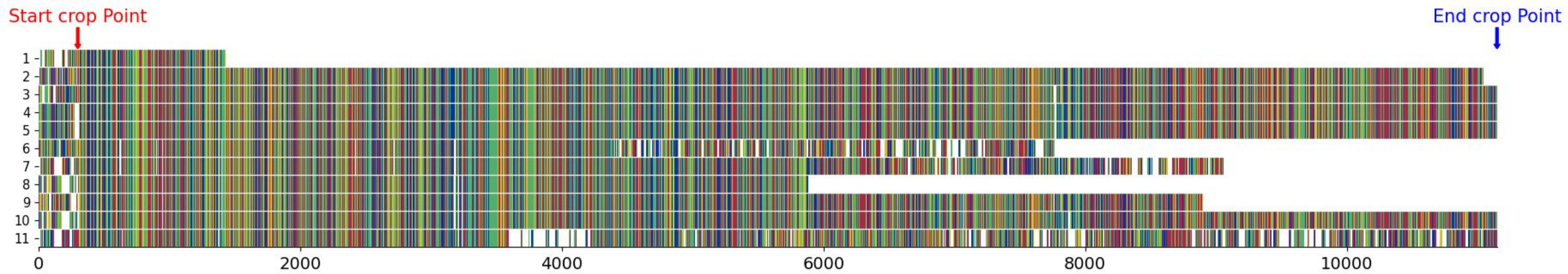


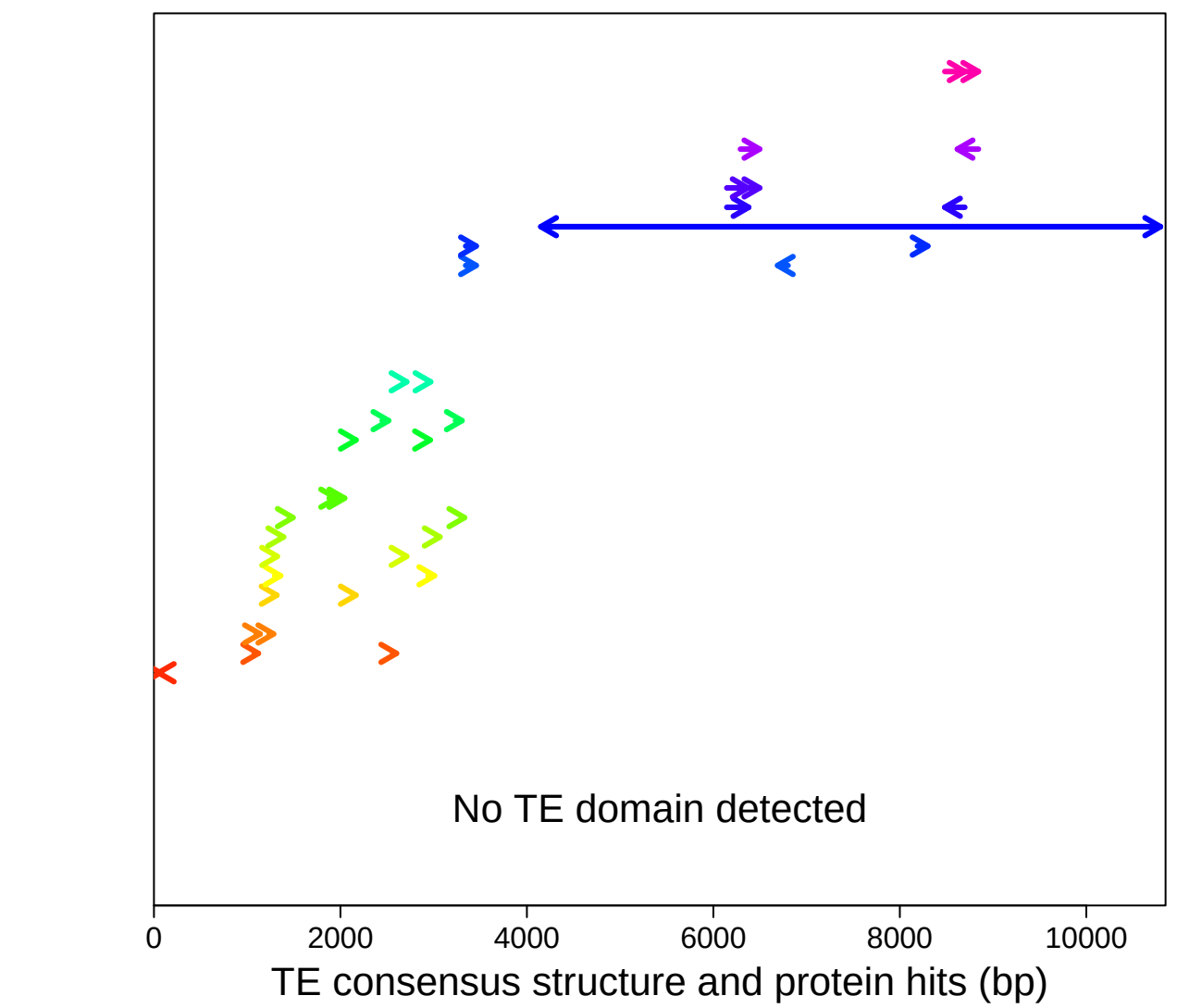
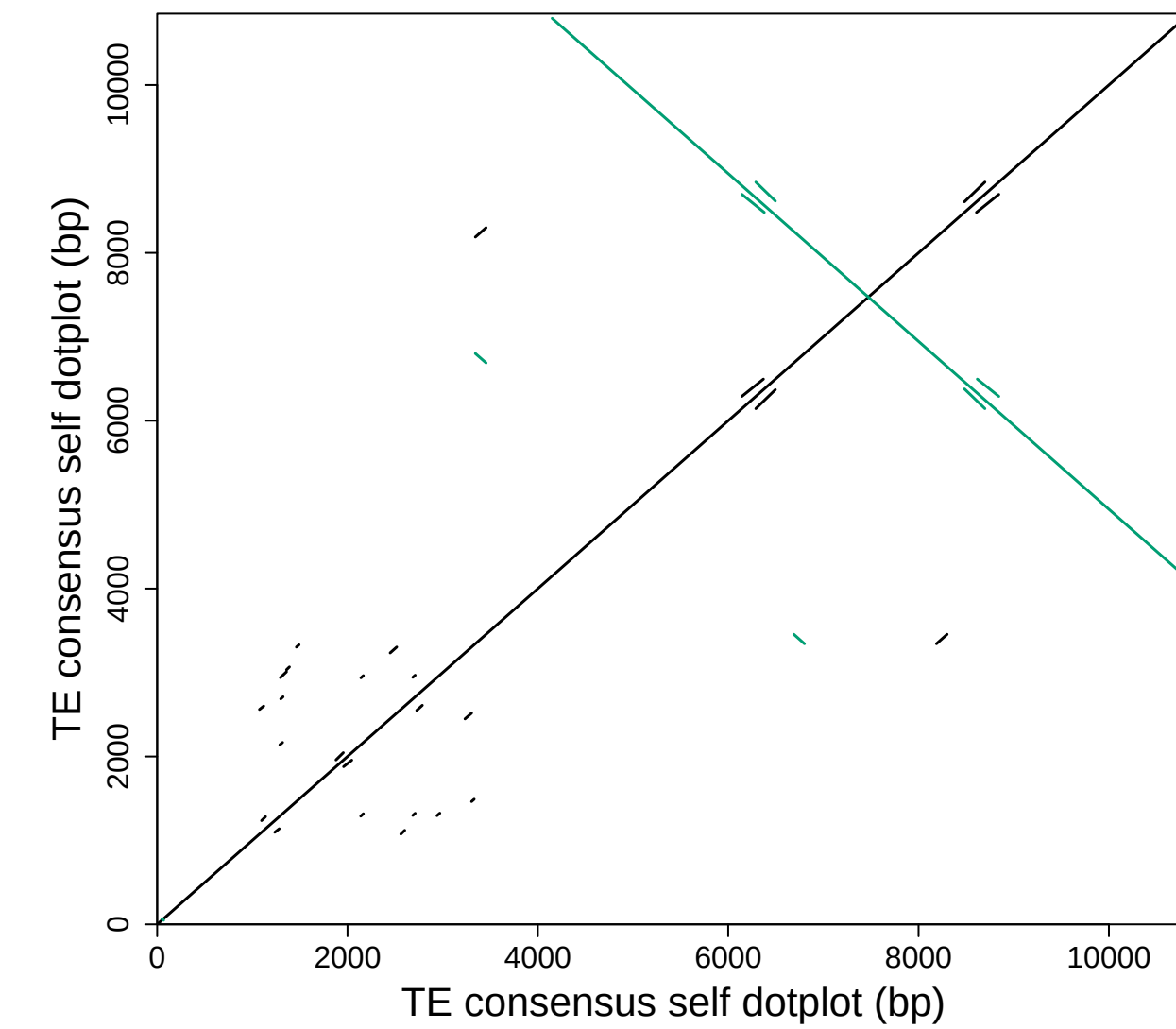
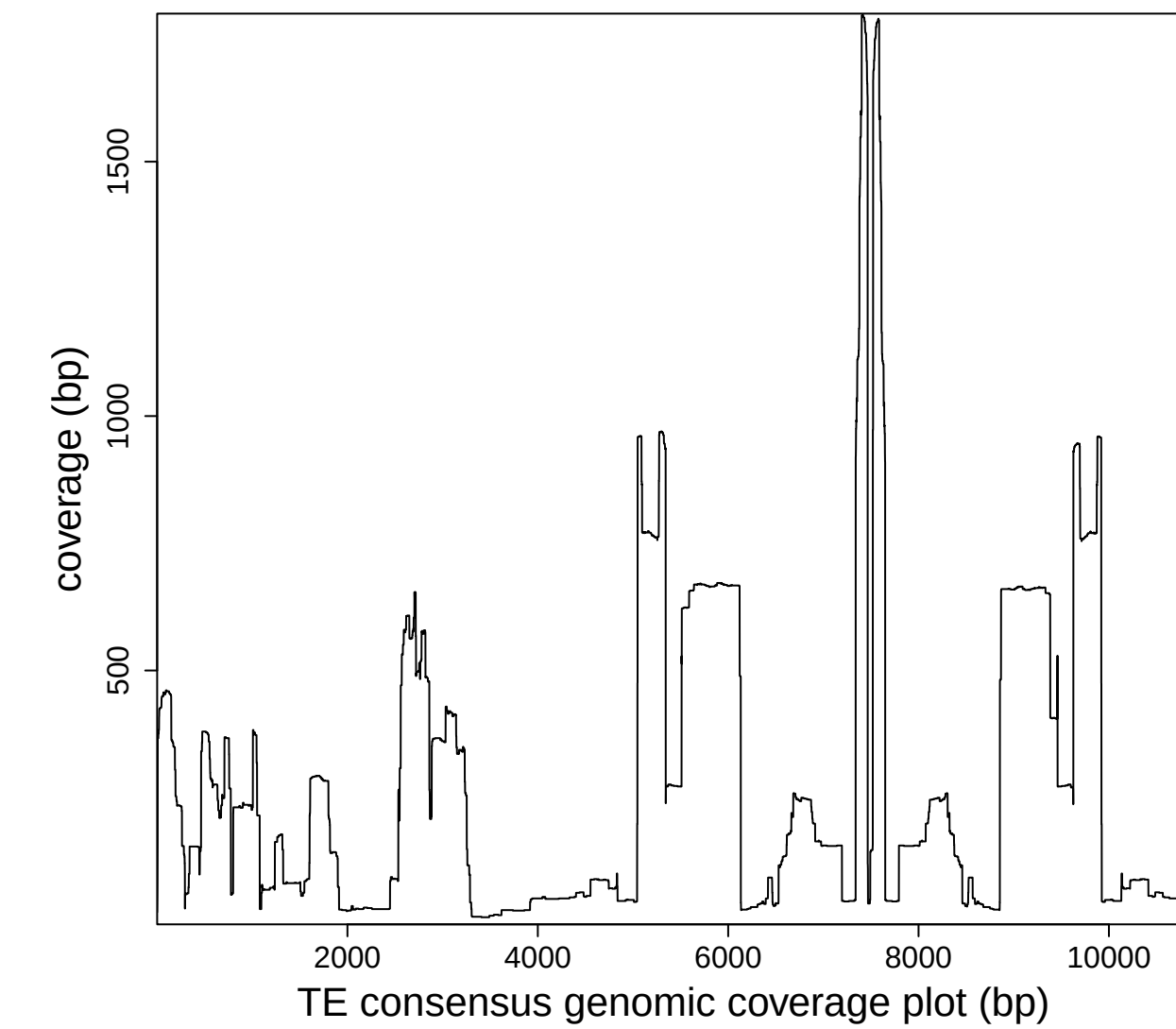
