

Start crop Point

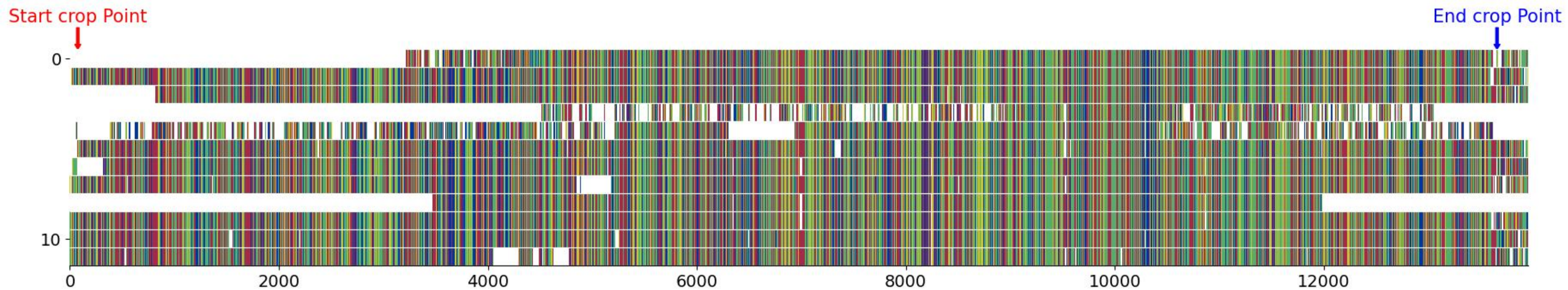


End crop Point



