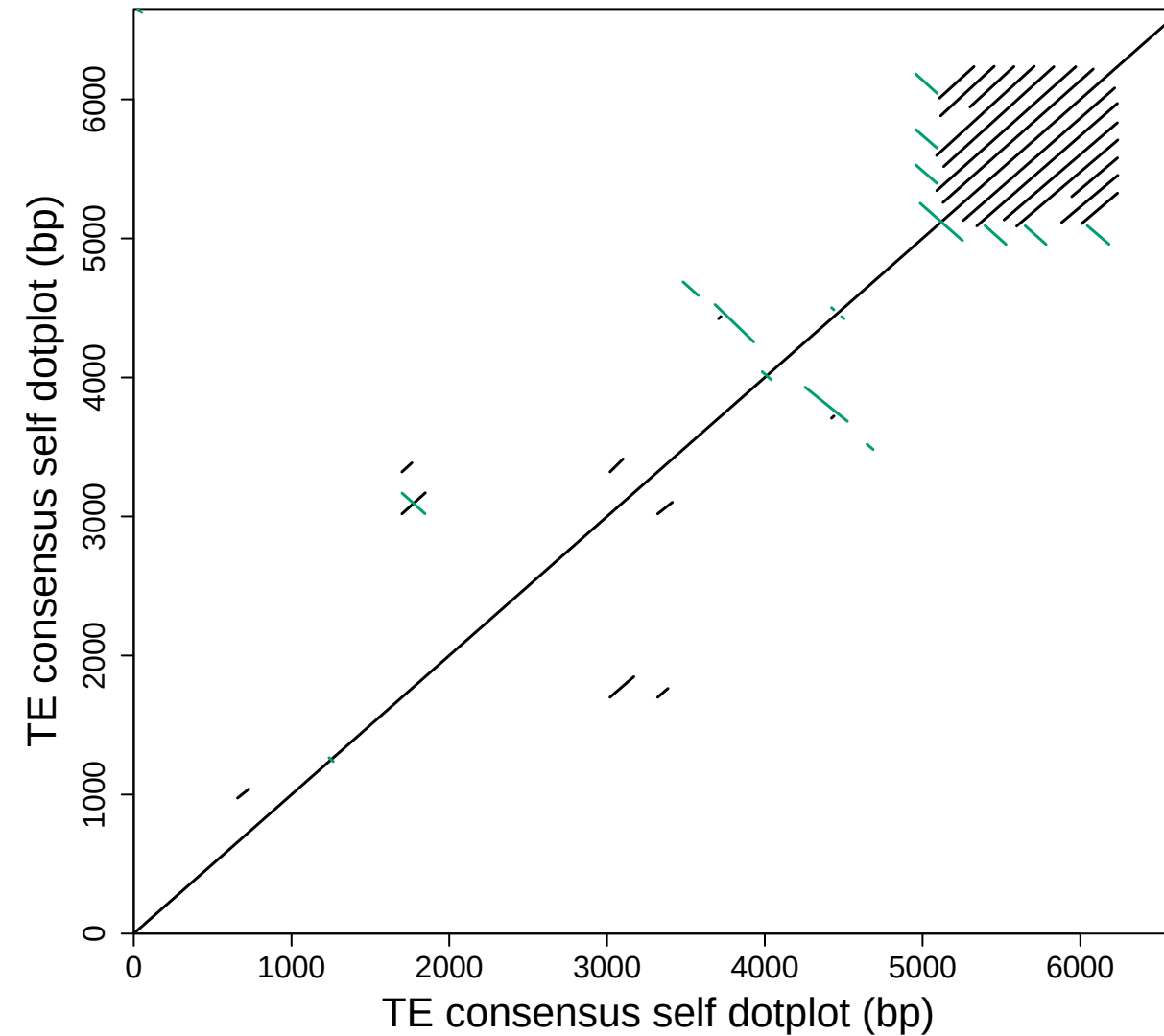
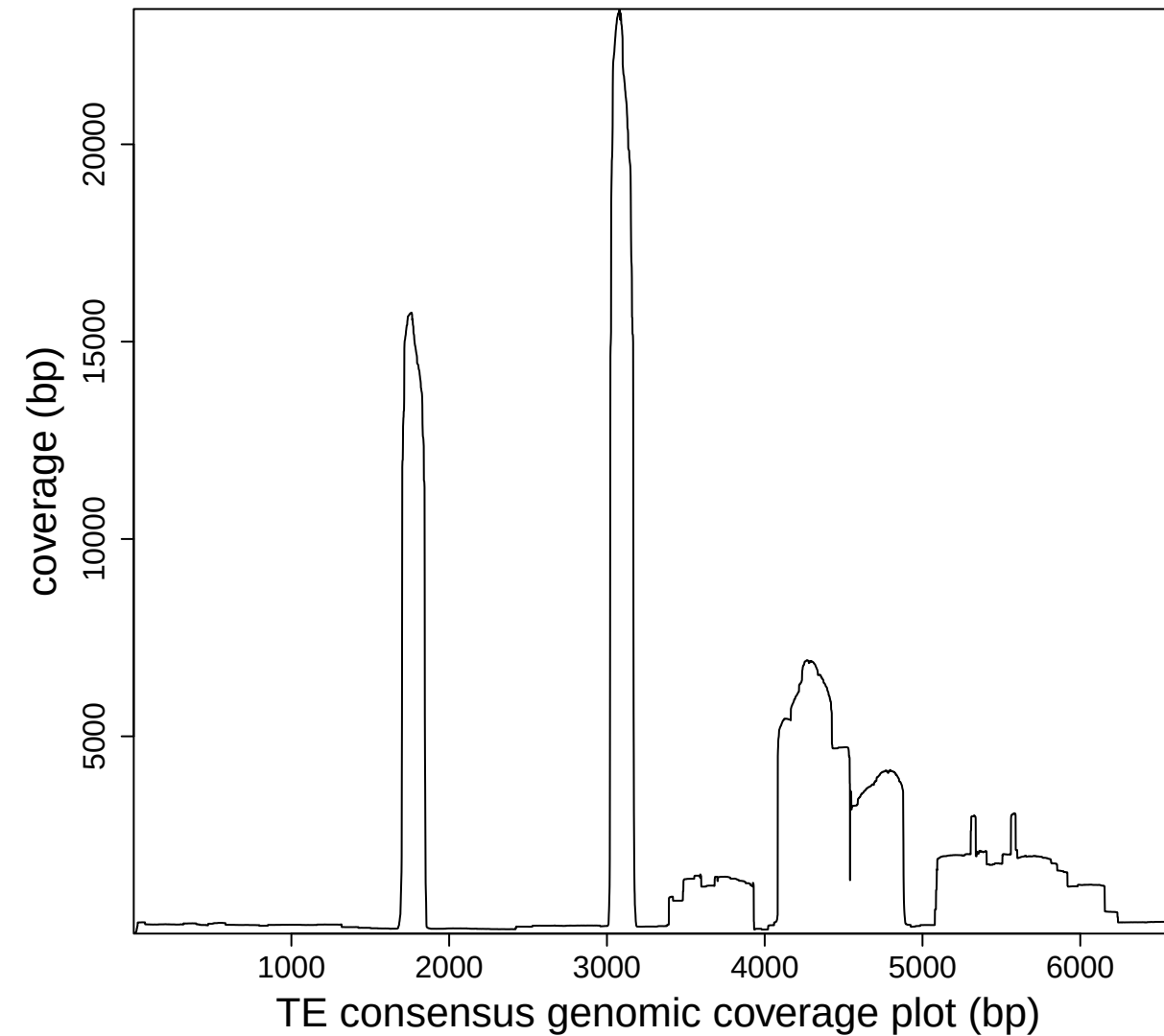
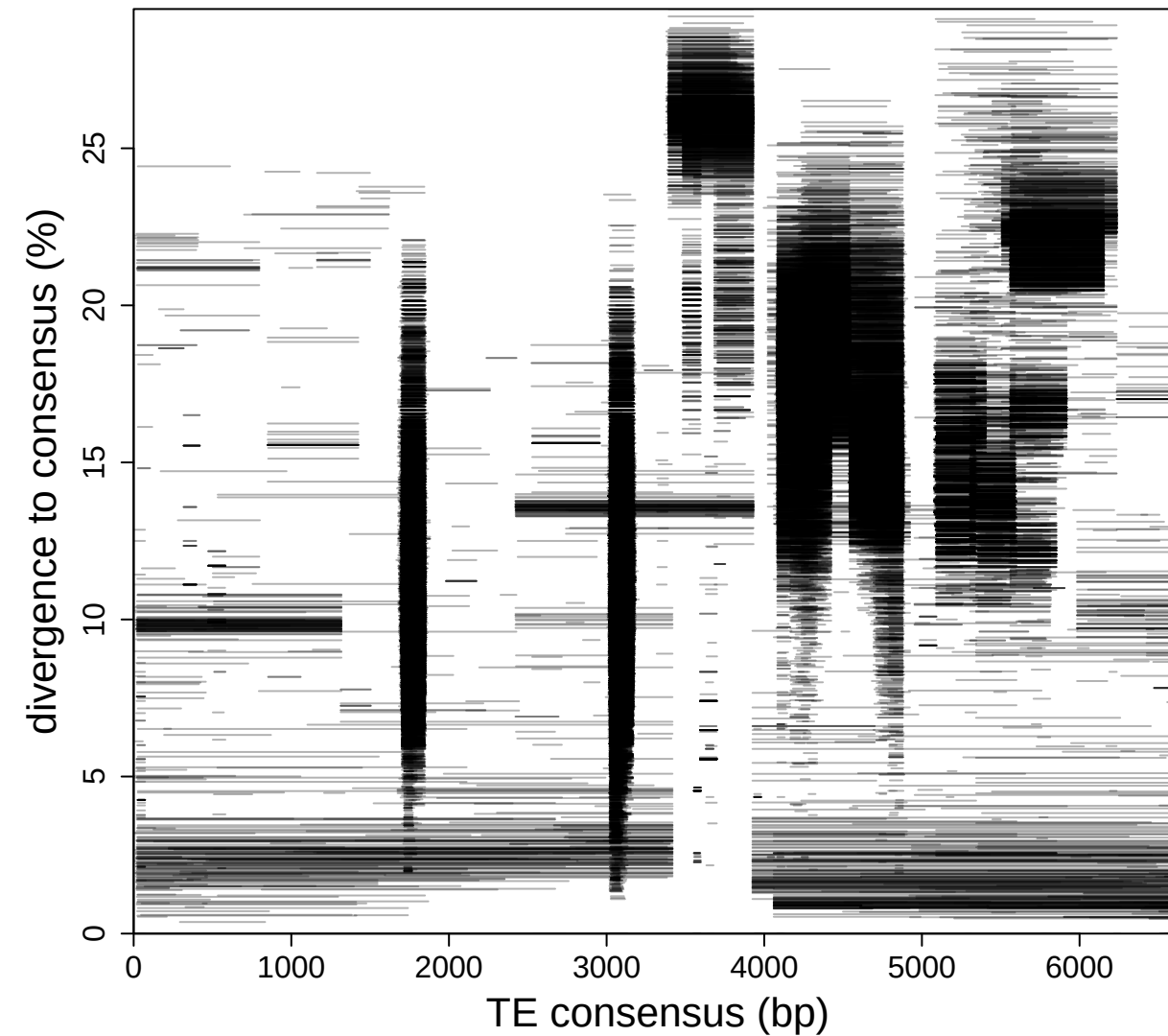