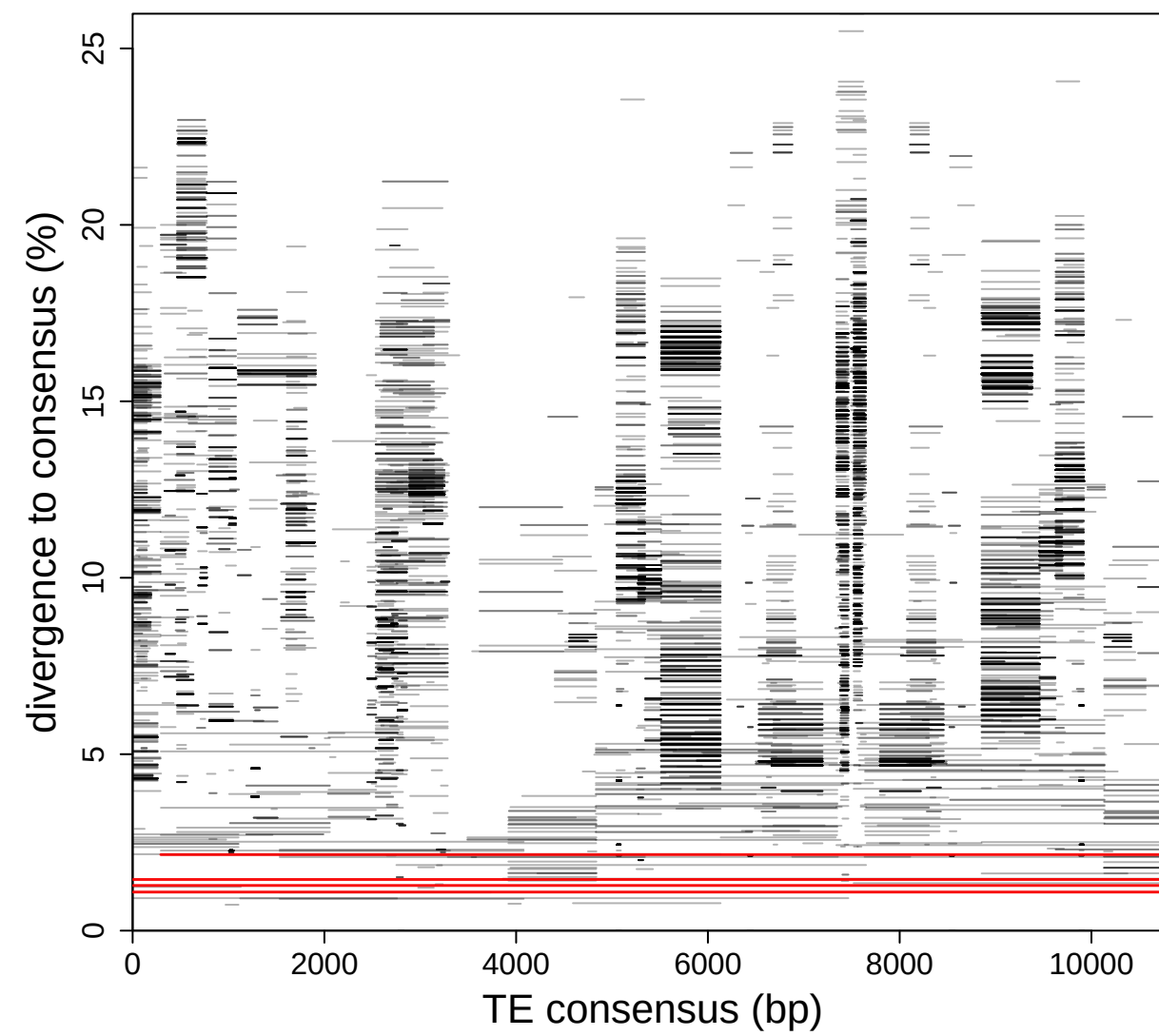
End crop Point

MSA length = 10850



g_3.bed fm_1.bed 0 0 n.bed g_1.bed fm_2.bed 0 0 bcln.fa ε
