

Start crop Point

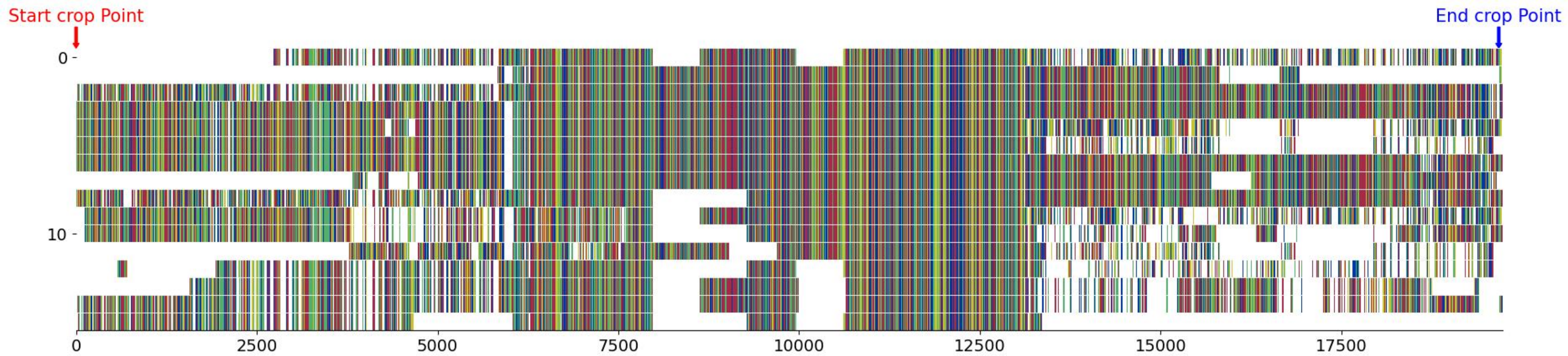


End crop Point



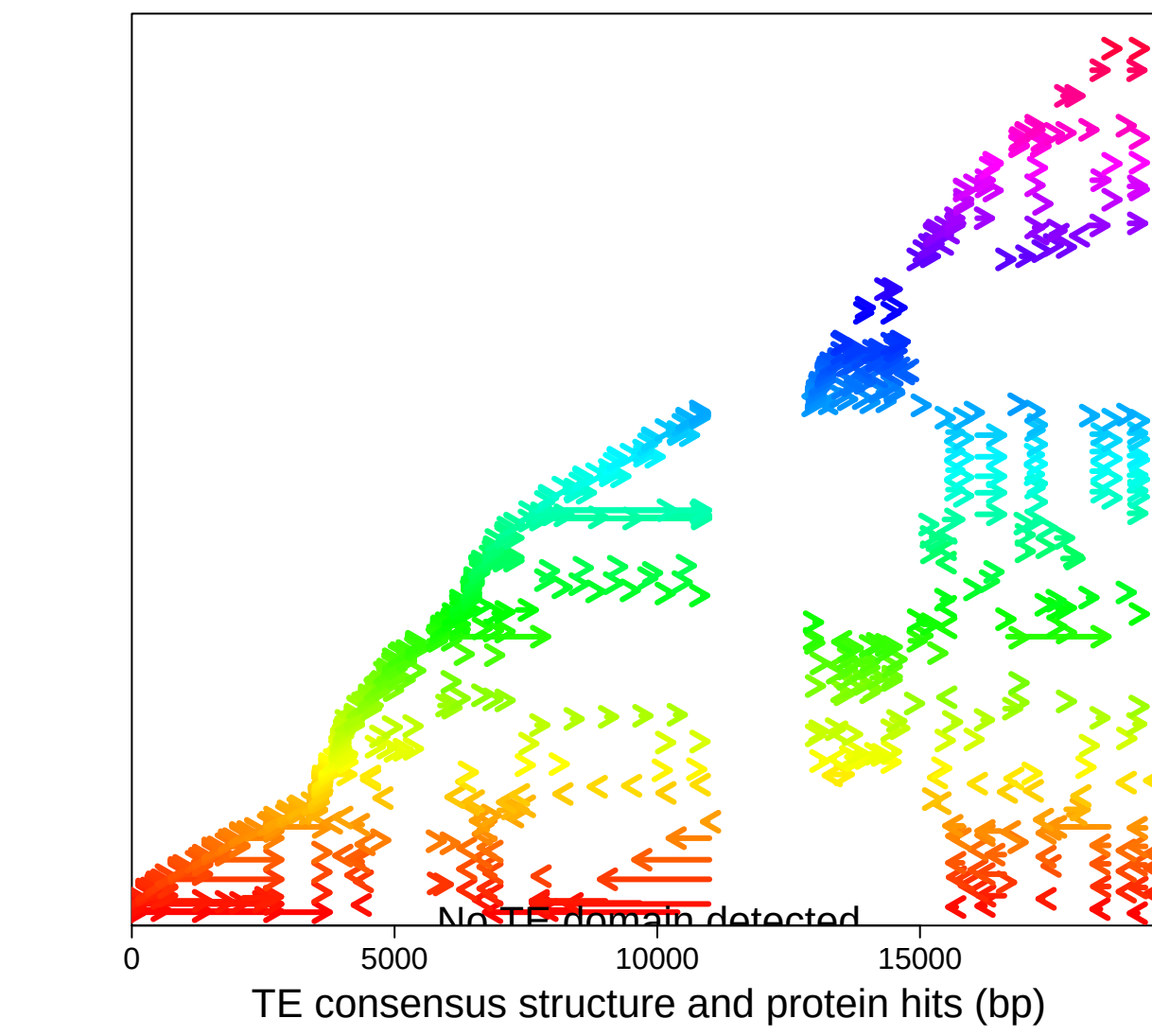
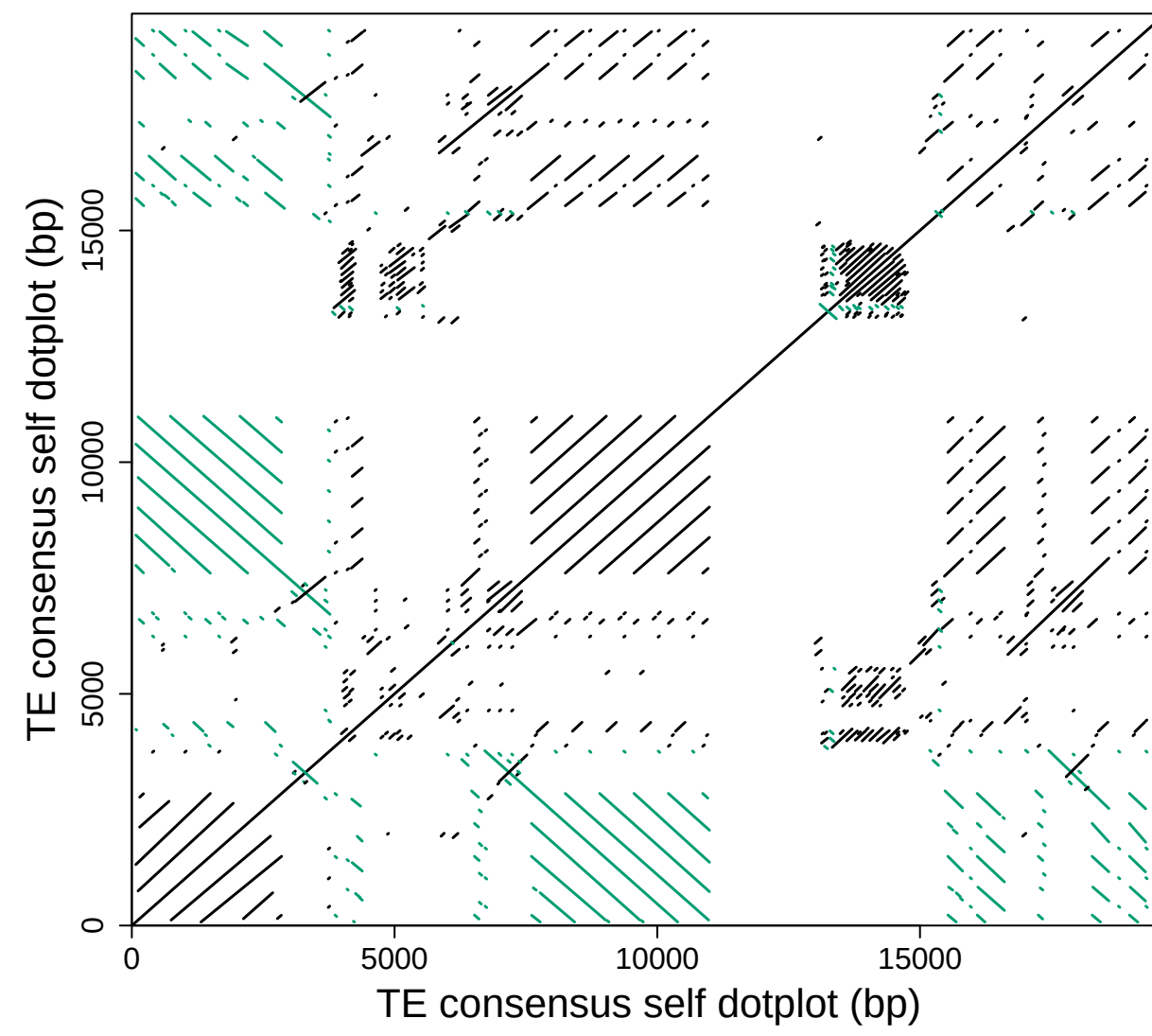