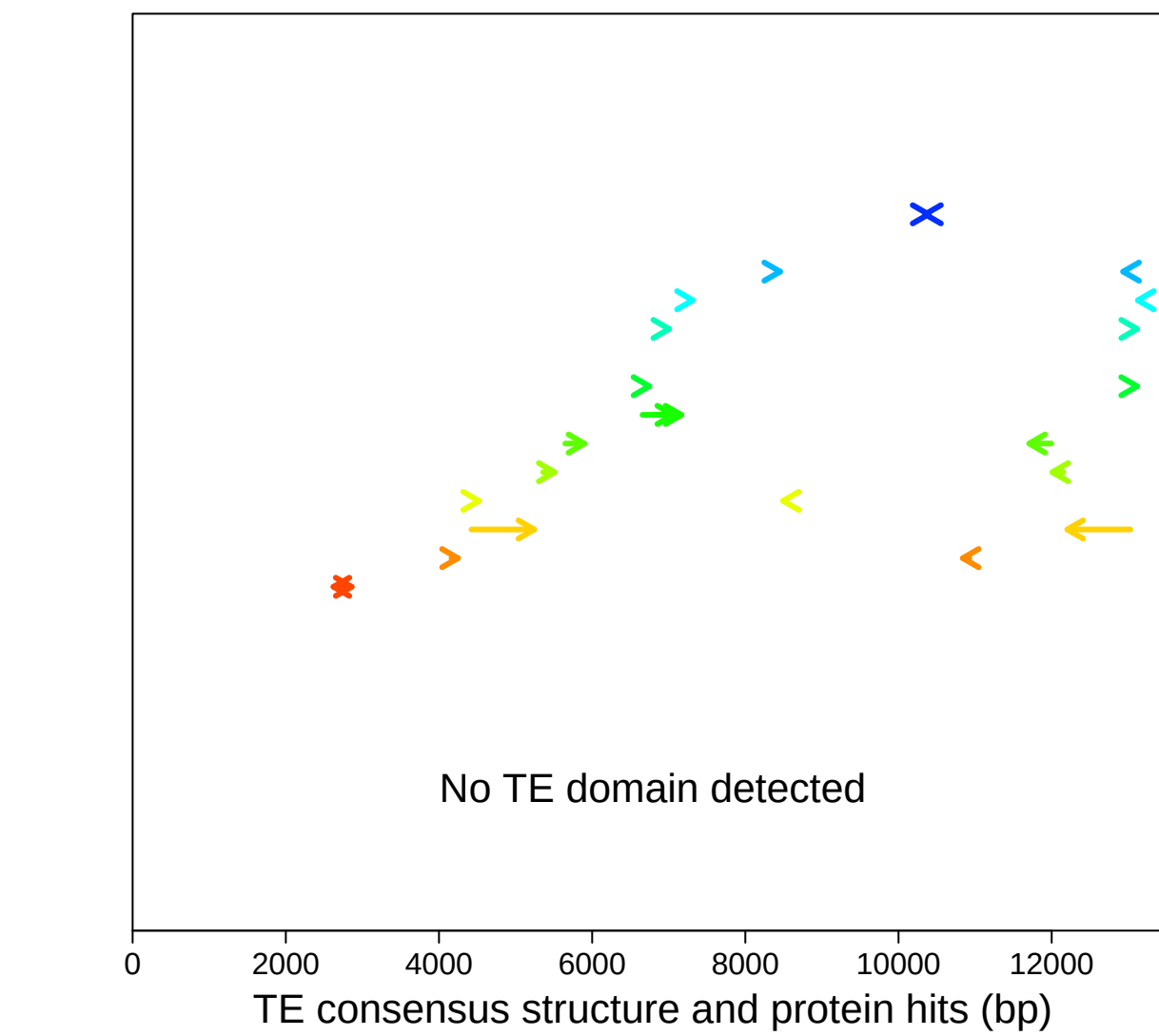
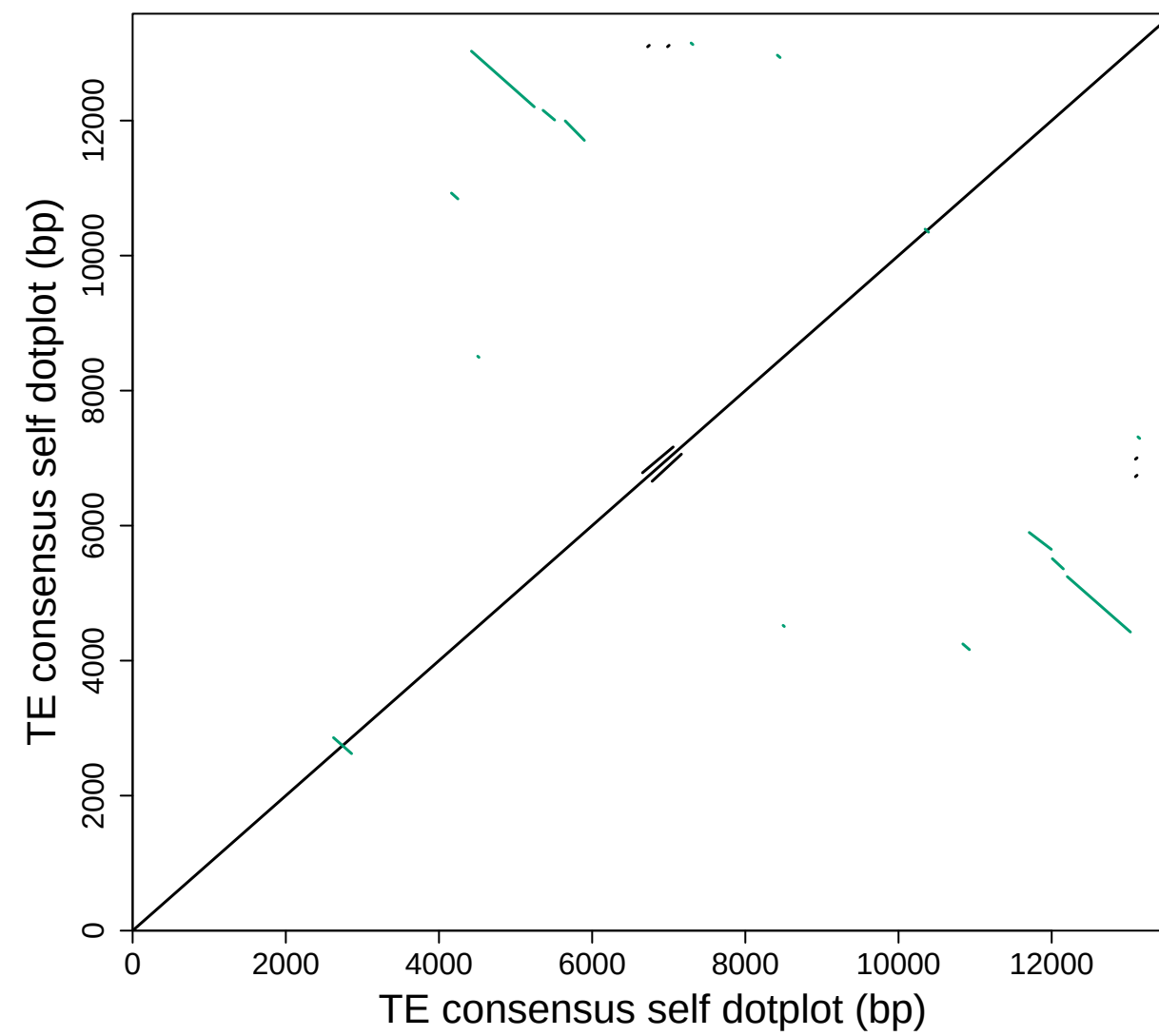