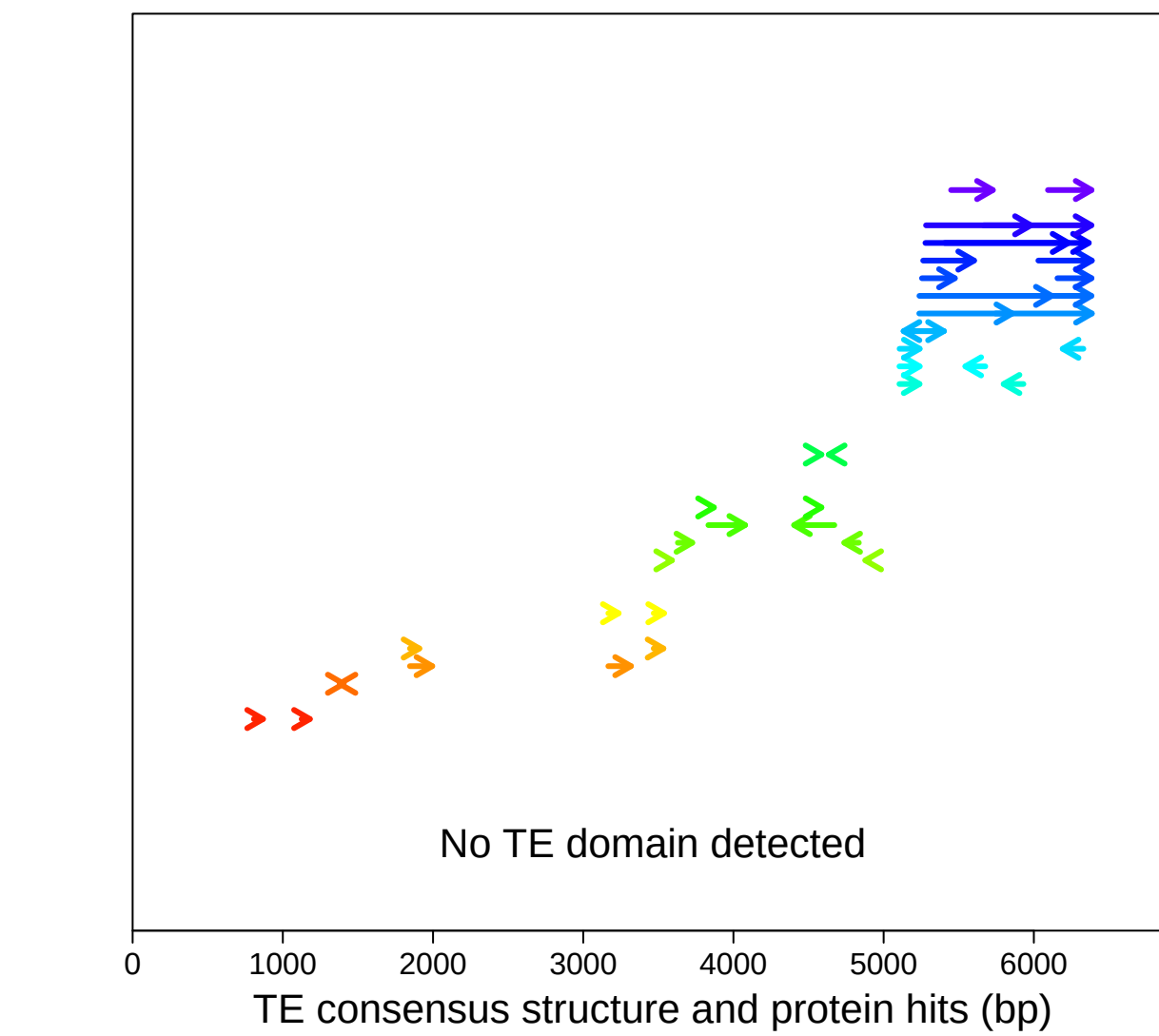
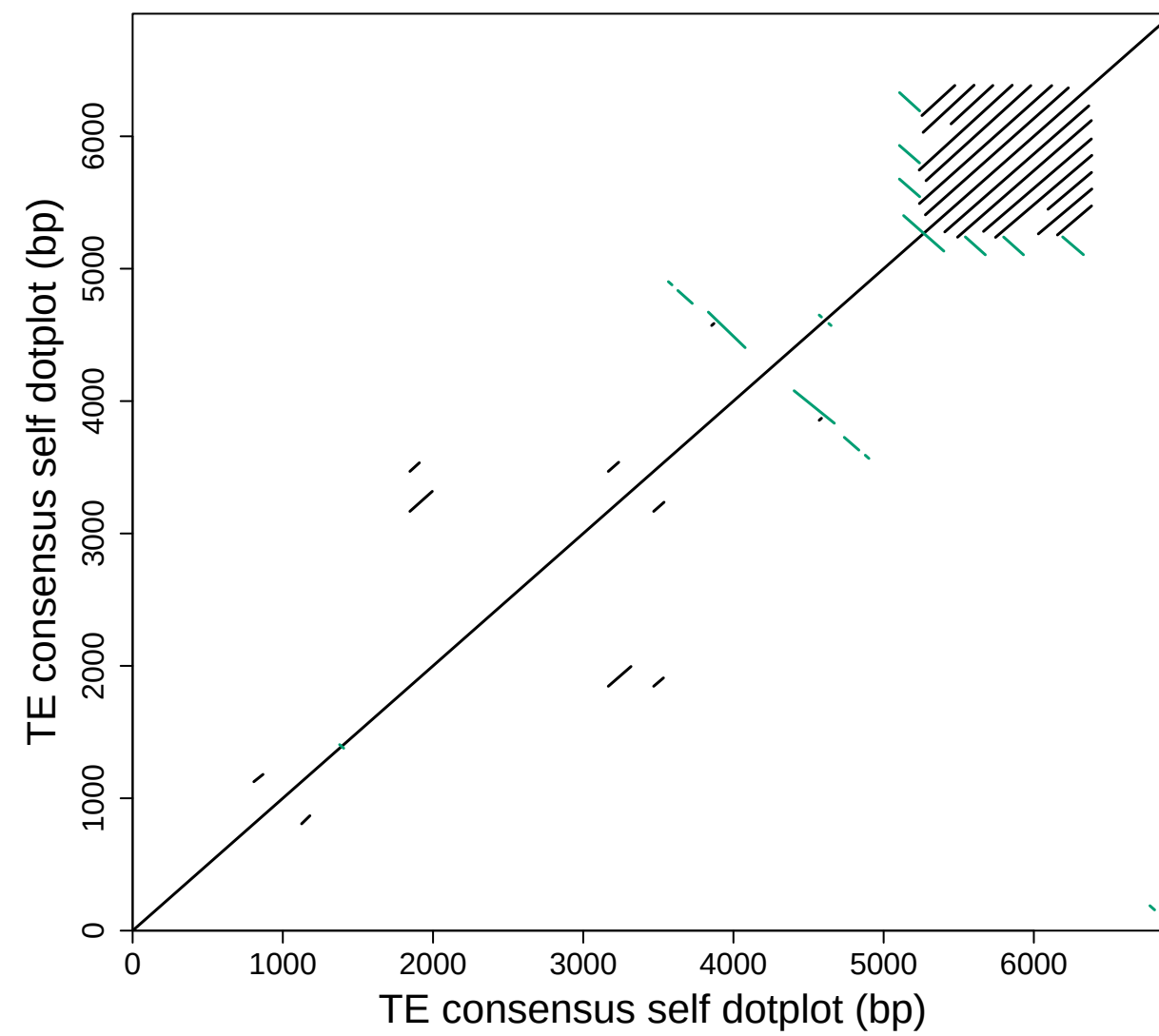
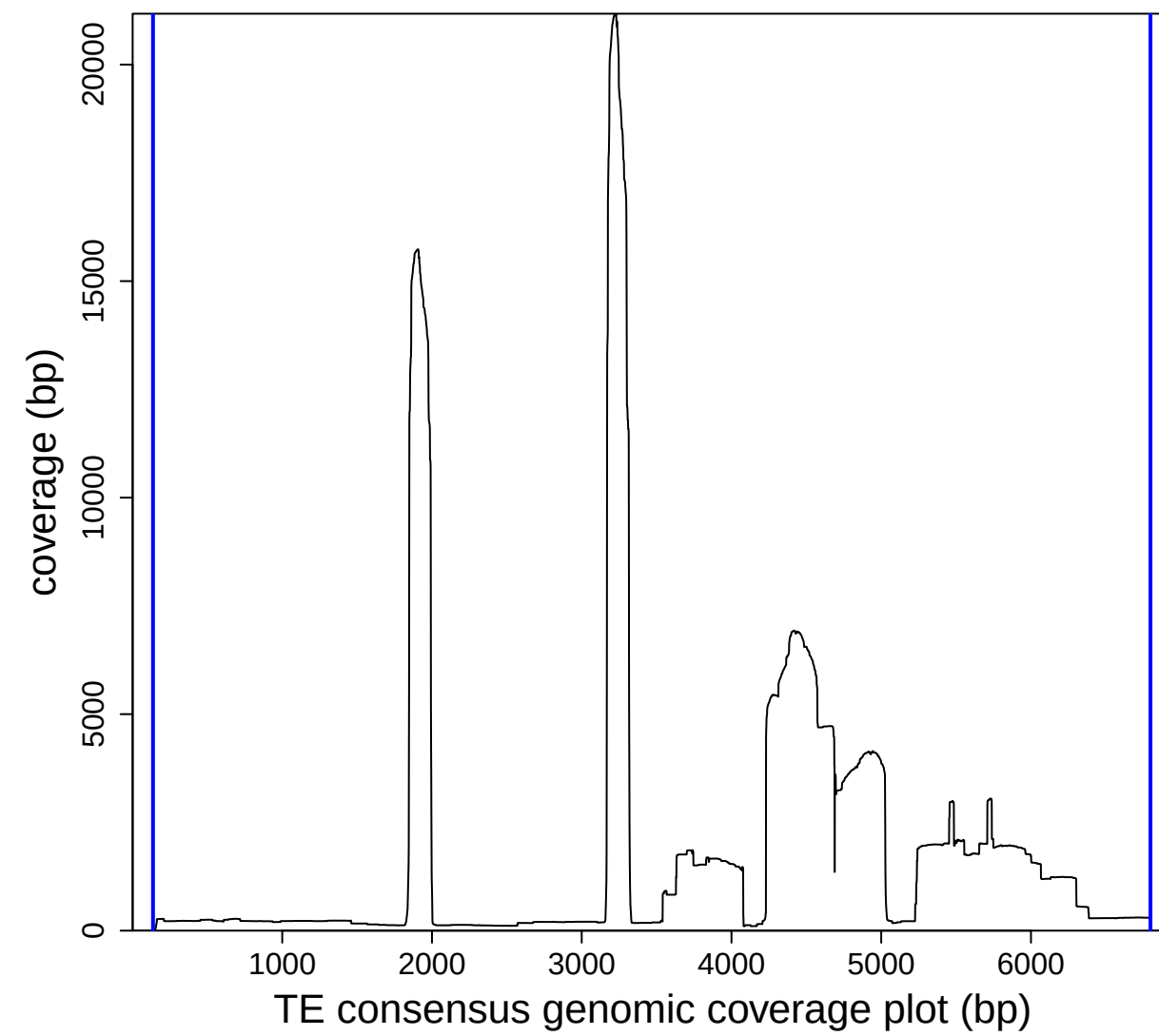
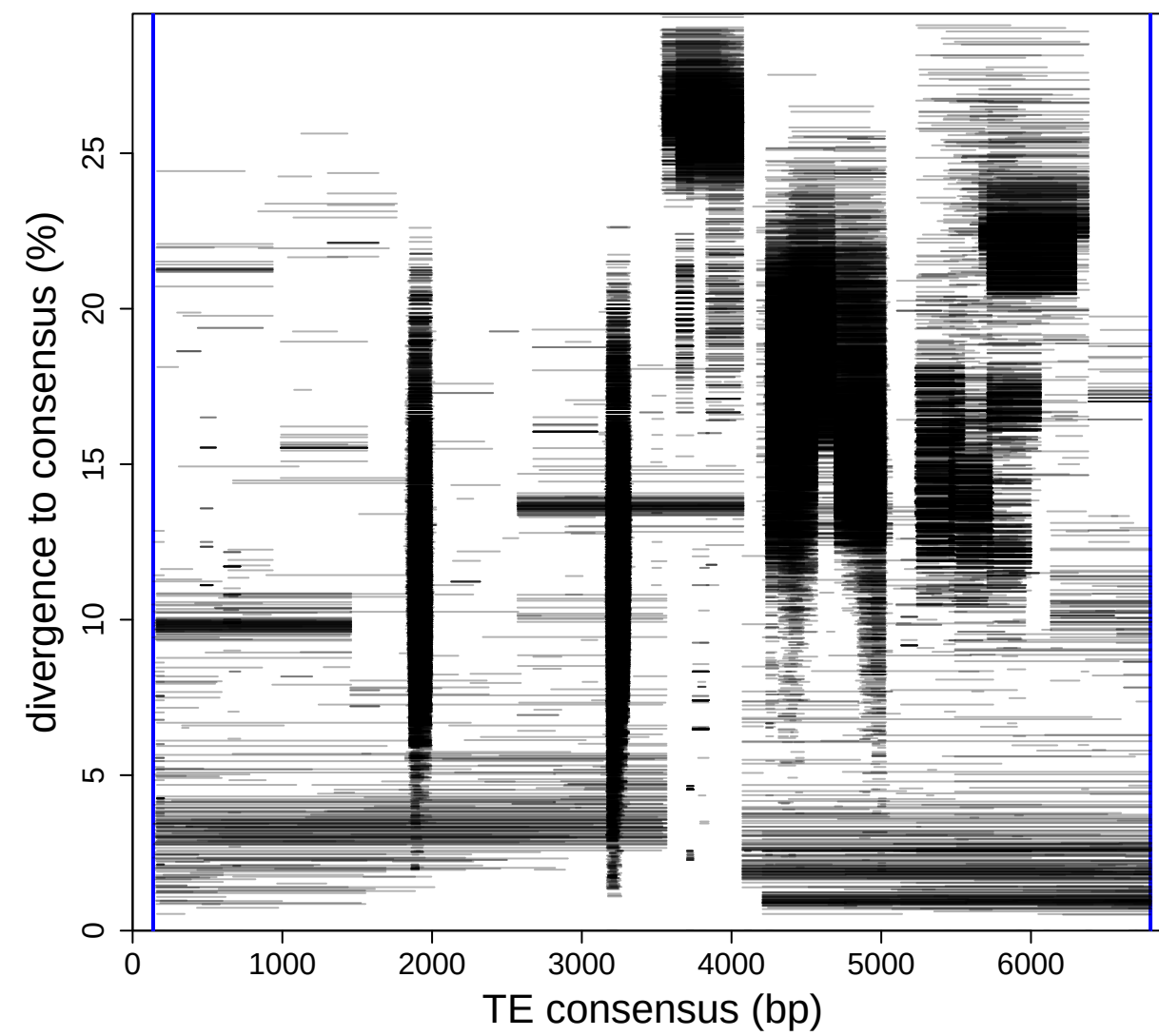