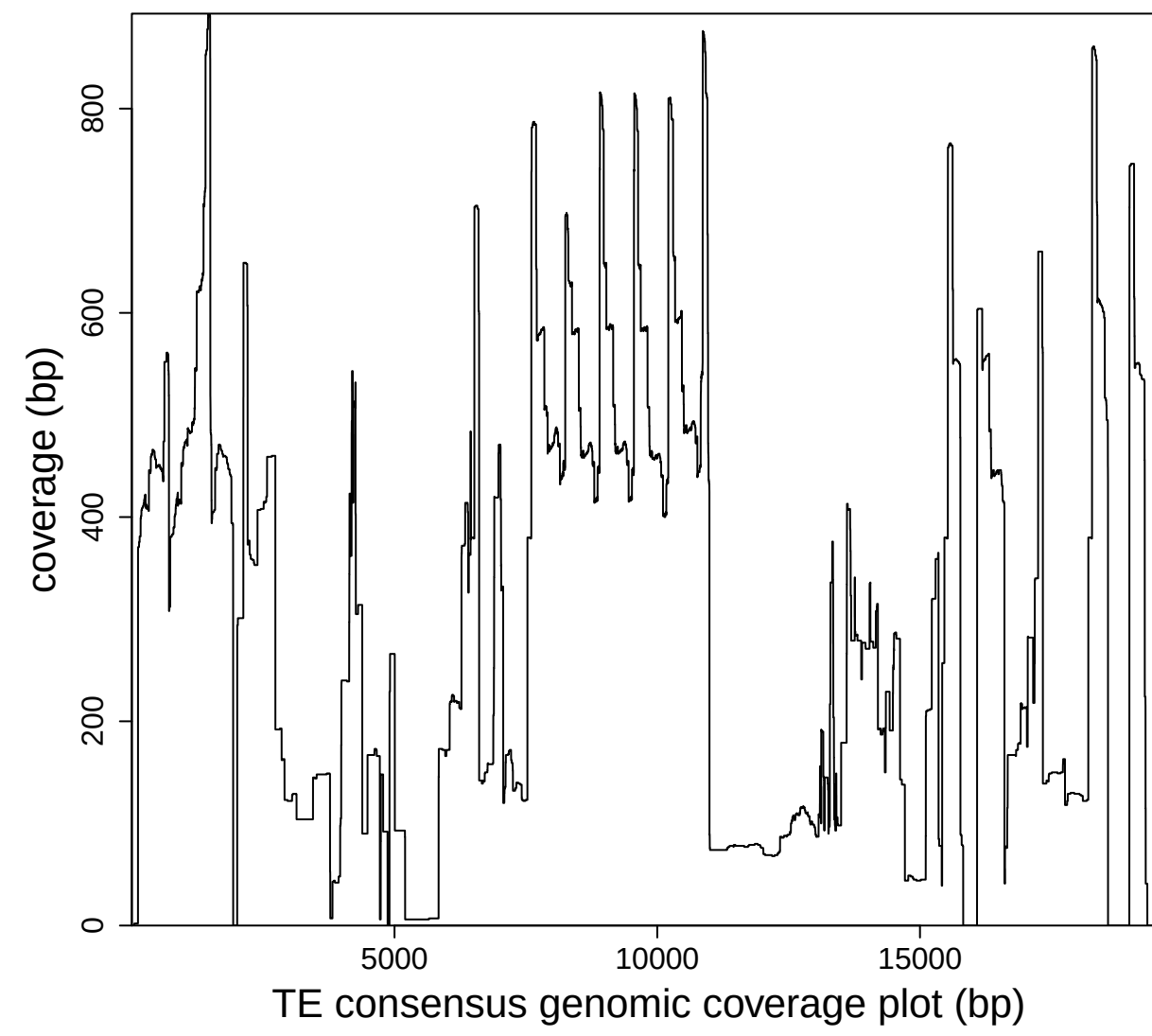
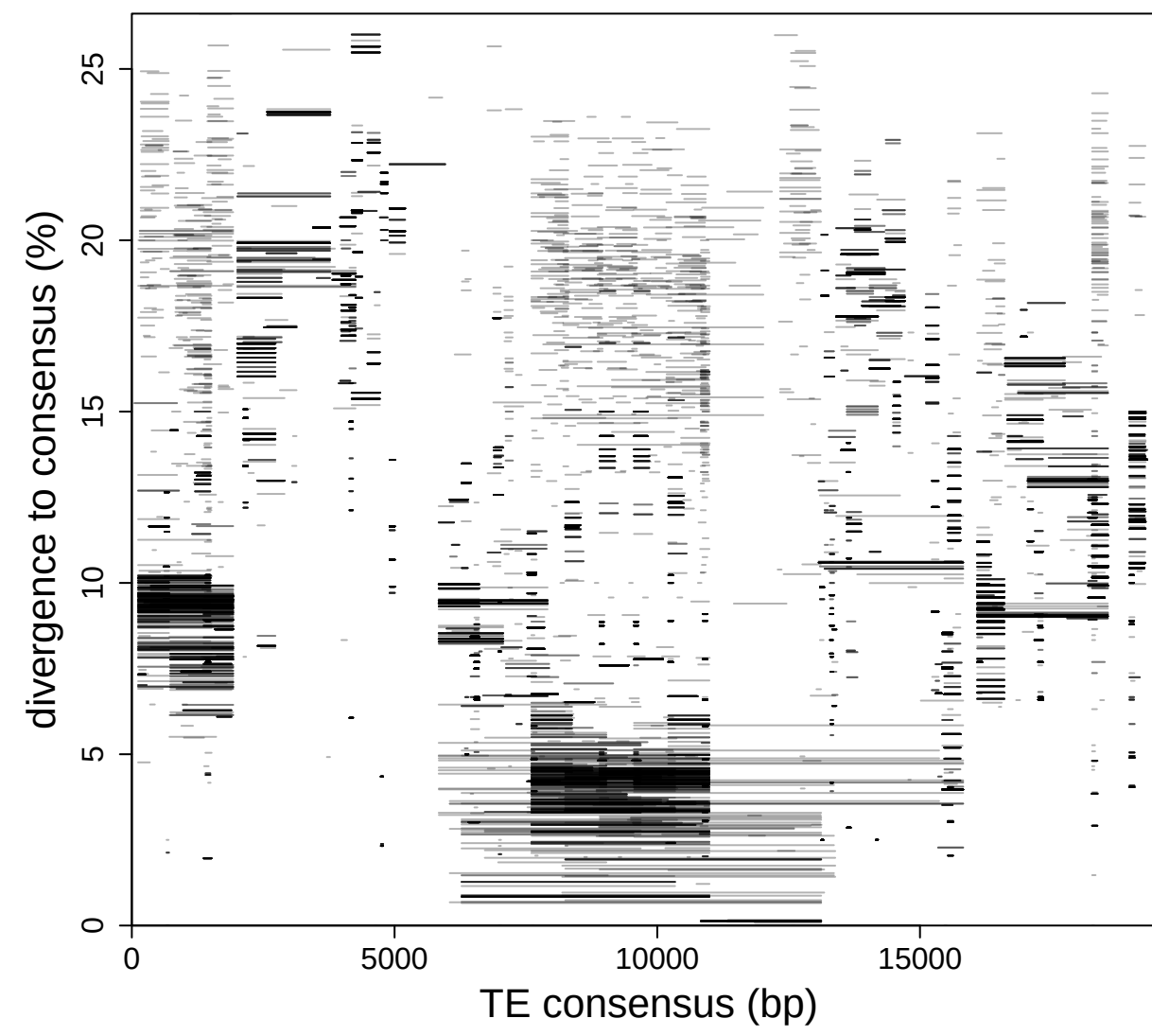
MSA length = 19679





g 1.bed fm 1.bed 0 0 n.bed g 1.bed fm 2.bed 0 0 bcln.fa a