size: 10850bp; fragments: 11182; full length: 4 (>=9765bp)

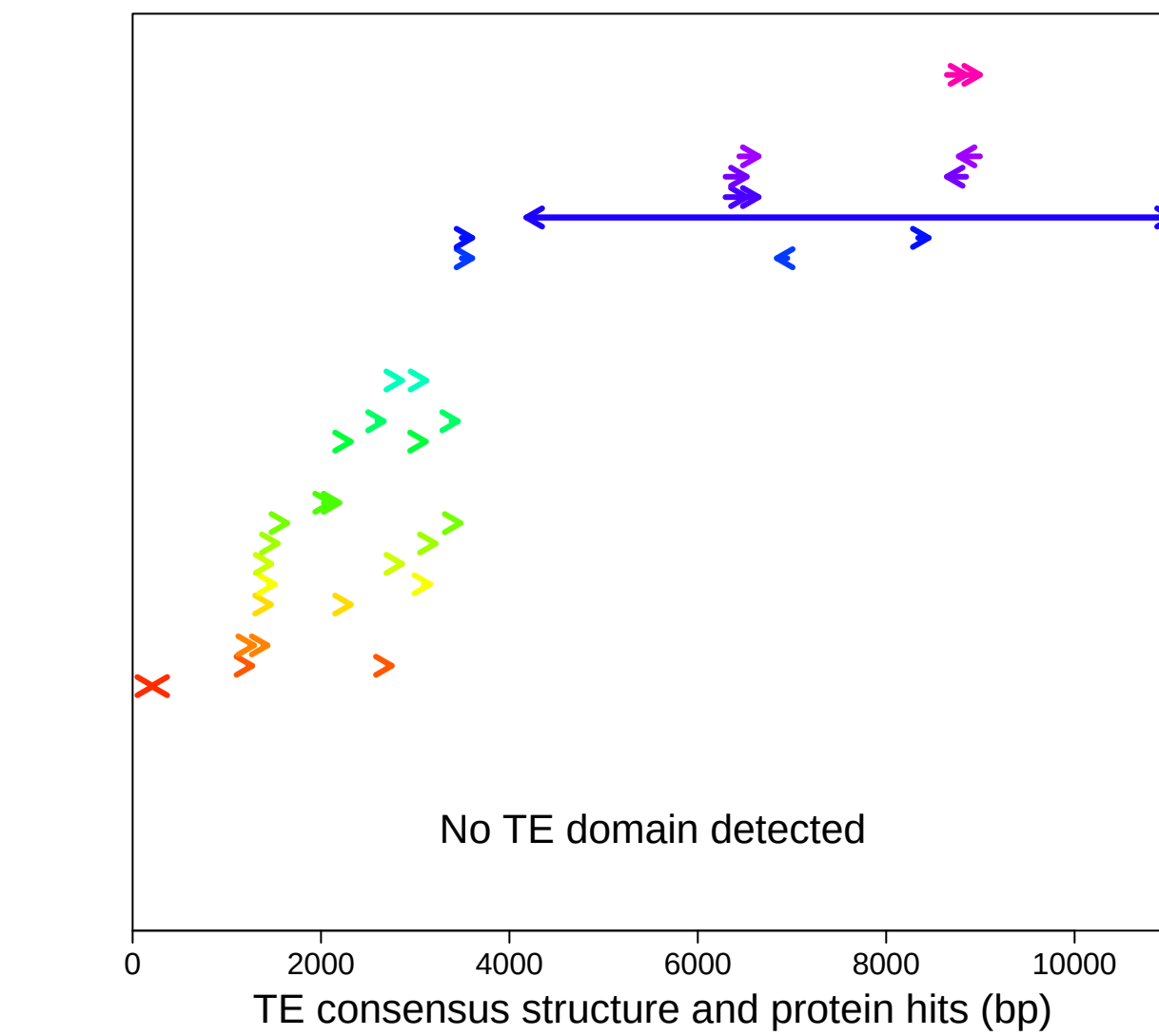
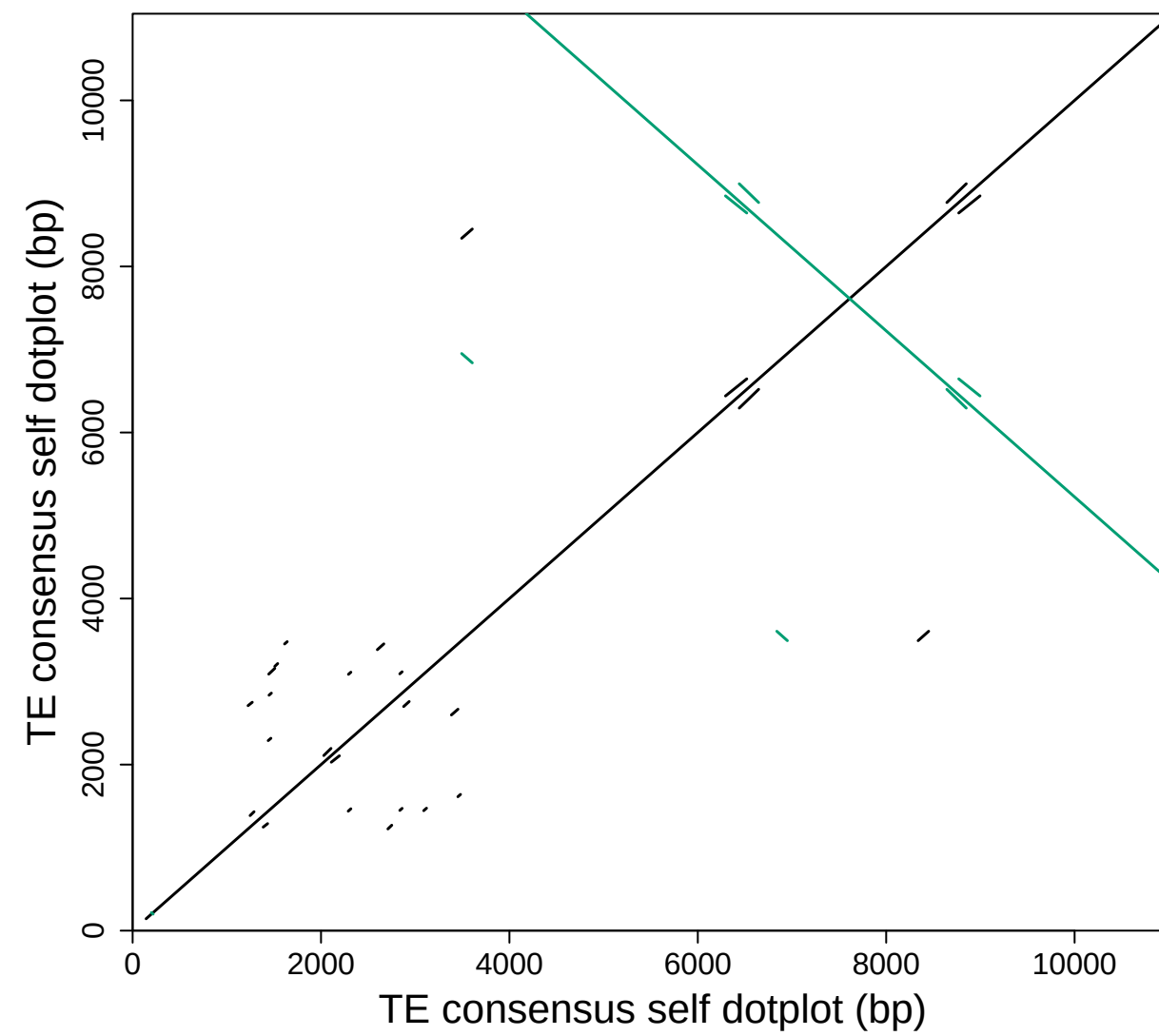