g_1.bed fm_1.bed 0_0 n.bed g_1.bed fm_2.bed 0_0 bcln.fa a
size: 6650bp; fragments: 58092; full length: 0 (>=5985bp)

After TEtrimmer 6650 bp



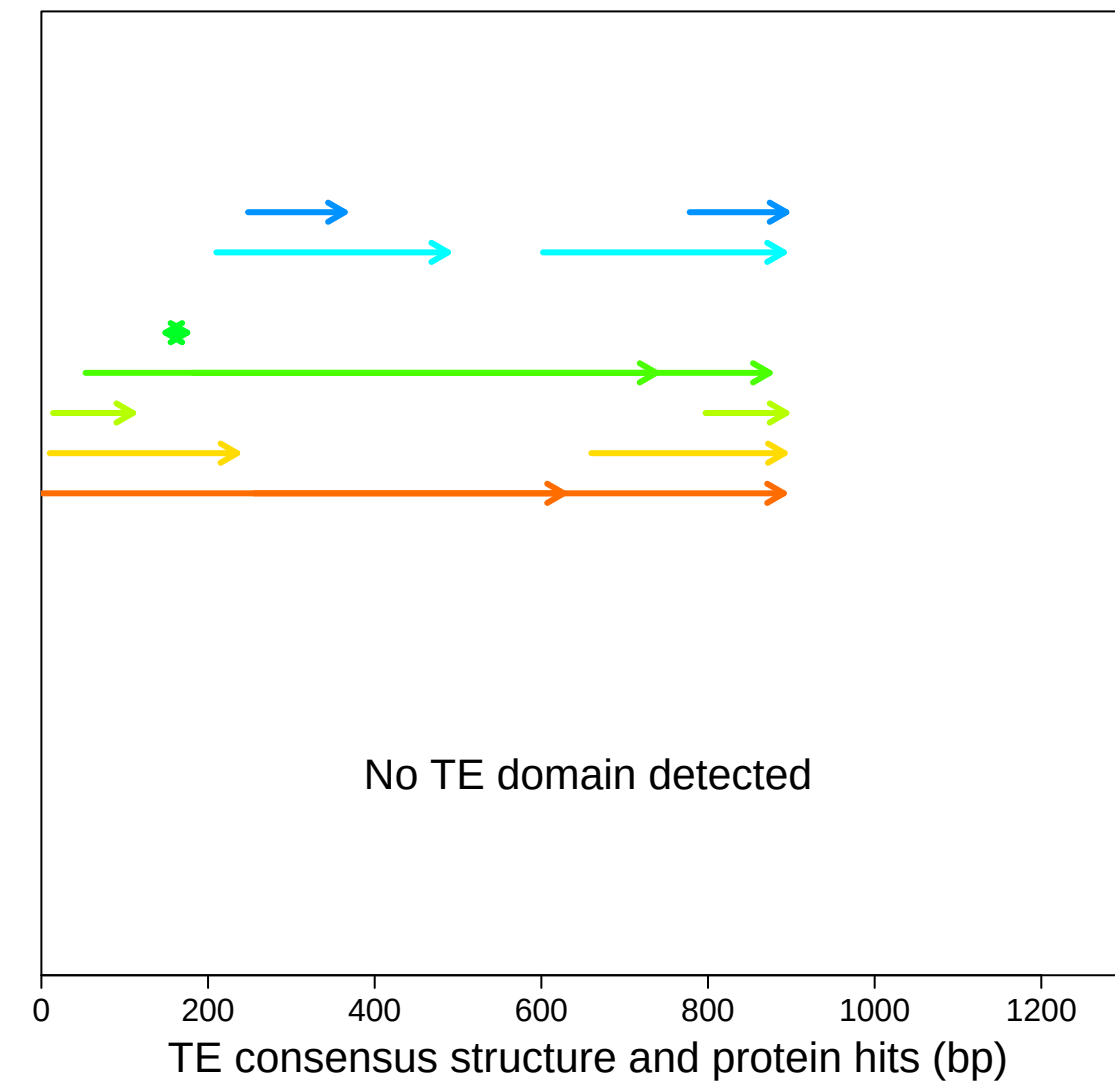
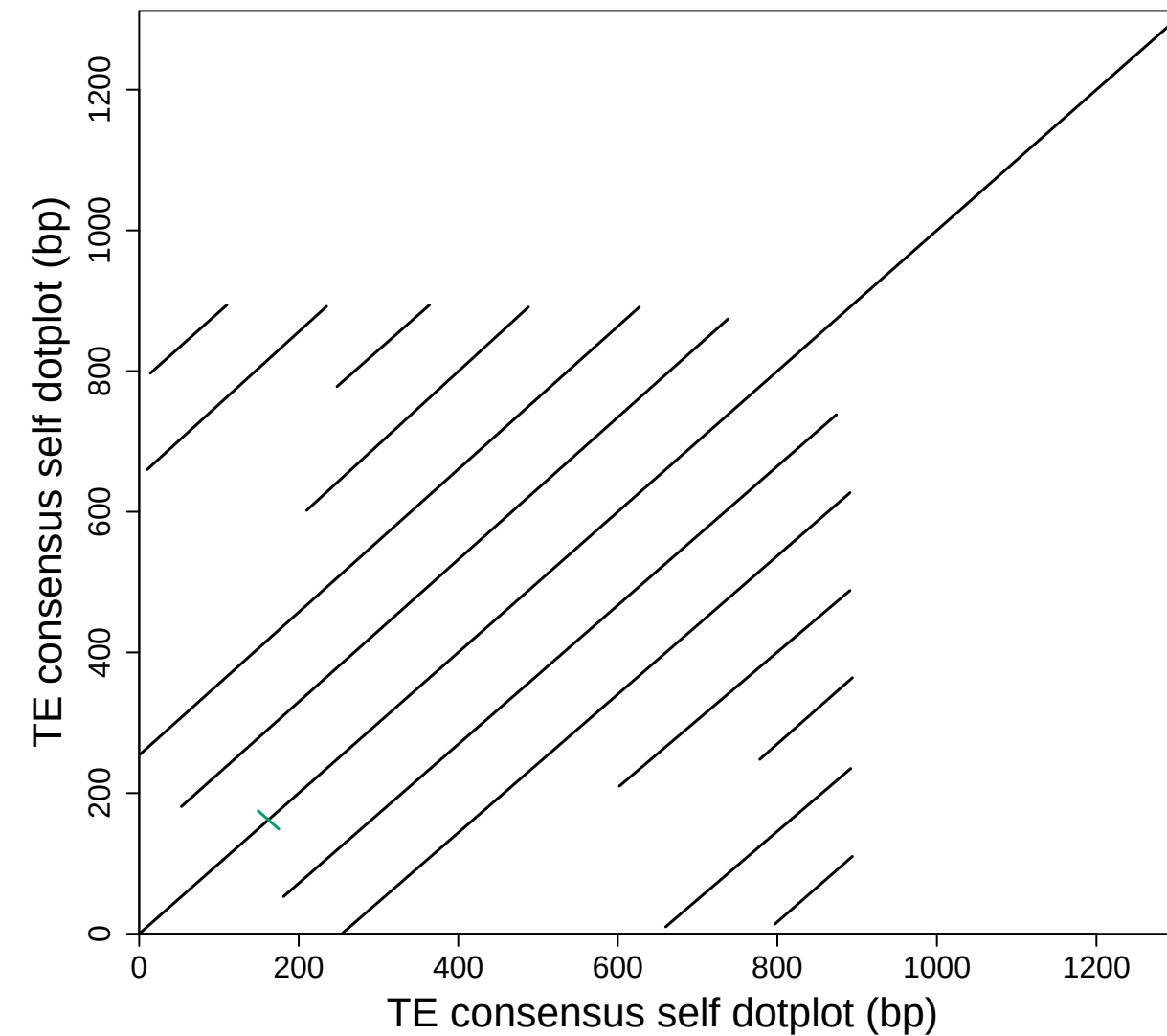
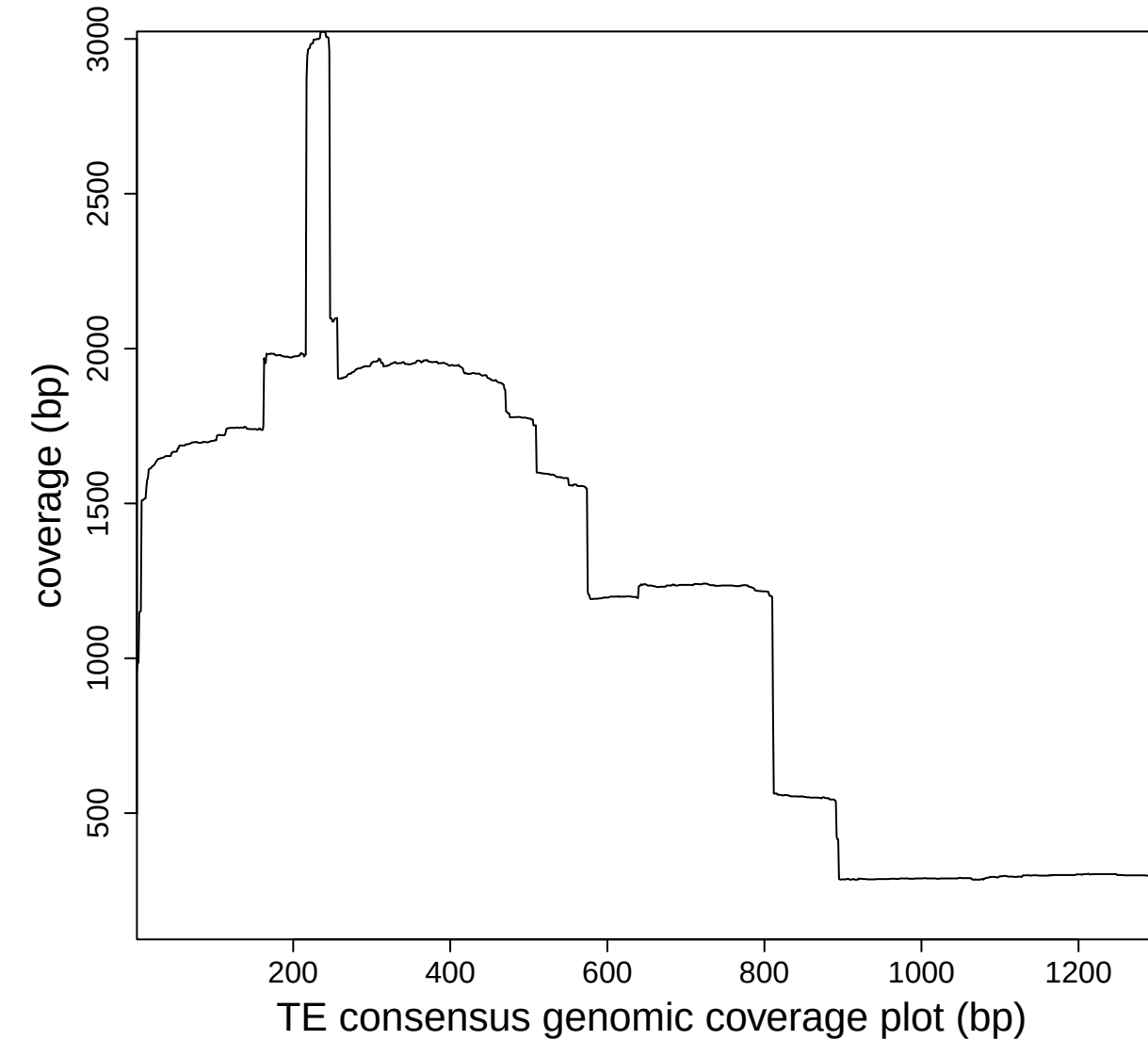