size: 19679bp; fragments: 13064; full length: 0 (>=17711.1bp)

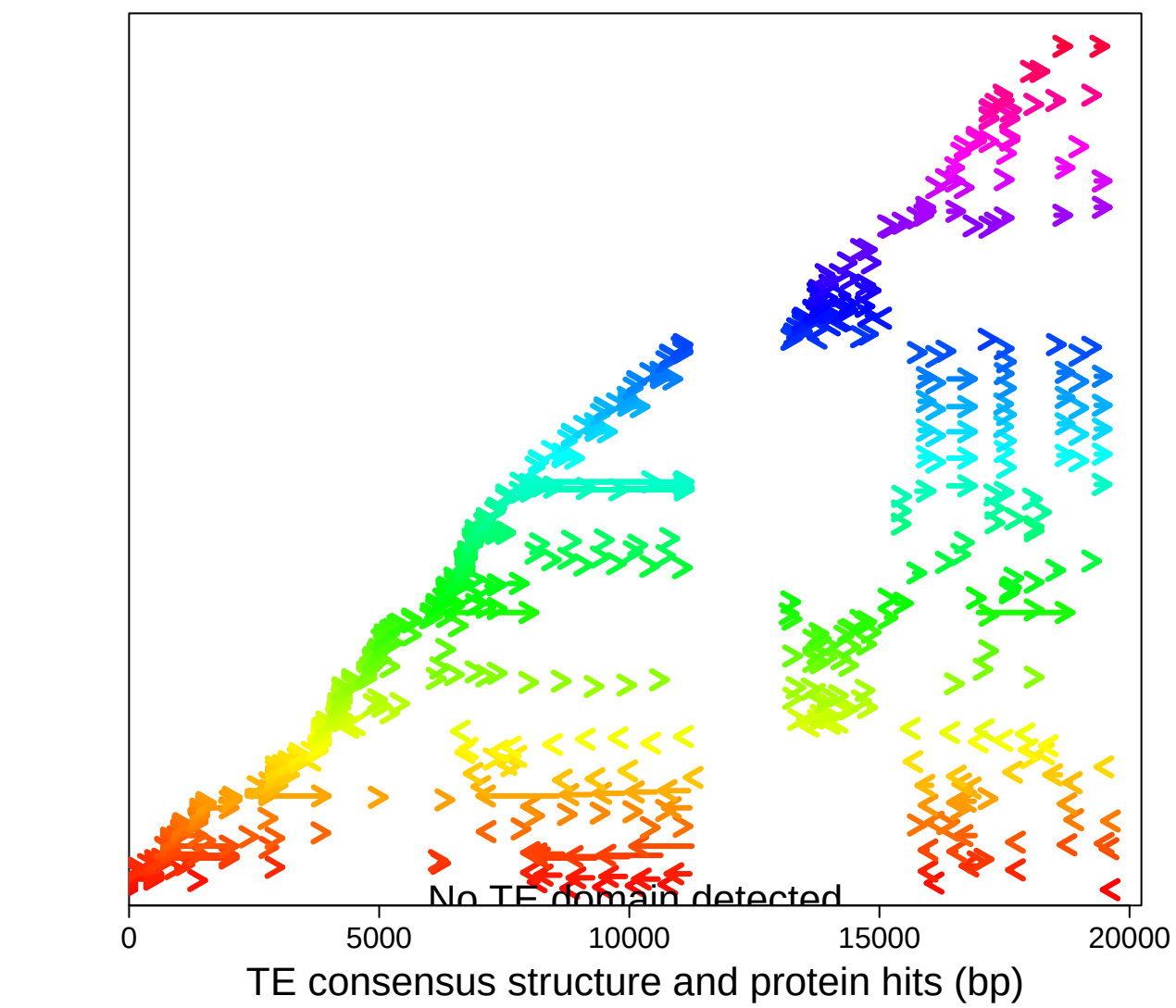
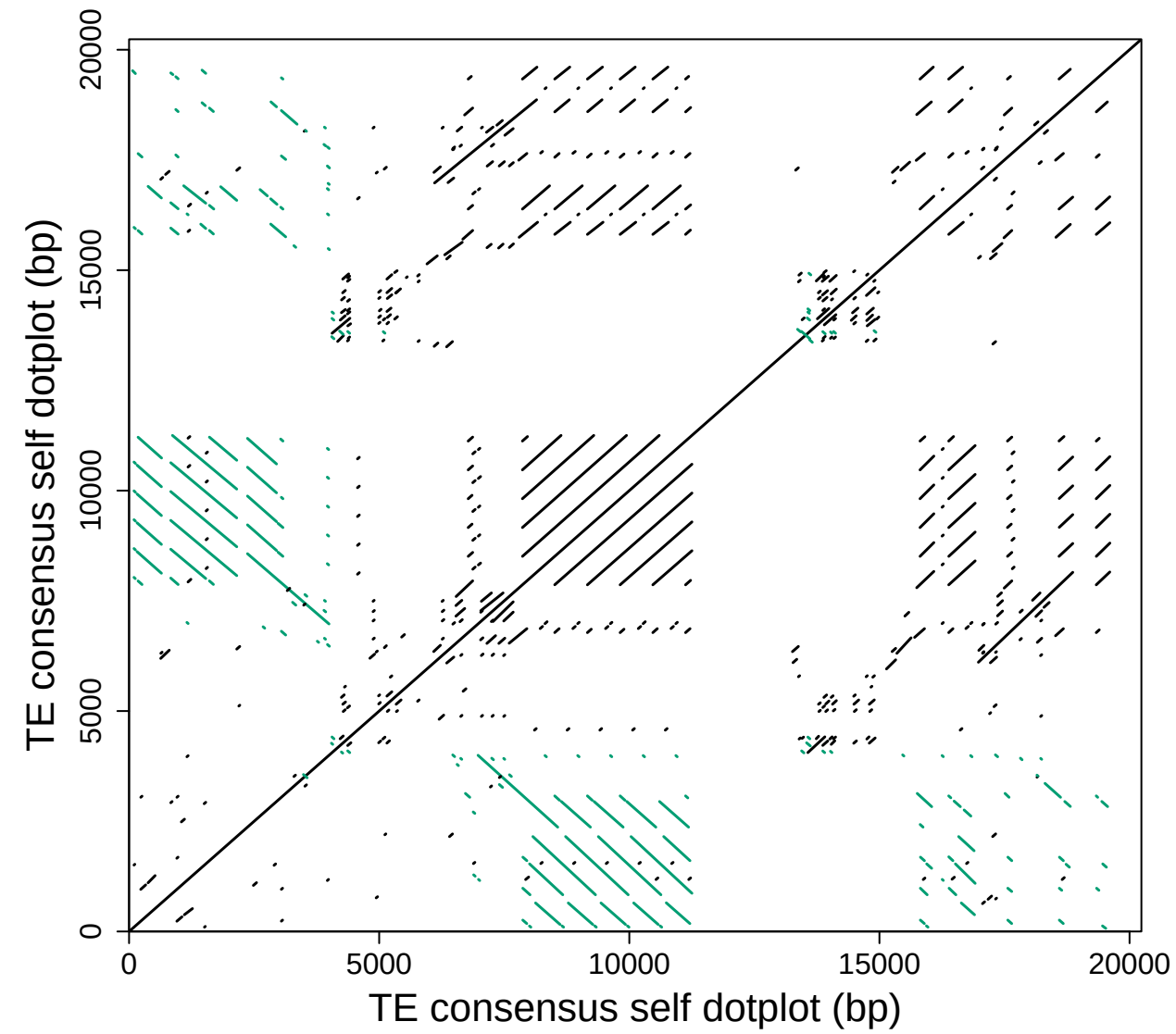