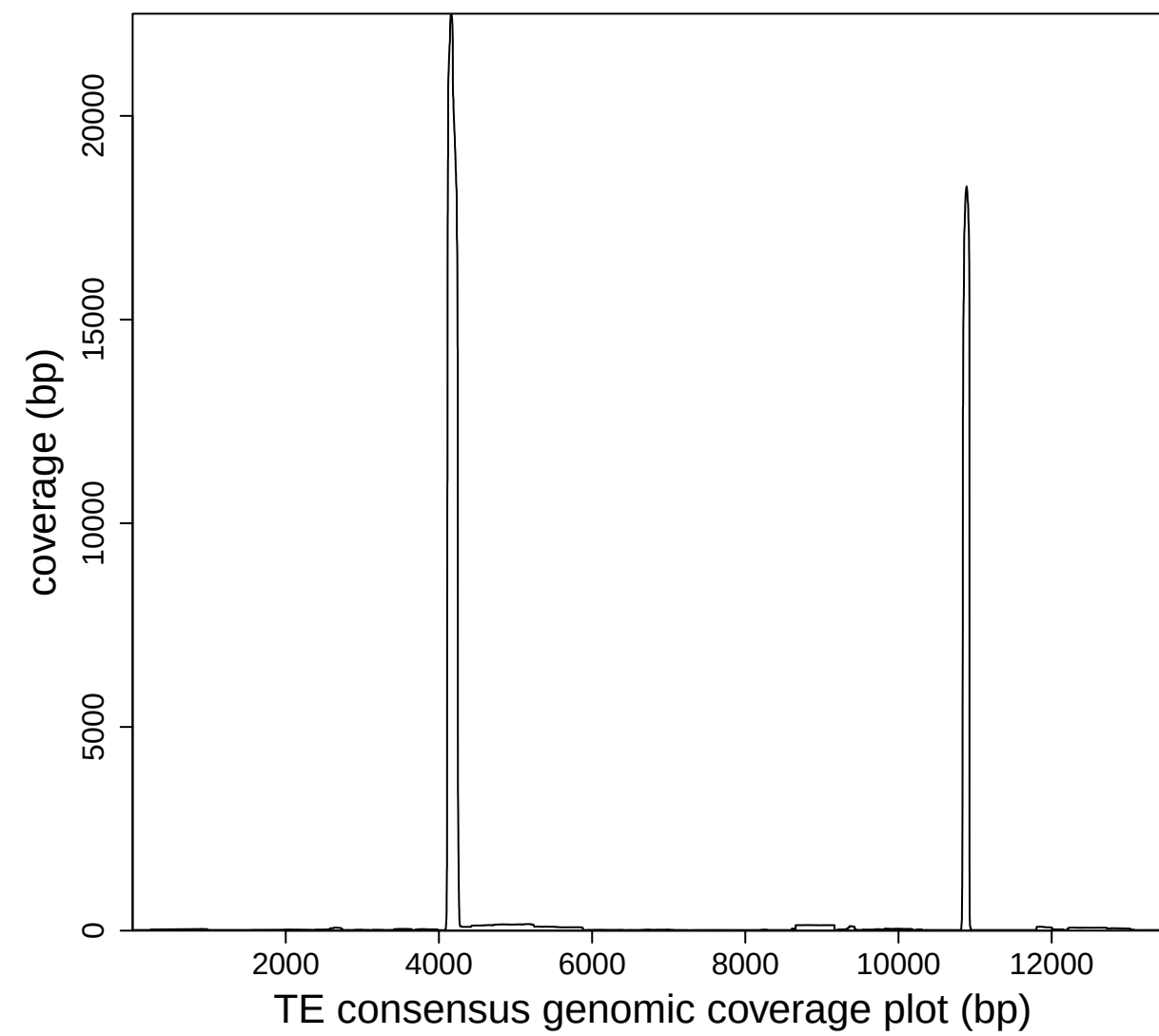
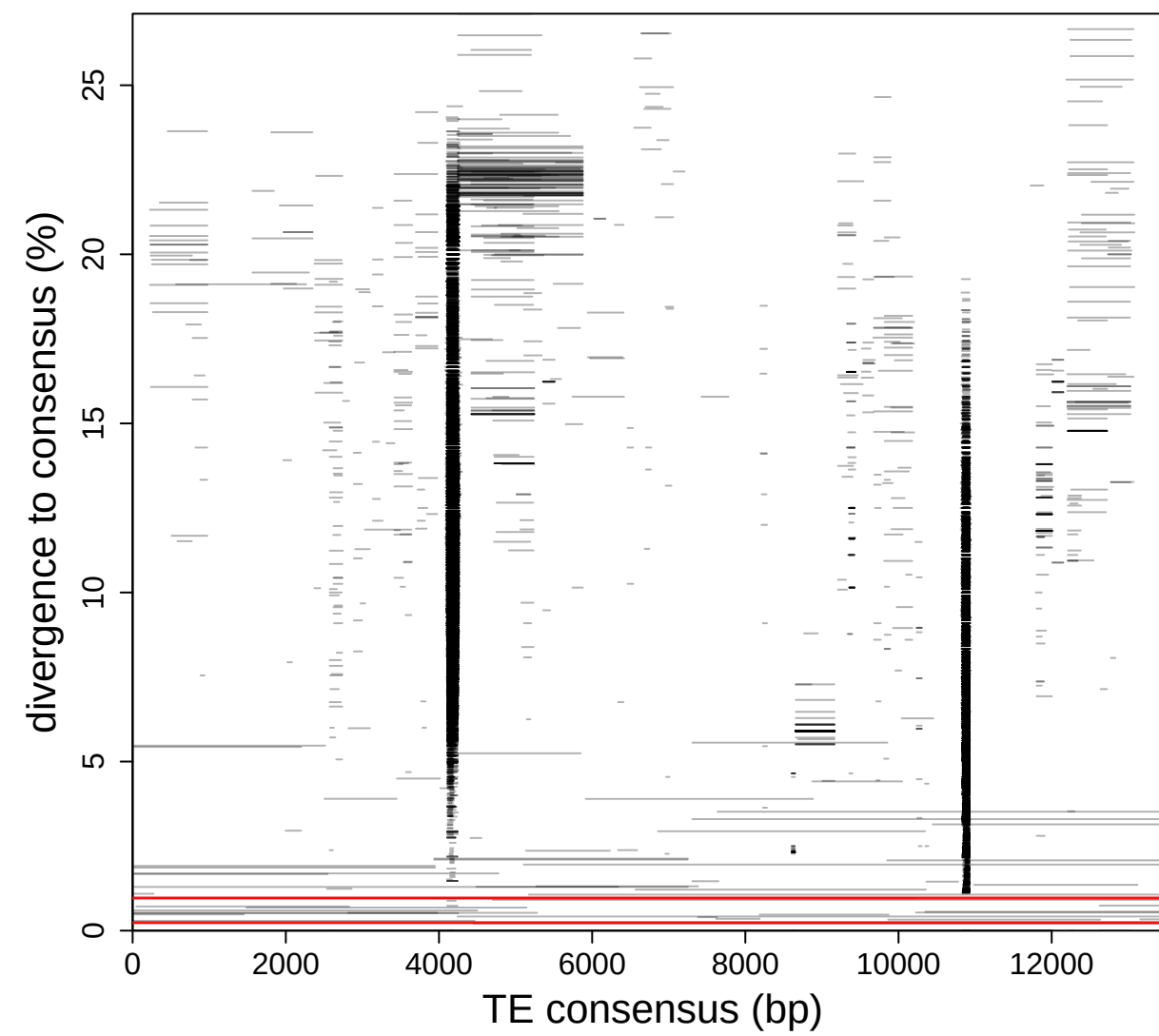