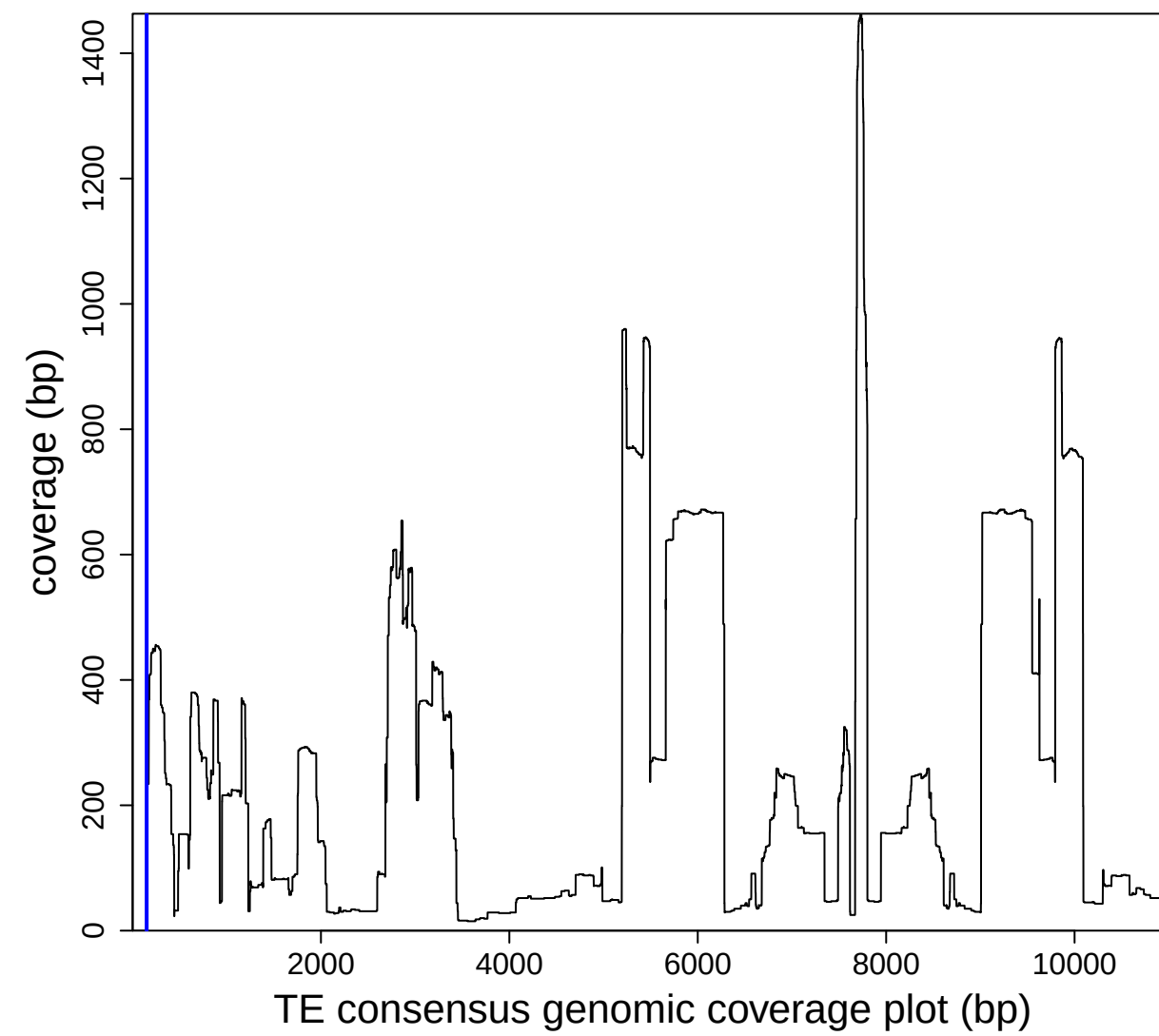
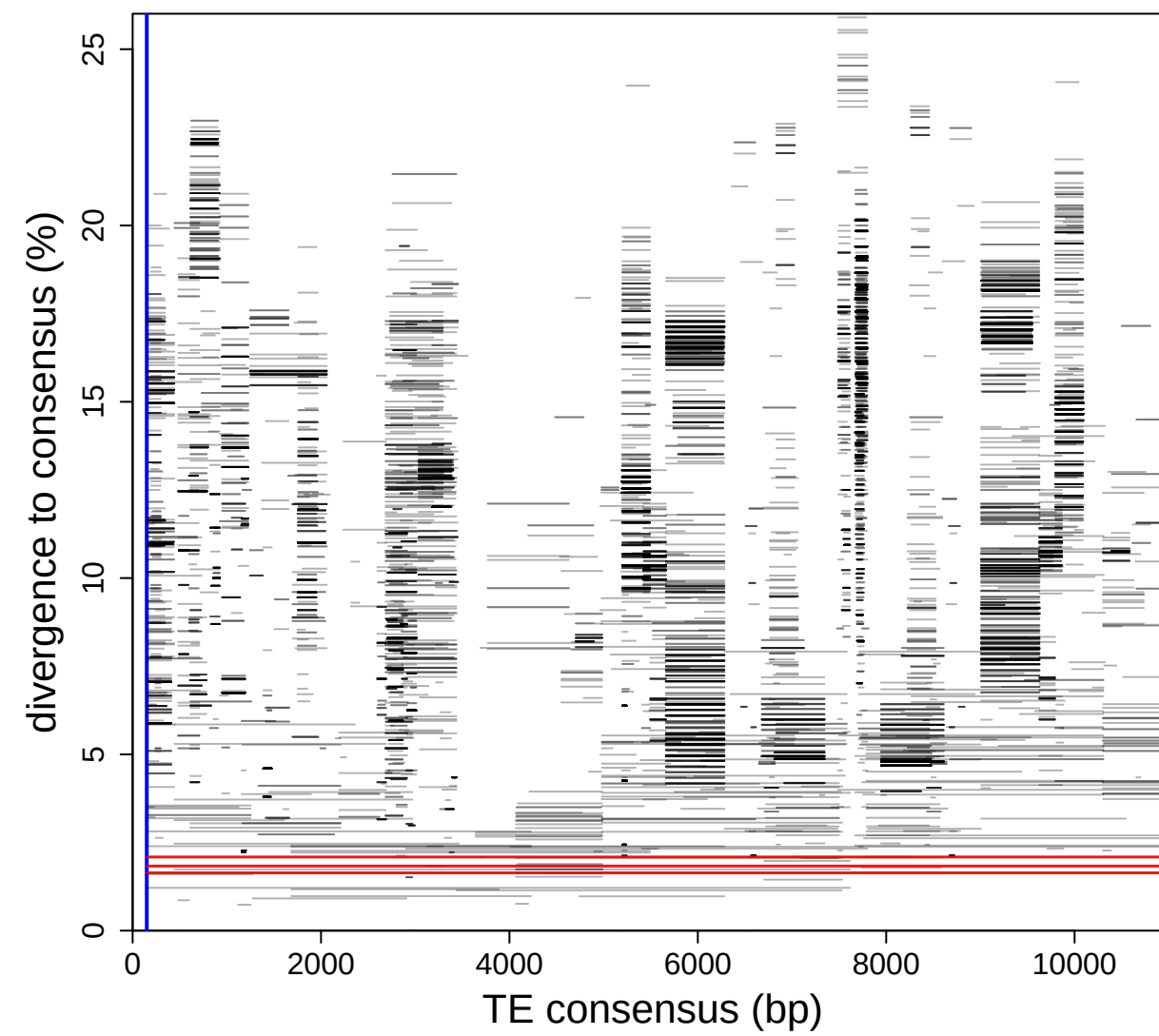
After TEtrimmer 10850 bp



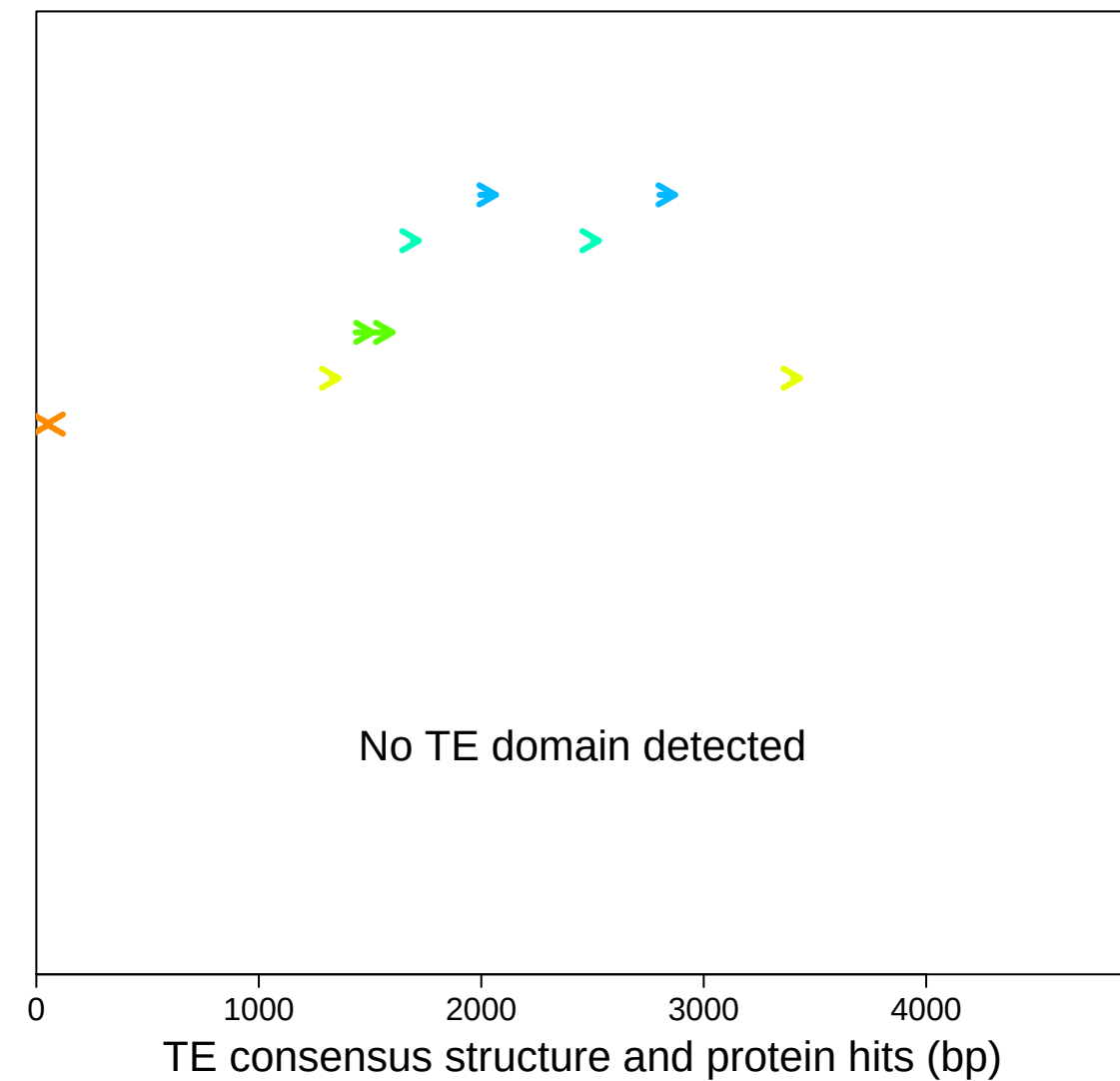
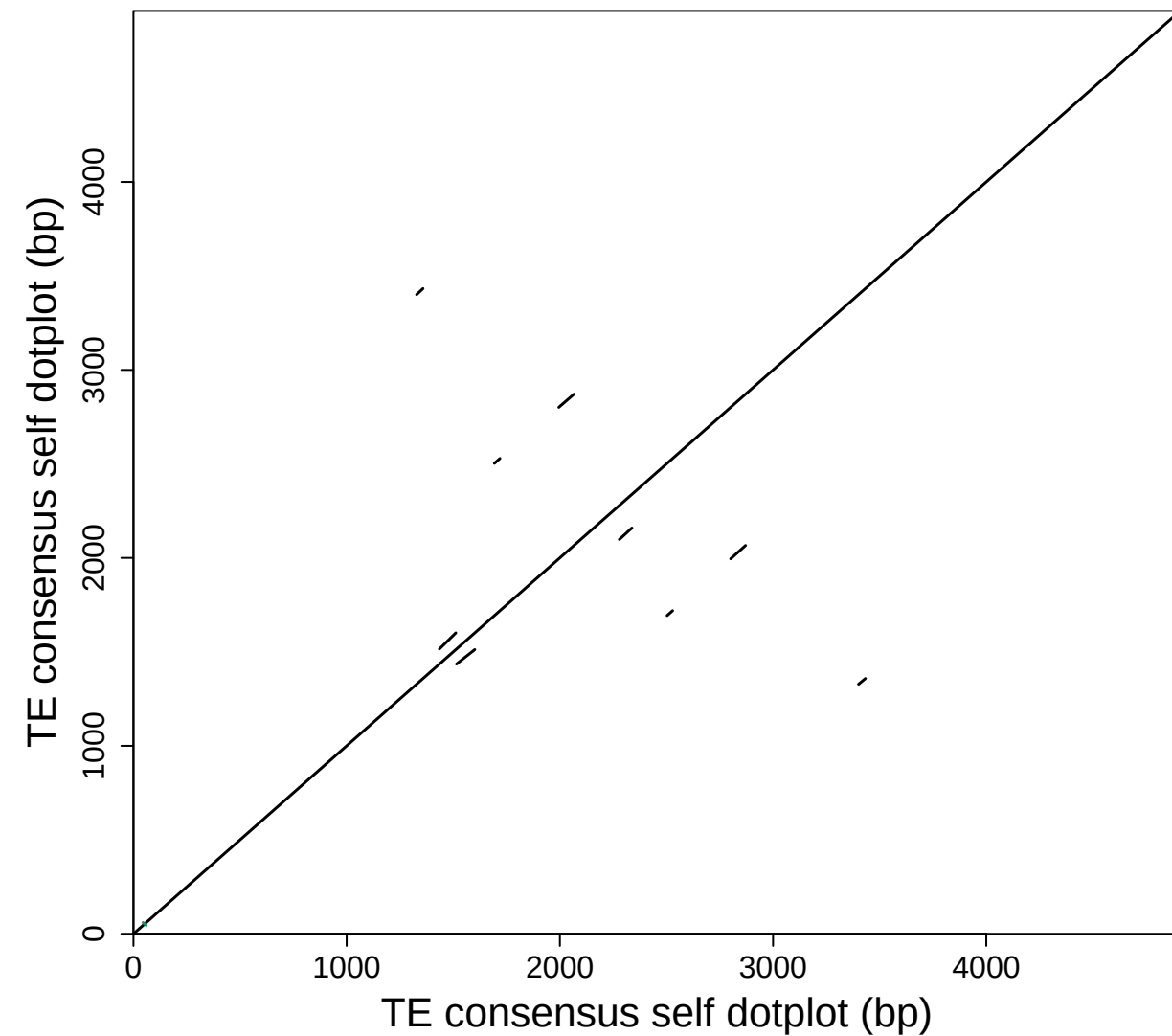