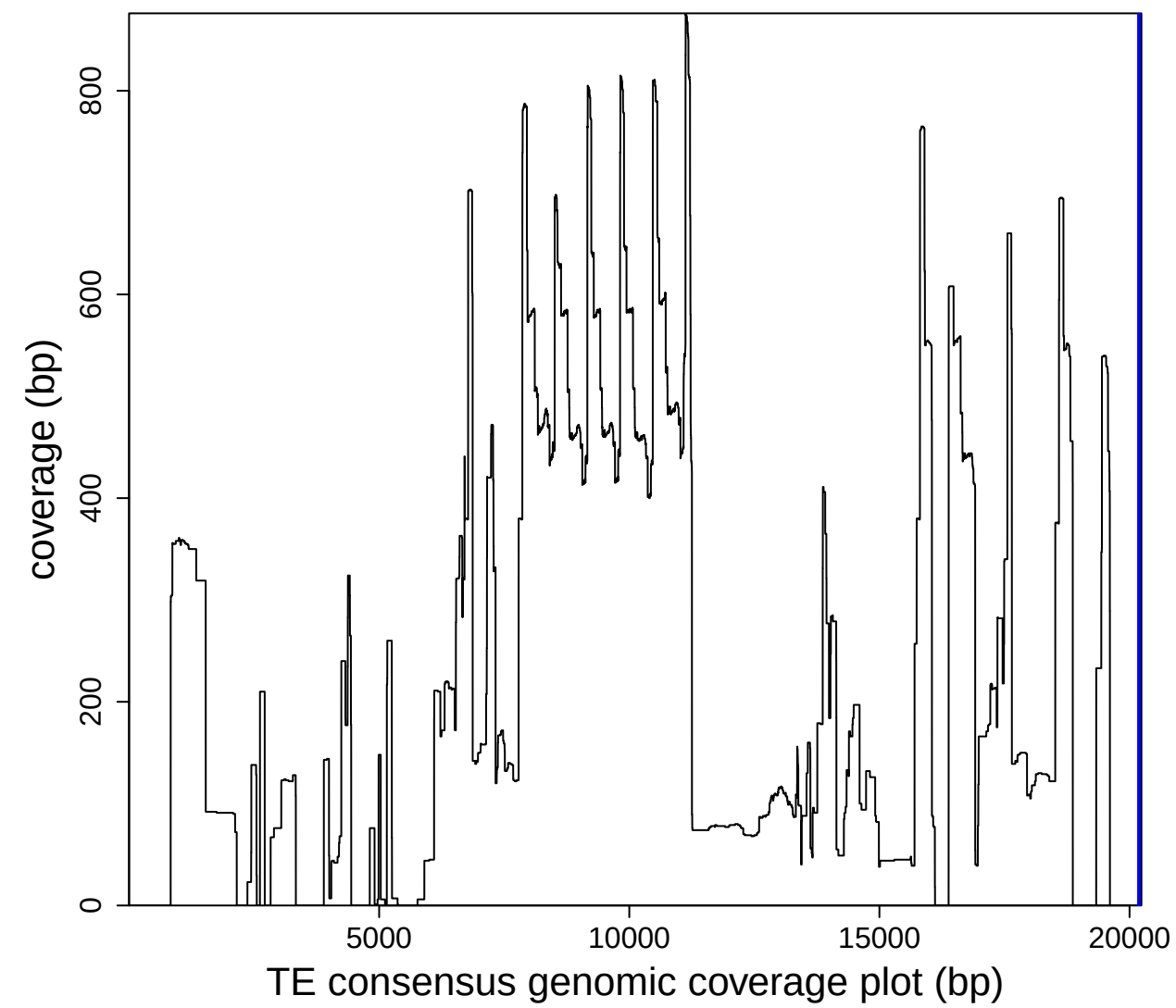
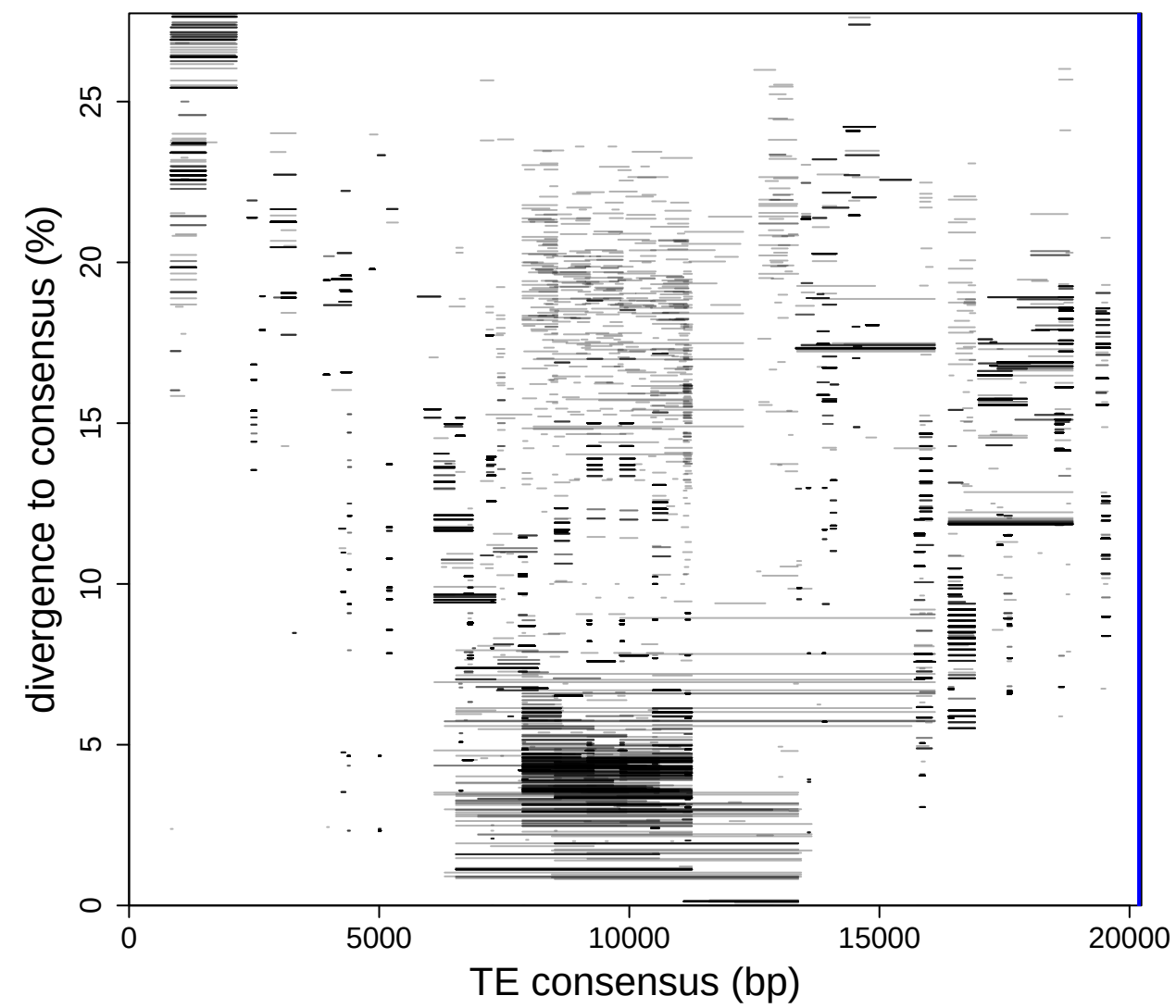
After TEtrimmer 19679 bp



f.bed_g_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bc