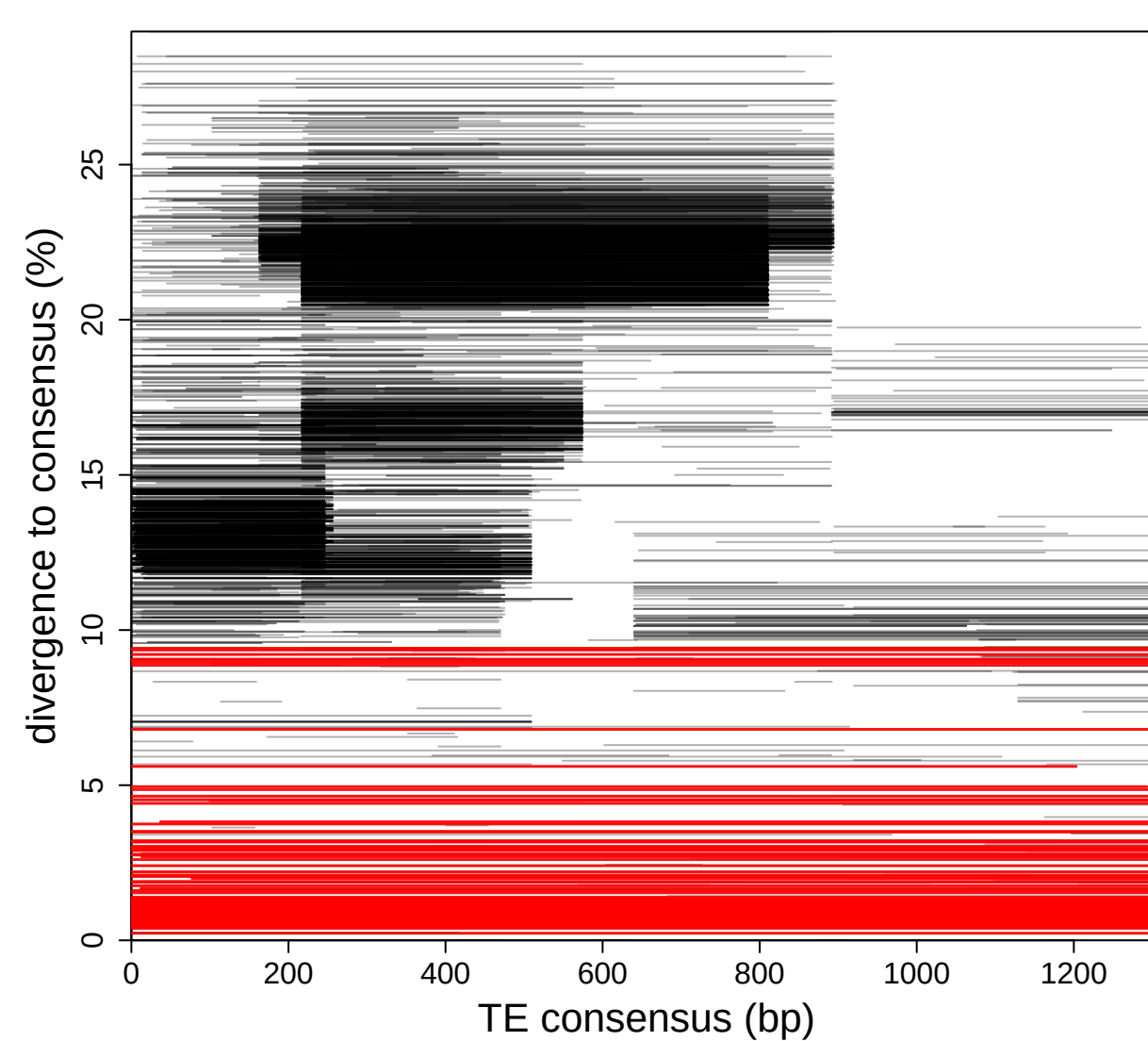
f.bed_g_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcl
size: 6948bp; fragments: 55912; full