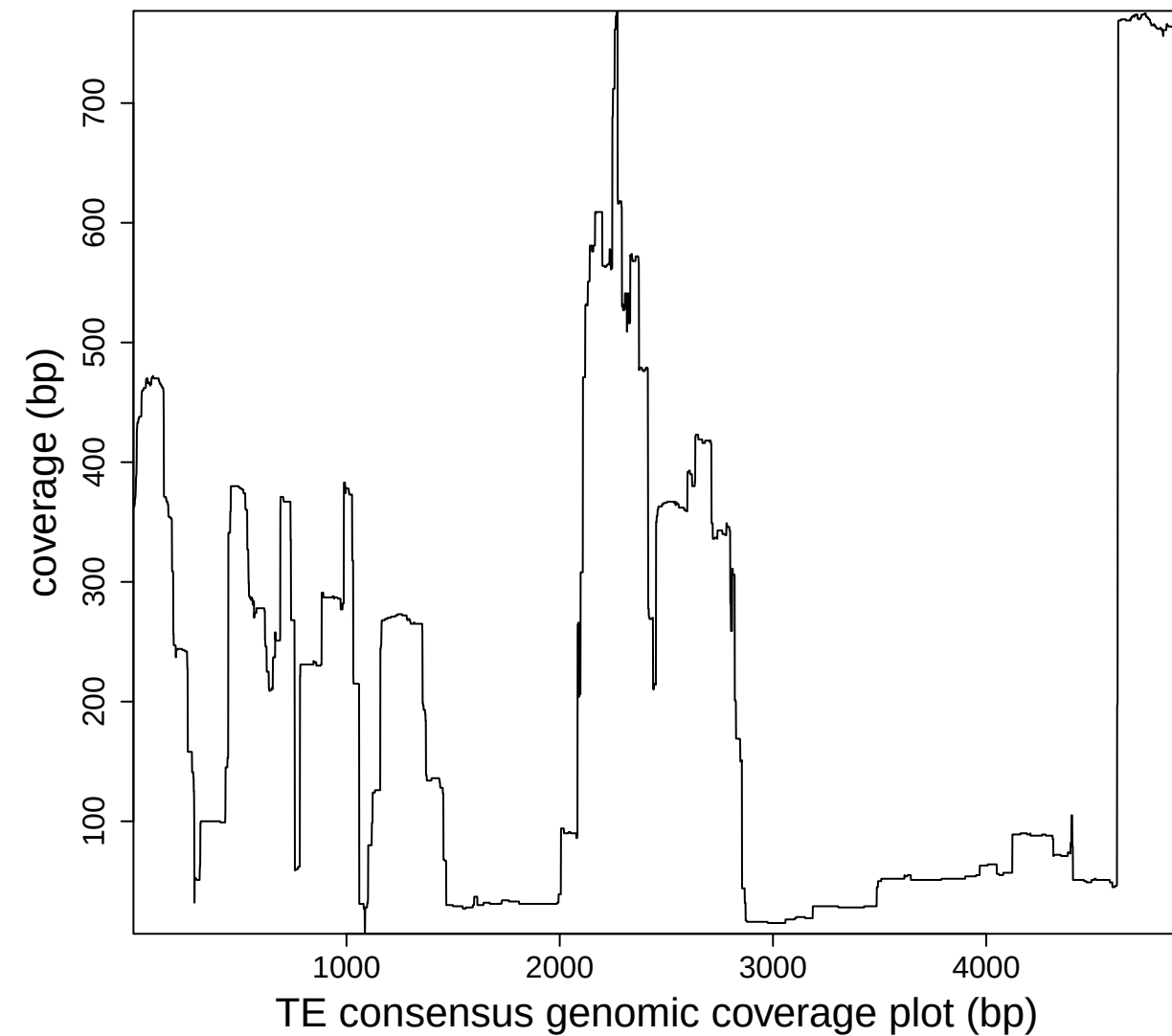
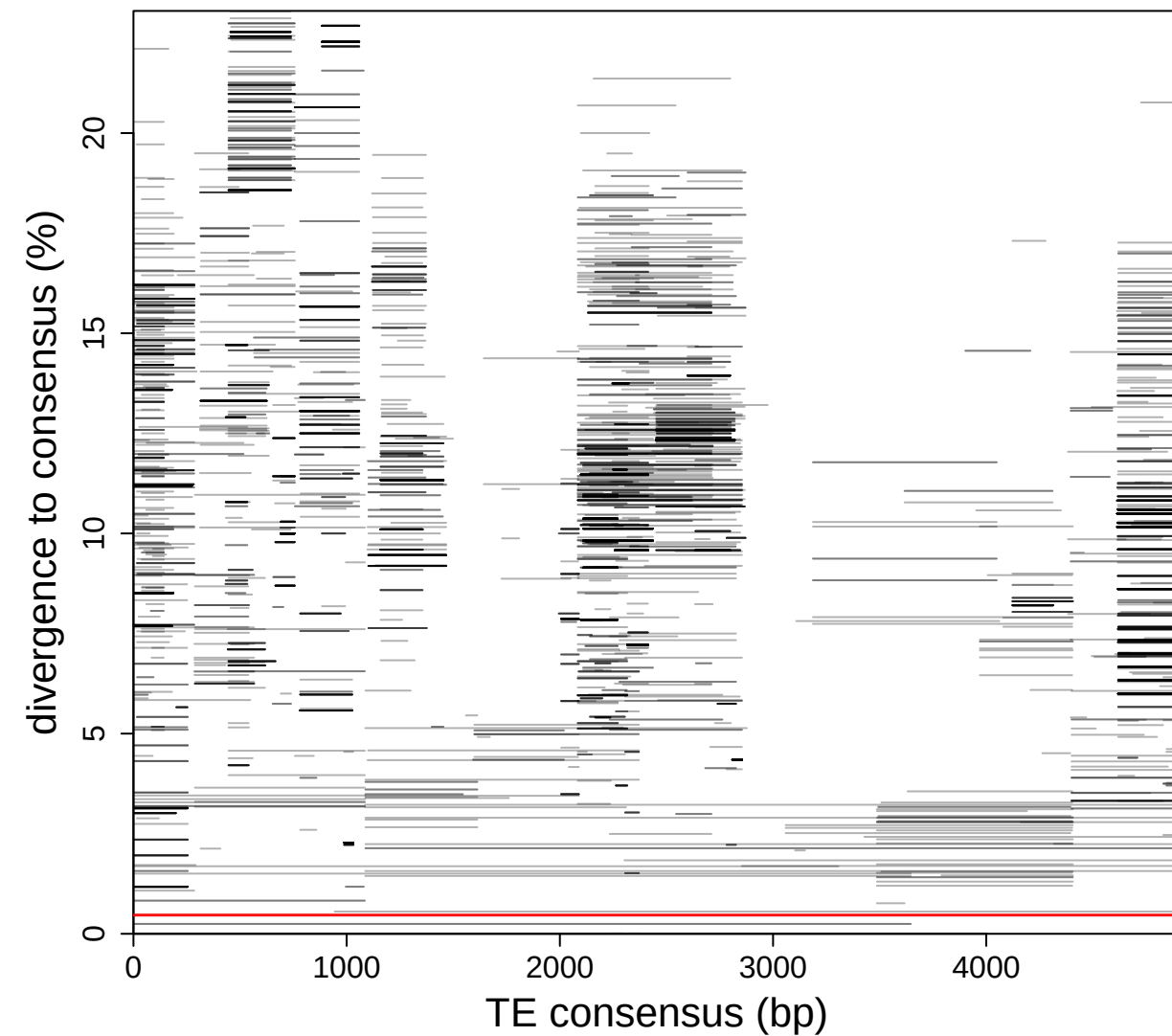
f.bed_g_3.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bc