size: 20237bp; fragments: 10408; full length: 0 (>=18213.3bp)

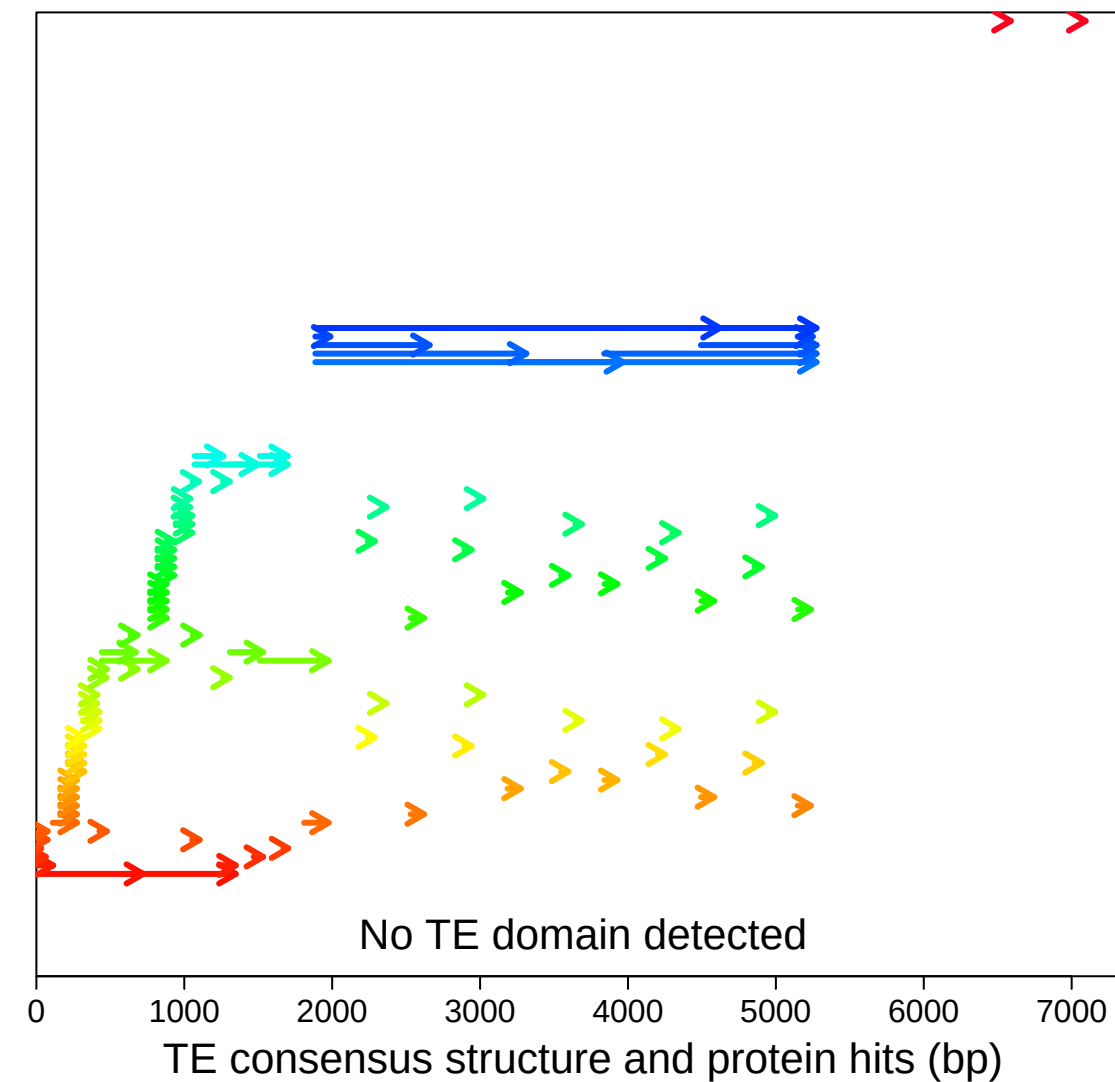
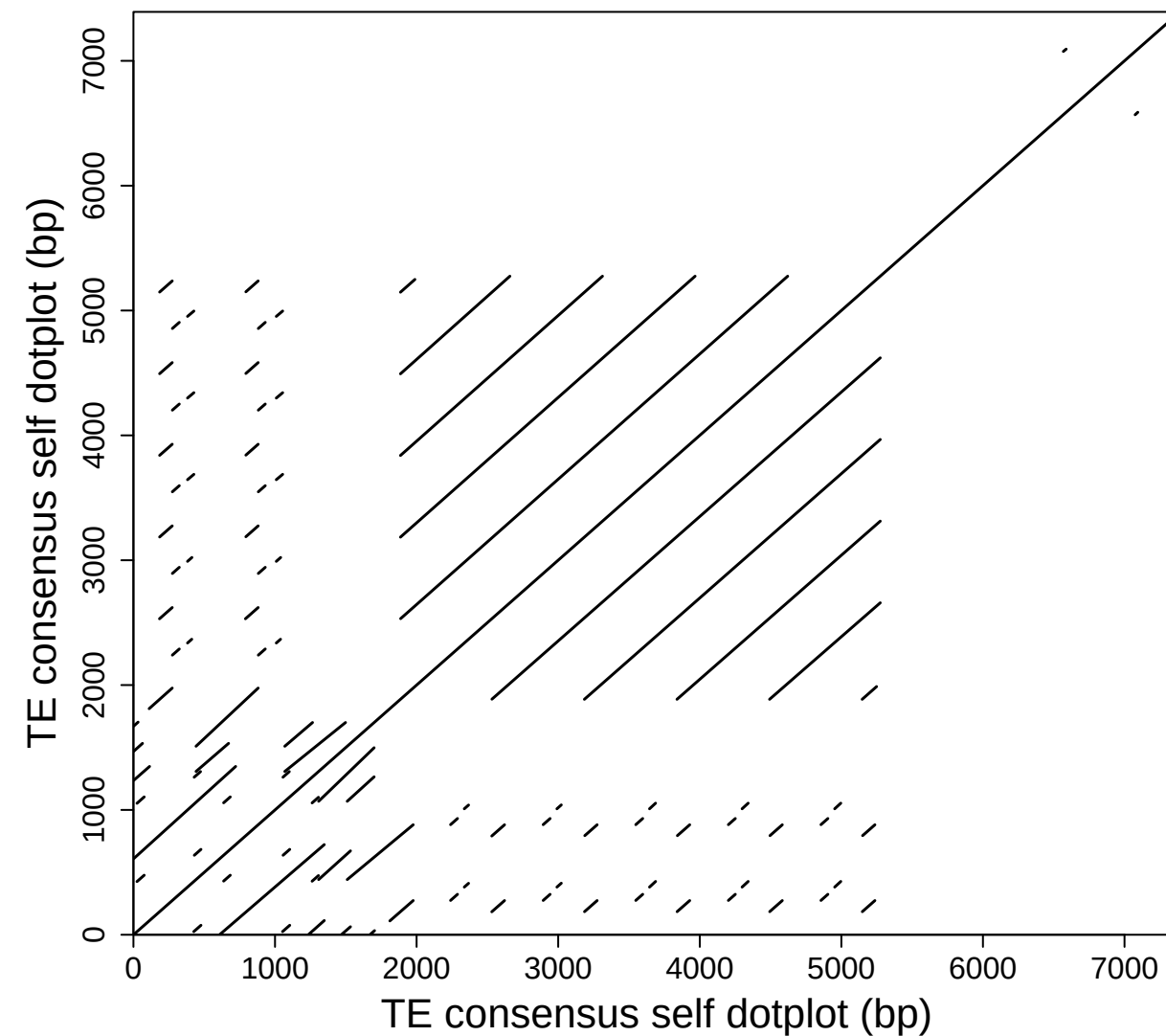