MSA length = 13583





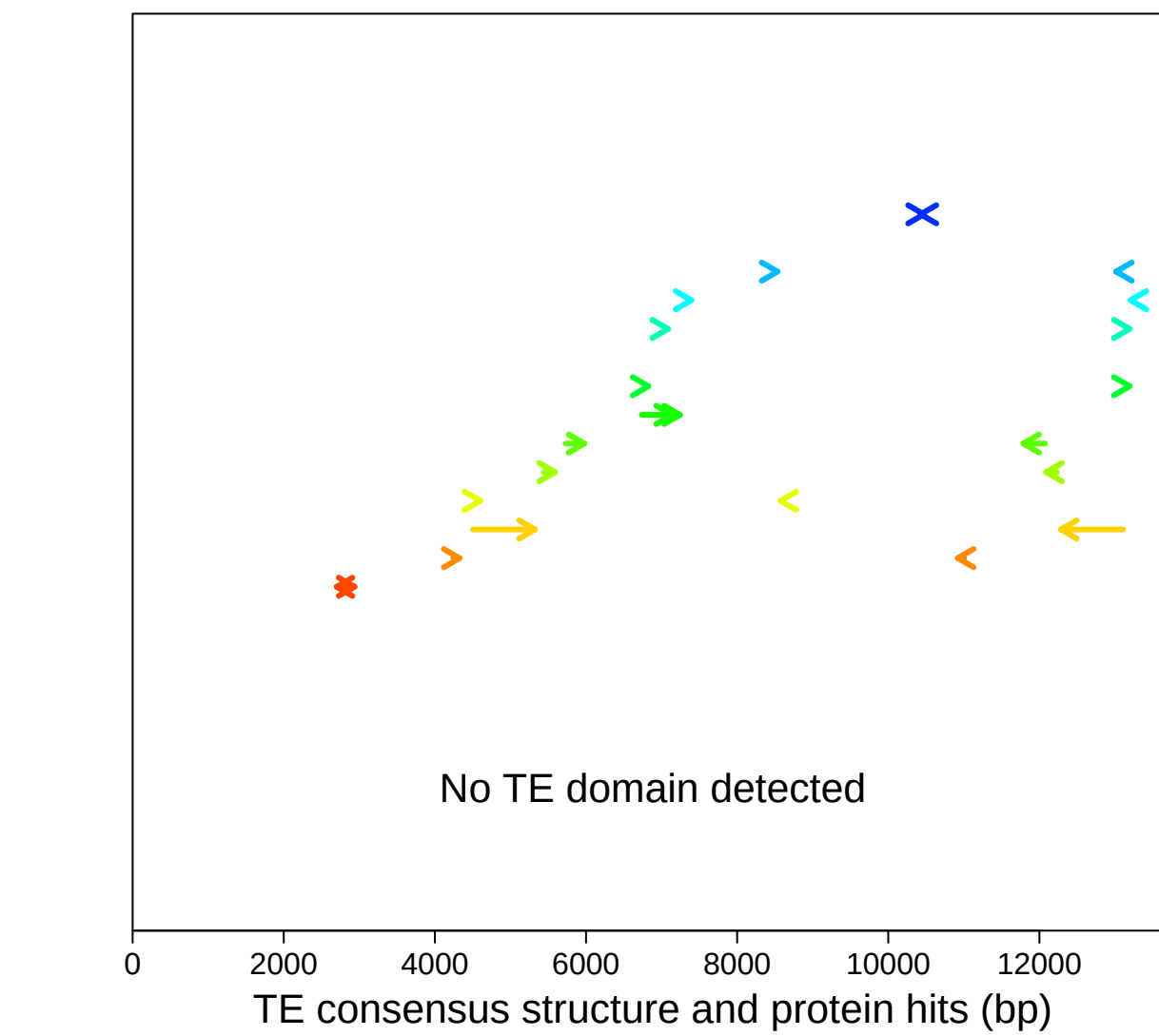