length: 0 (>=6253.2bp)

After TEtrimmer Extended plot Blue lines are boundaries

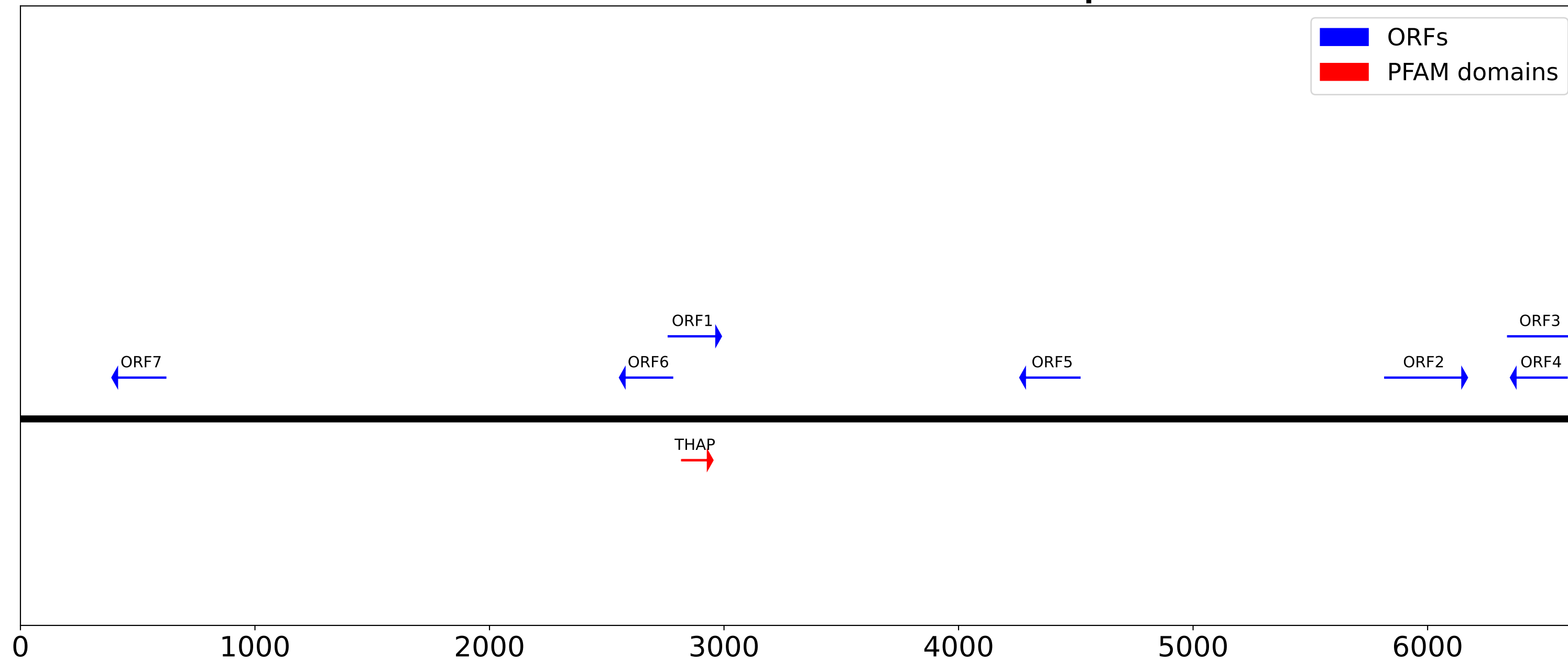


TE: rnd_1_family_615