size: 11044bp; fragments: 9236; full length: 3 (>=9939.6bp)

After TEtrimmer Extended plot

Blue lines are boundaries

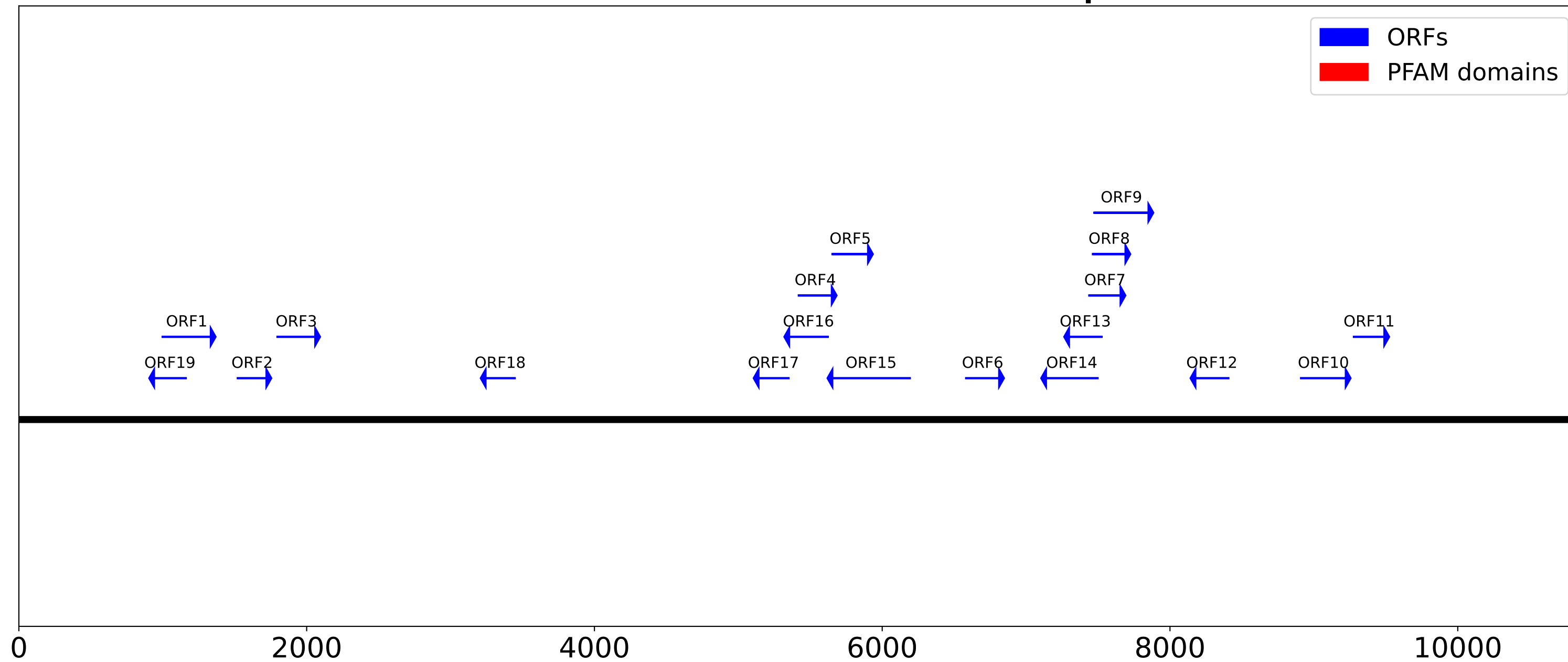