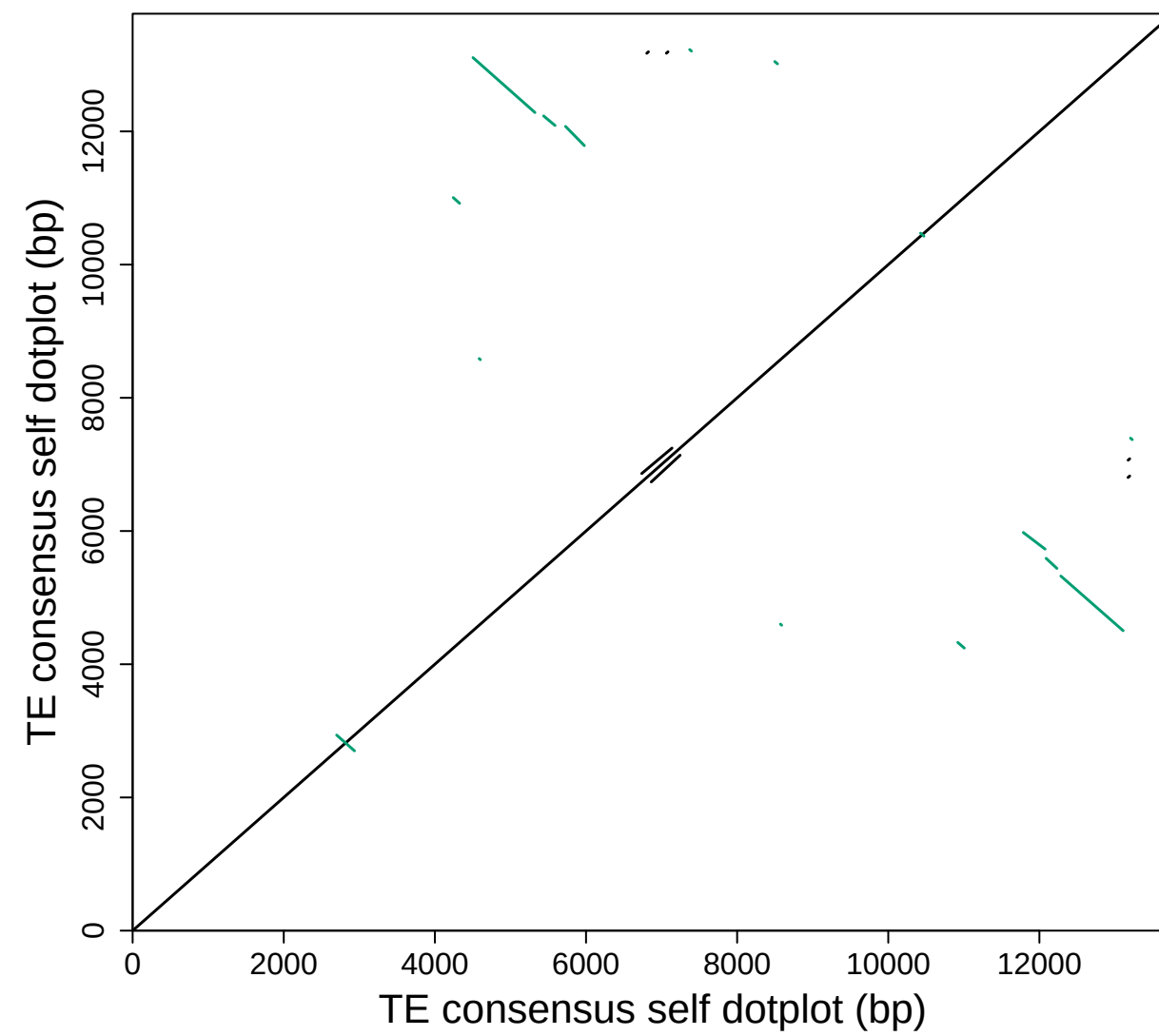
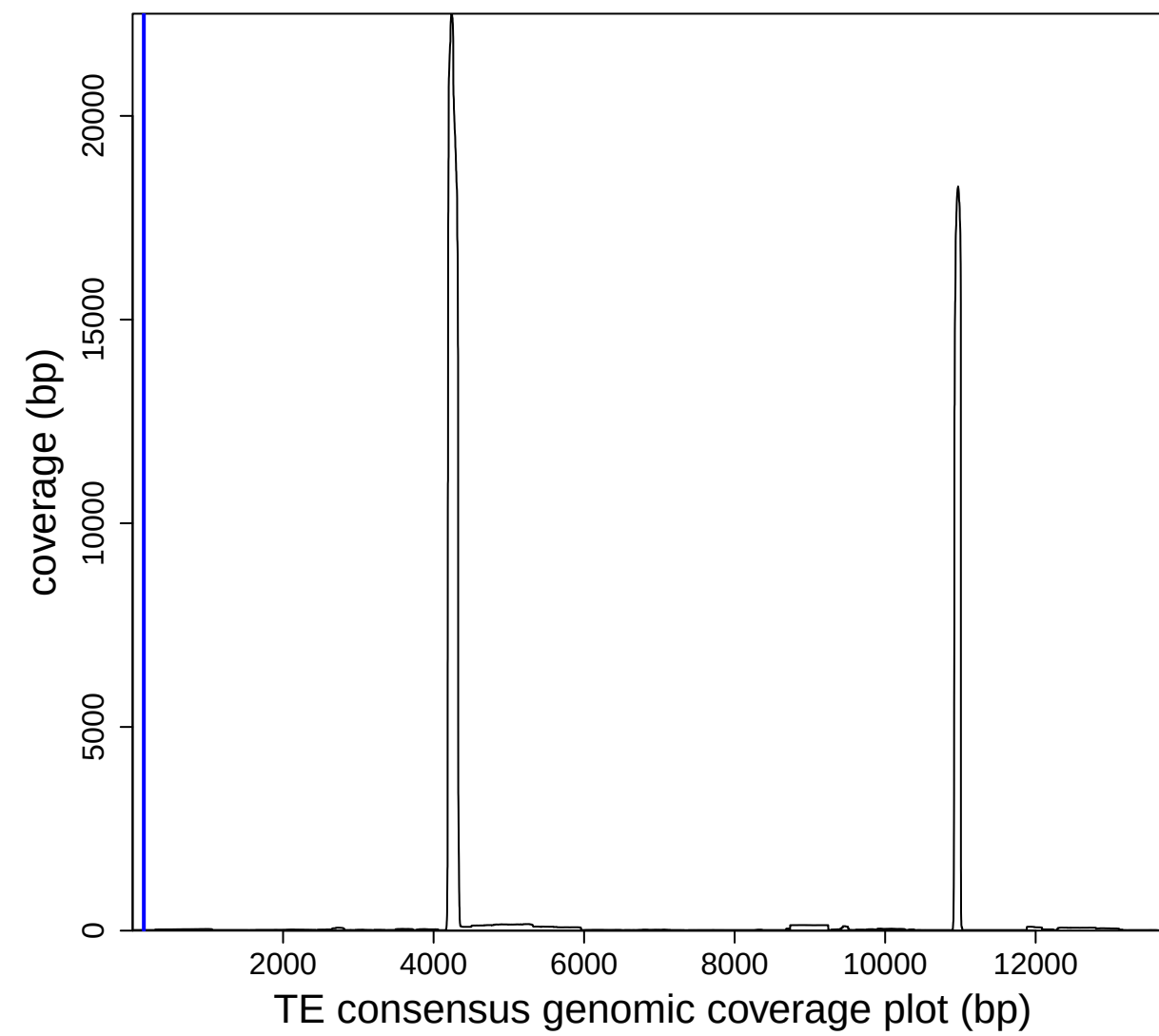