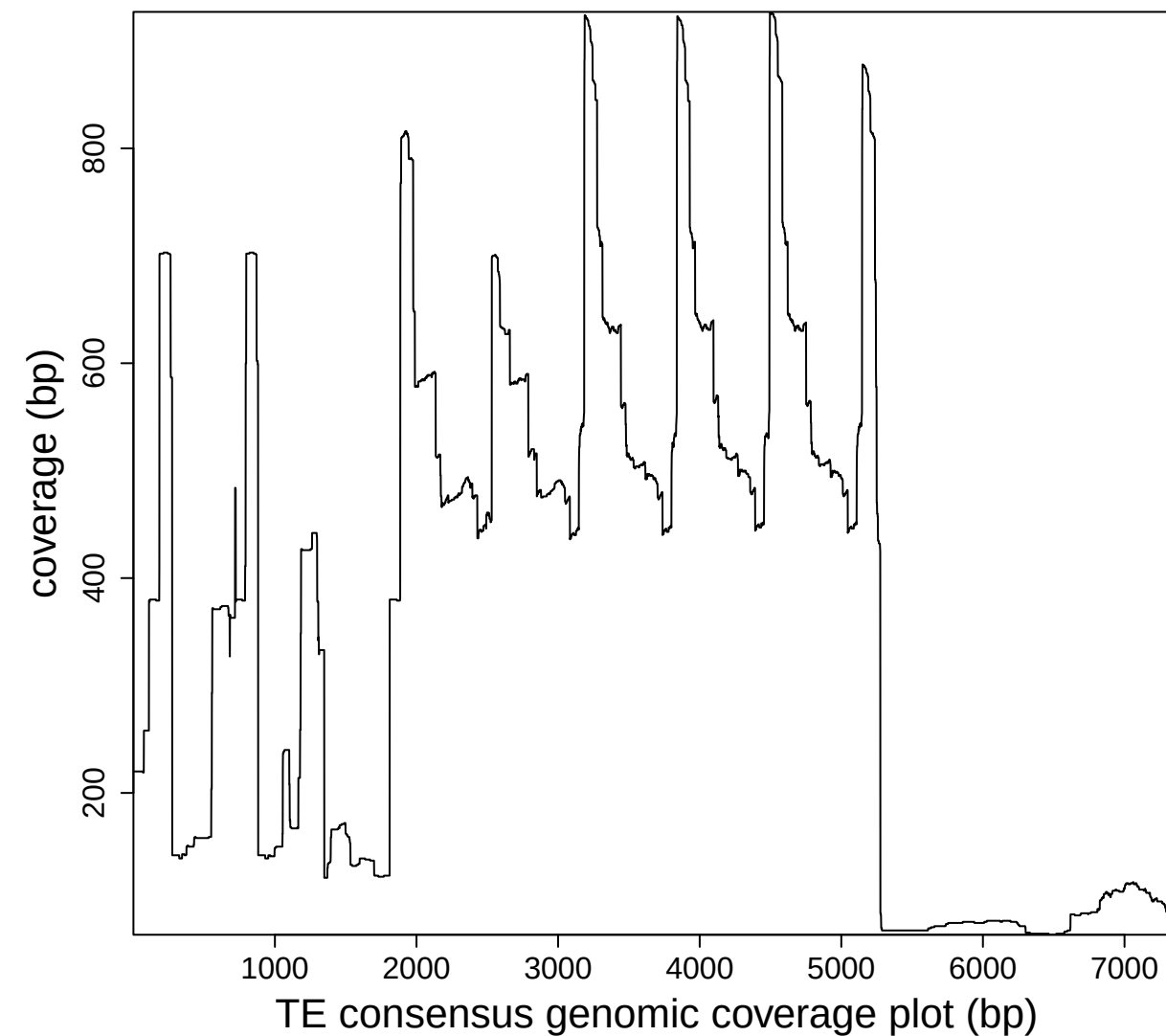
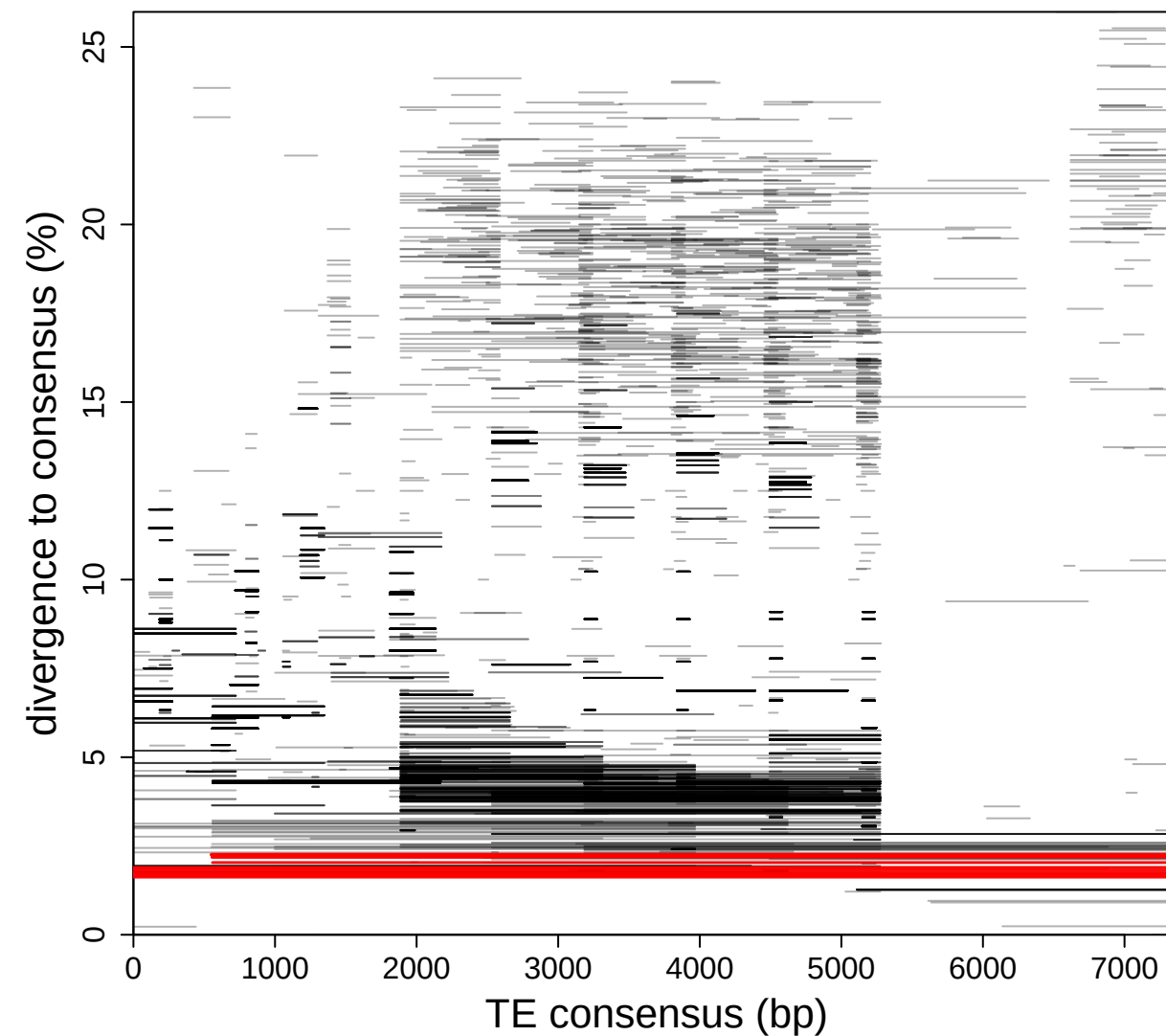
After TEtrimmer Extended plot

Blue lines are boundaries