TE: rnd_5_family_3948
size: 4910bp; fragments: 3997; full length: 1 (>=4419bp)

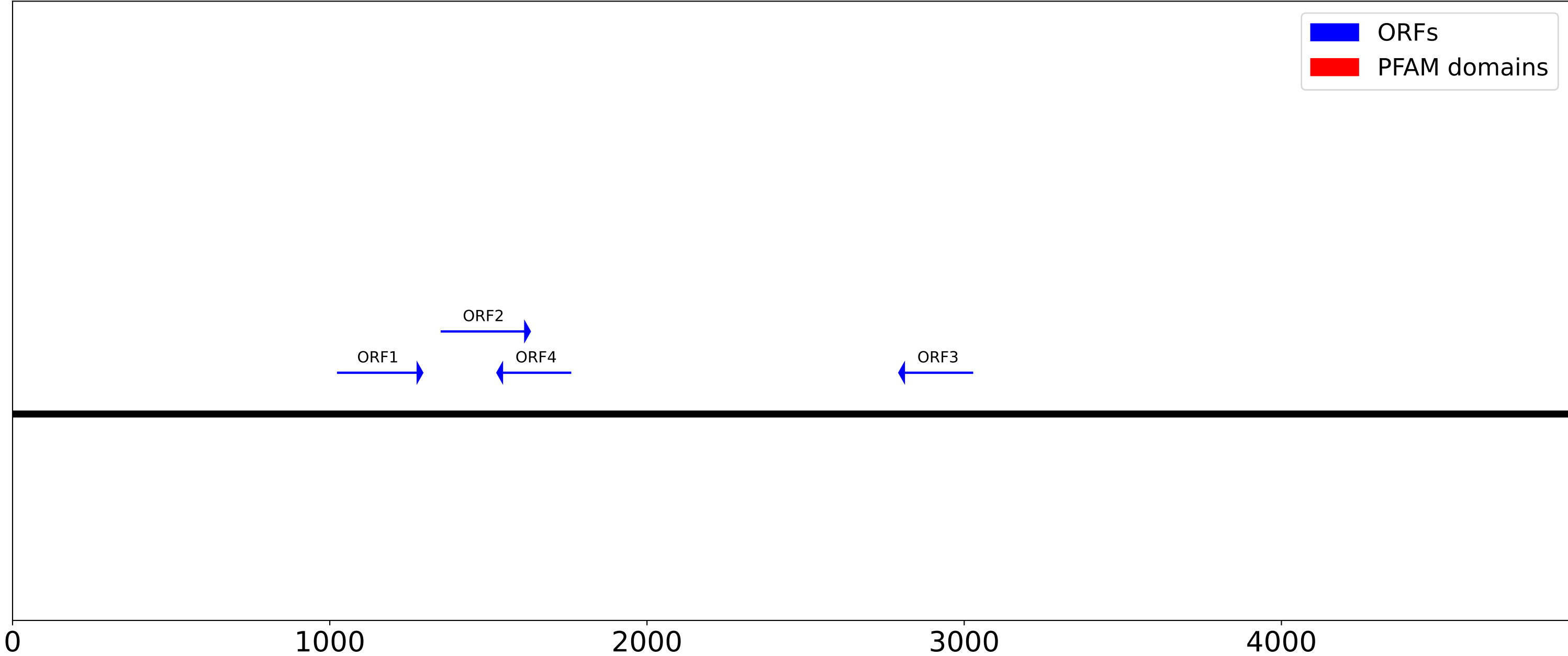