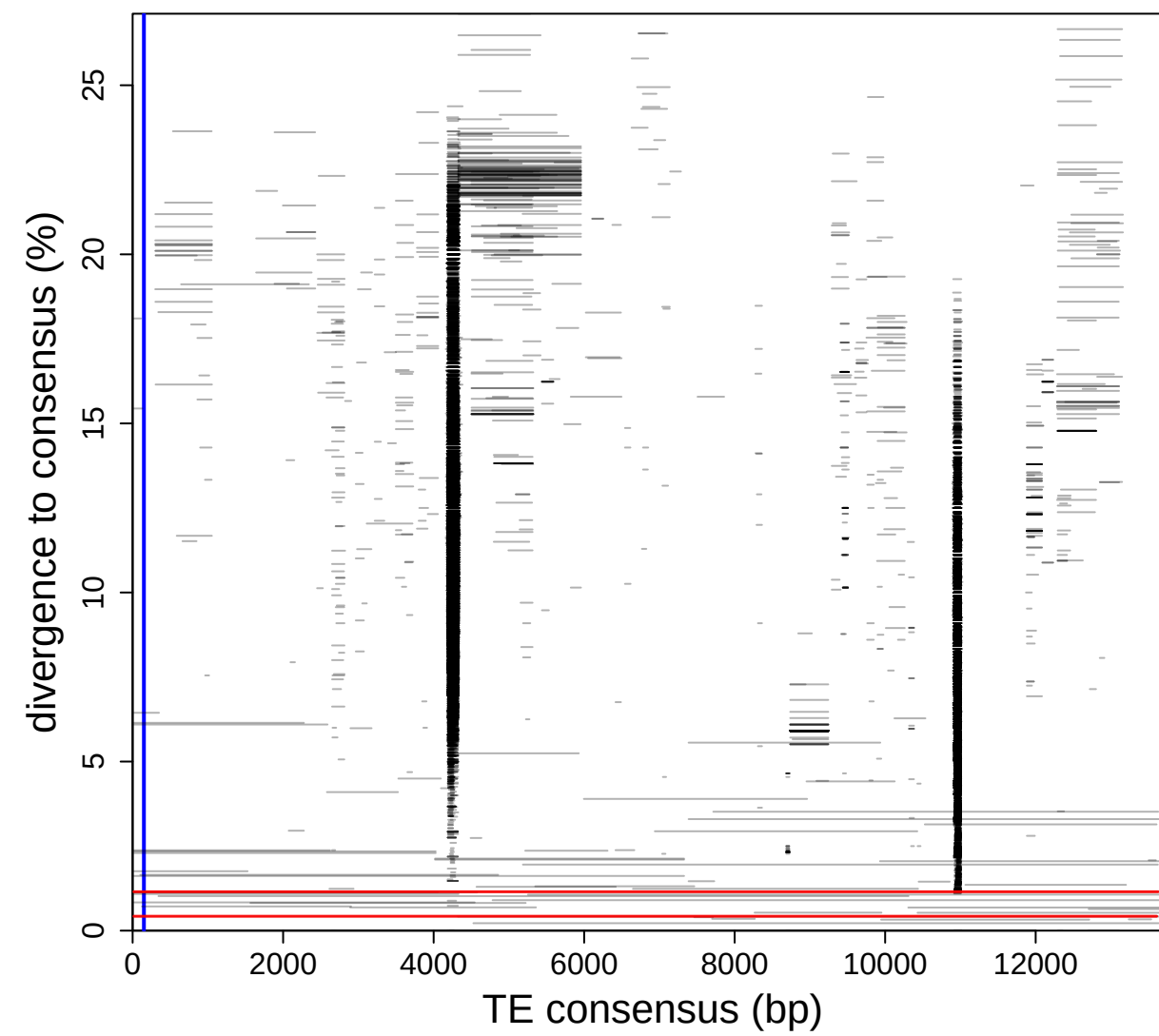
g_1.bed fm_1.bed 0 0 n.bed g_1.bed