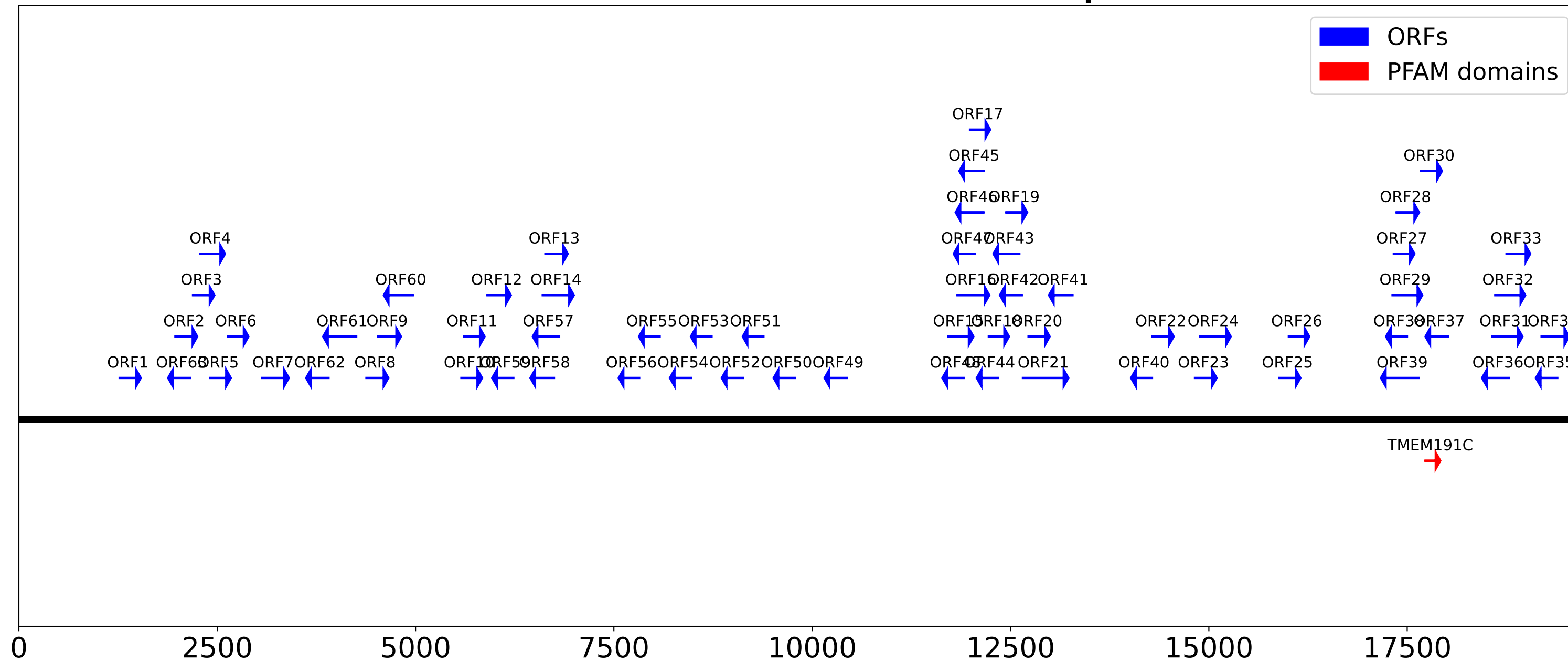


TE: rnd_5_family_4748
size: 7392bp; fragments: 5293; full length: 18 (≥ 6652.8 bp)

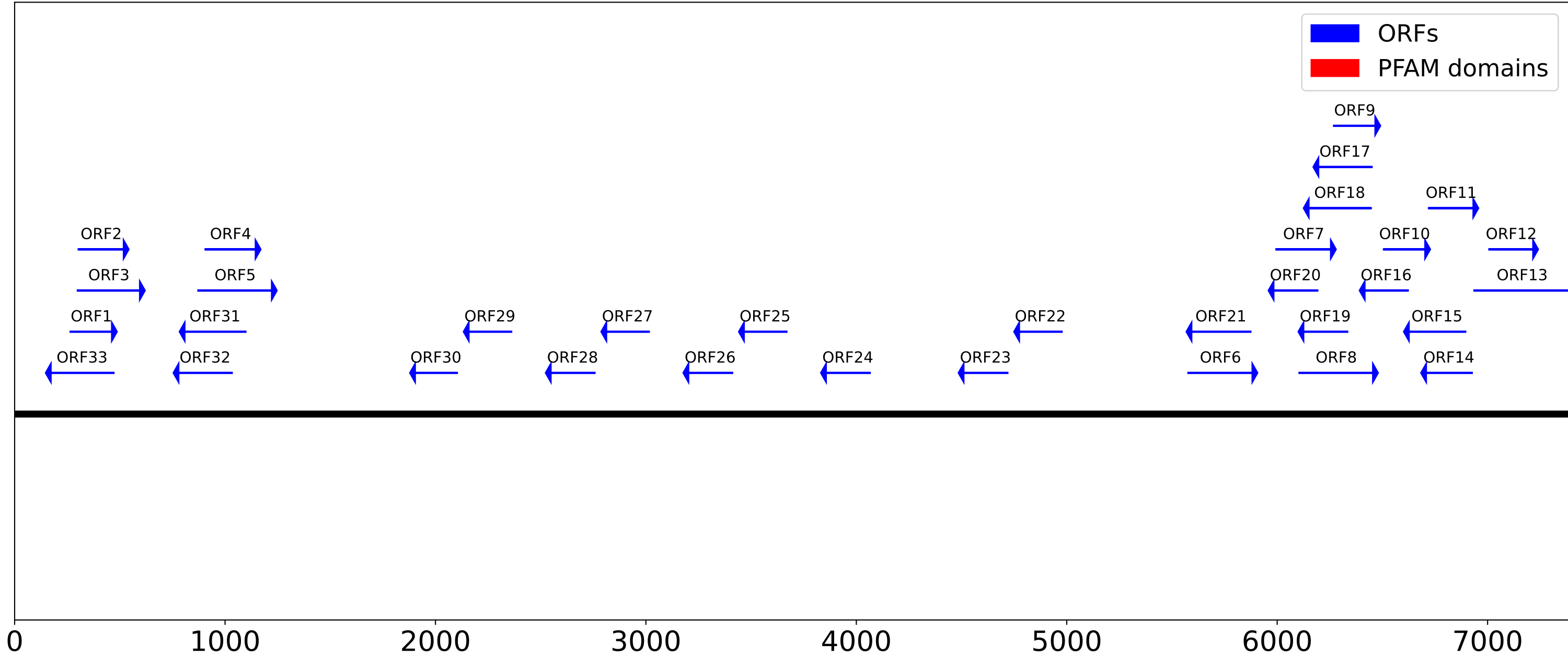