size: 1312bp; fragments: 3466; full length: 167 (≥ 1180.8 bp)

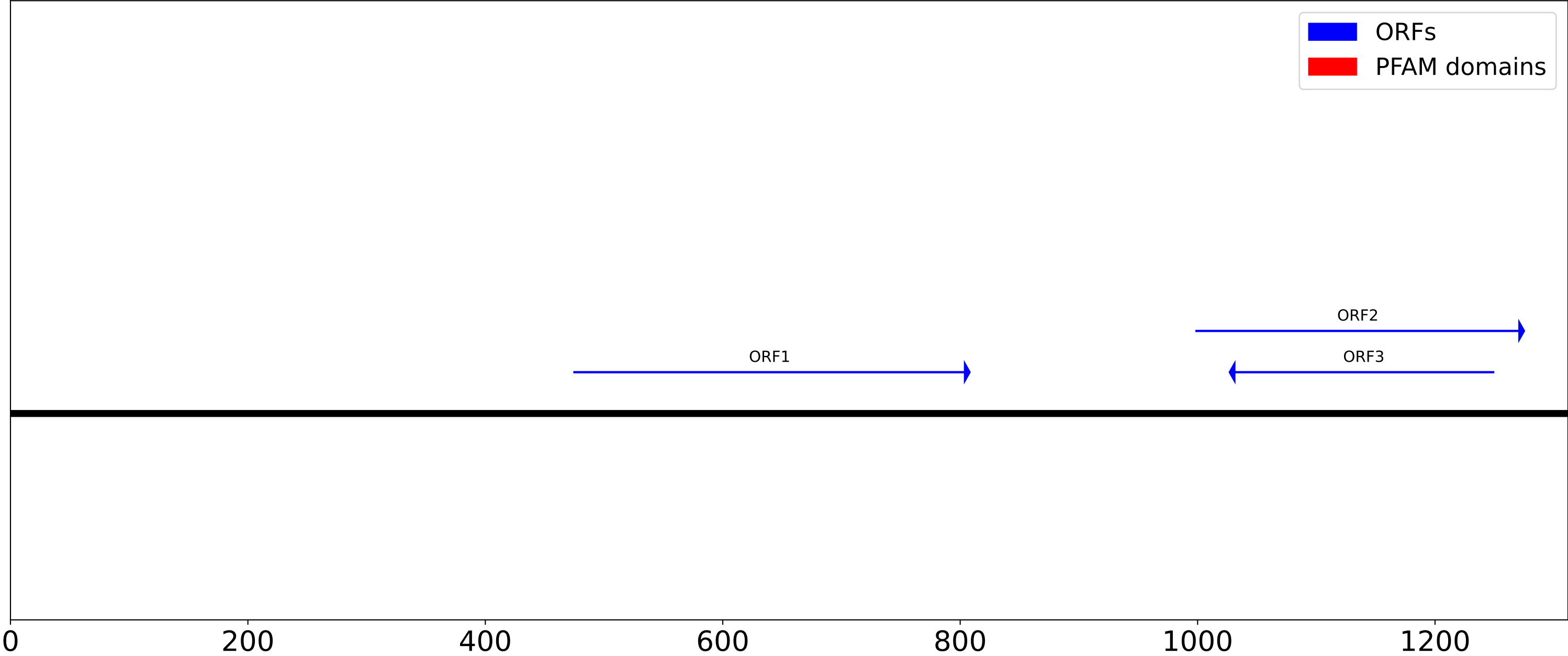
Before TEtrimmer 1312 bp