Before TETrimmer 4910 bp

After TEtrimmer ORF and PFAM domain plot



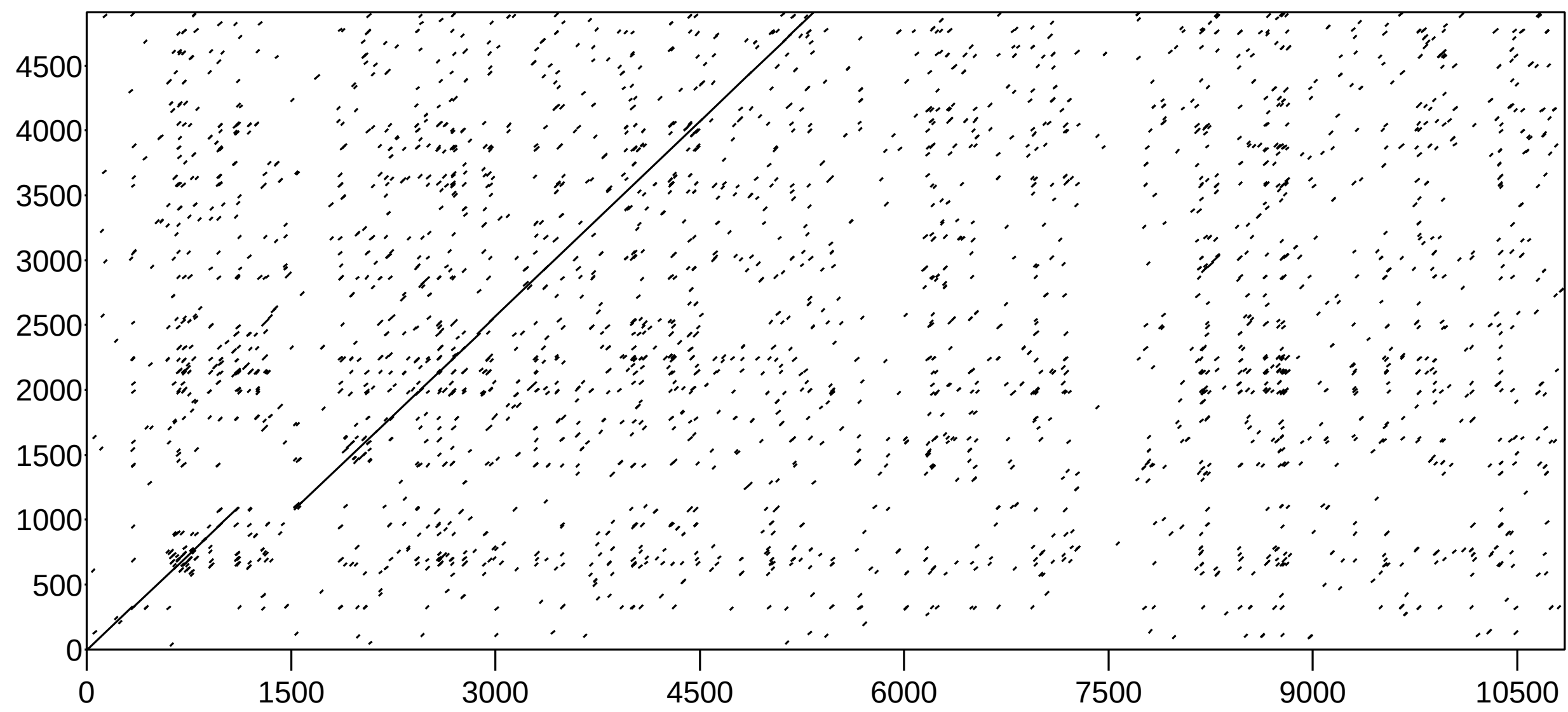
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)