Before TEtrimmer 7392 bp



After TETrimmer ORF and PFAM domain plot



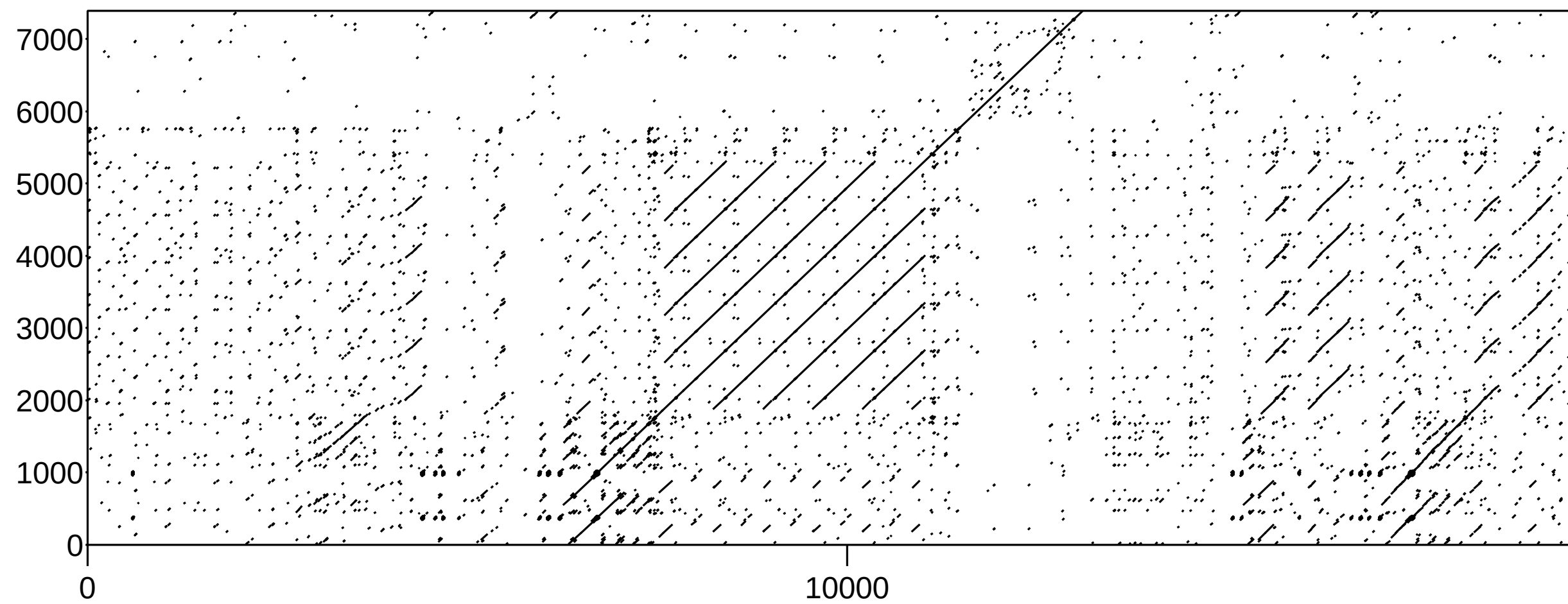
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)