After TEtrimmer ORF and PFAM domain plot



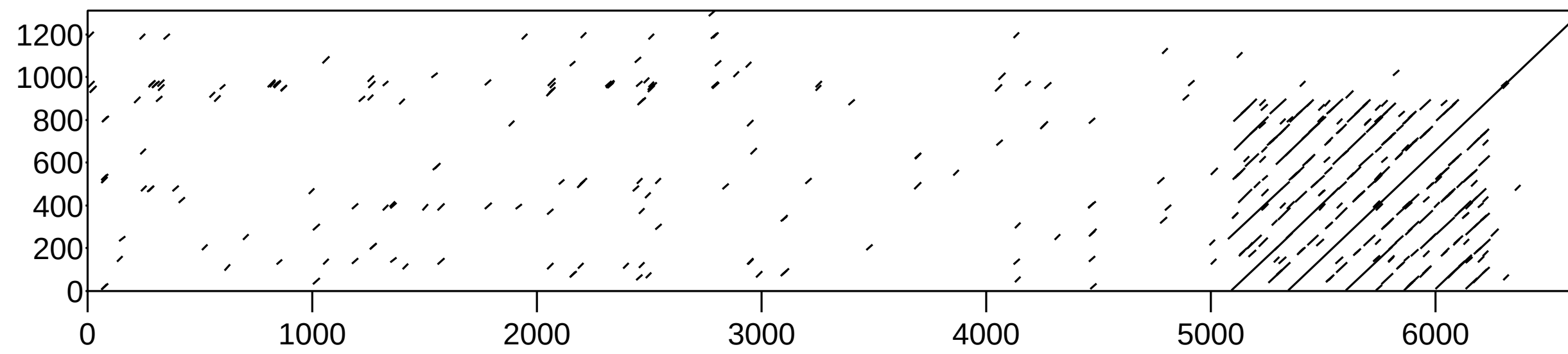
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)