fm_2.bed 0 0 bcln.fa a
size: 13583bp; fragments: 41778; full length: 2 (>=12224.7bp)

After TEtrimmer 13583 bp

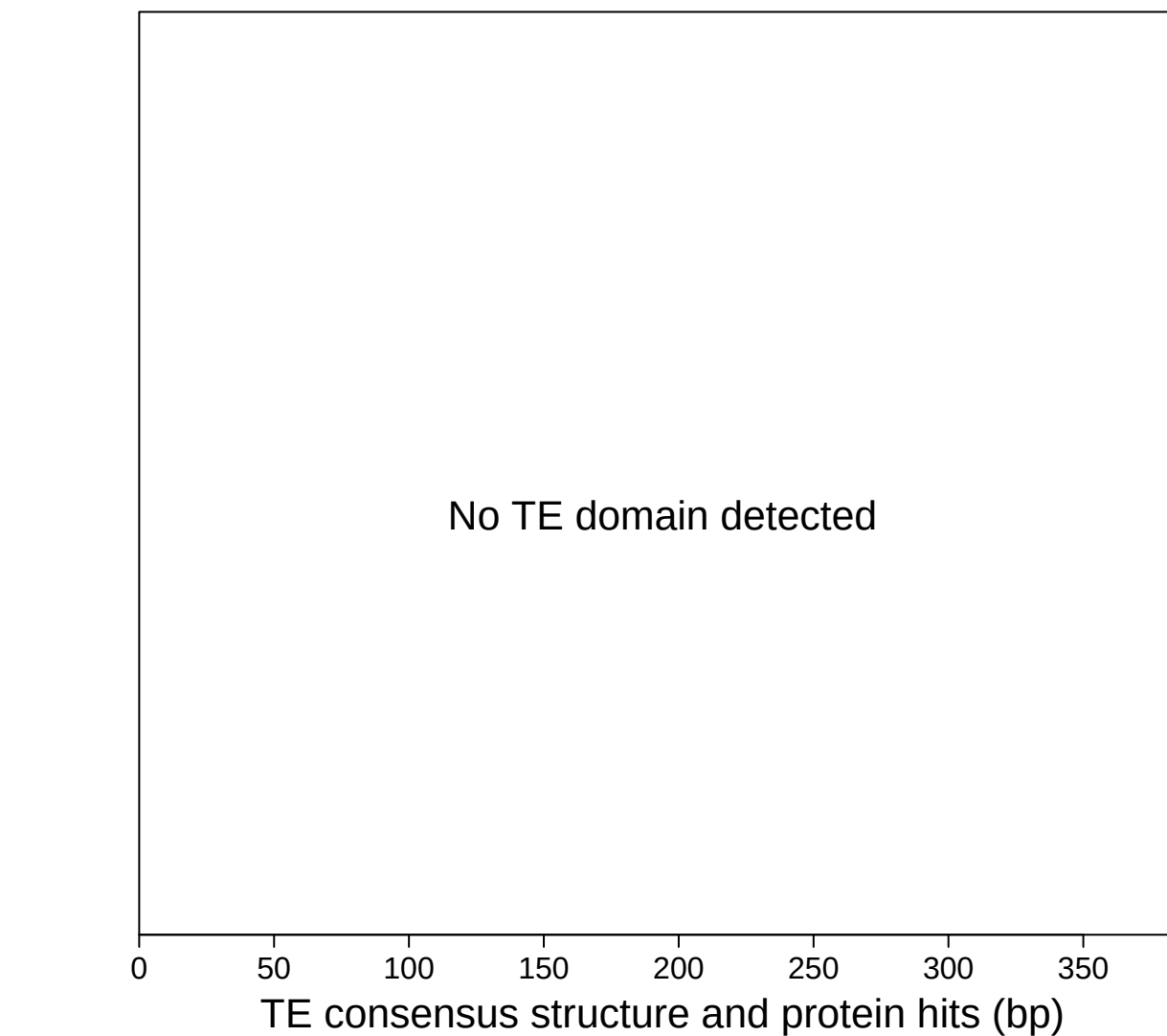
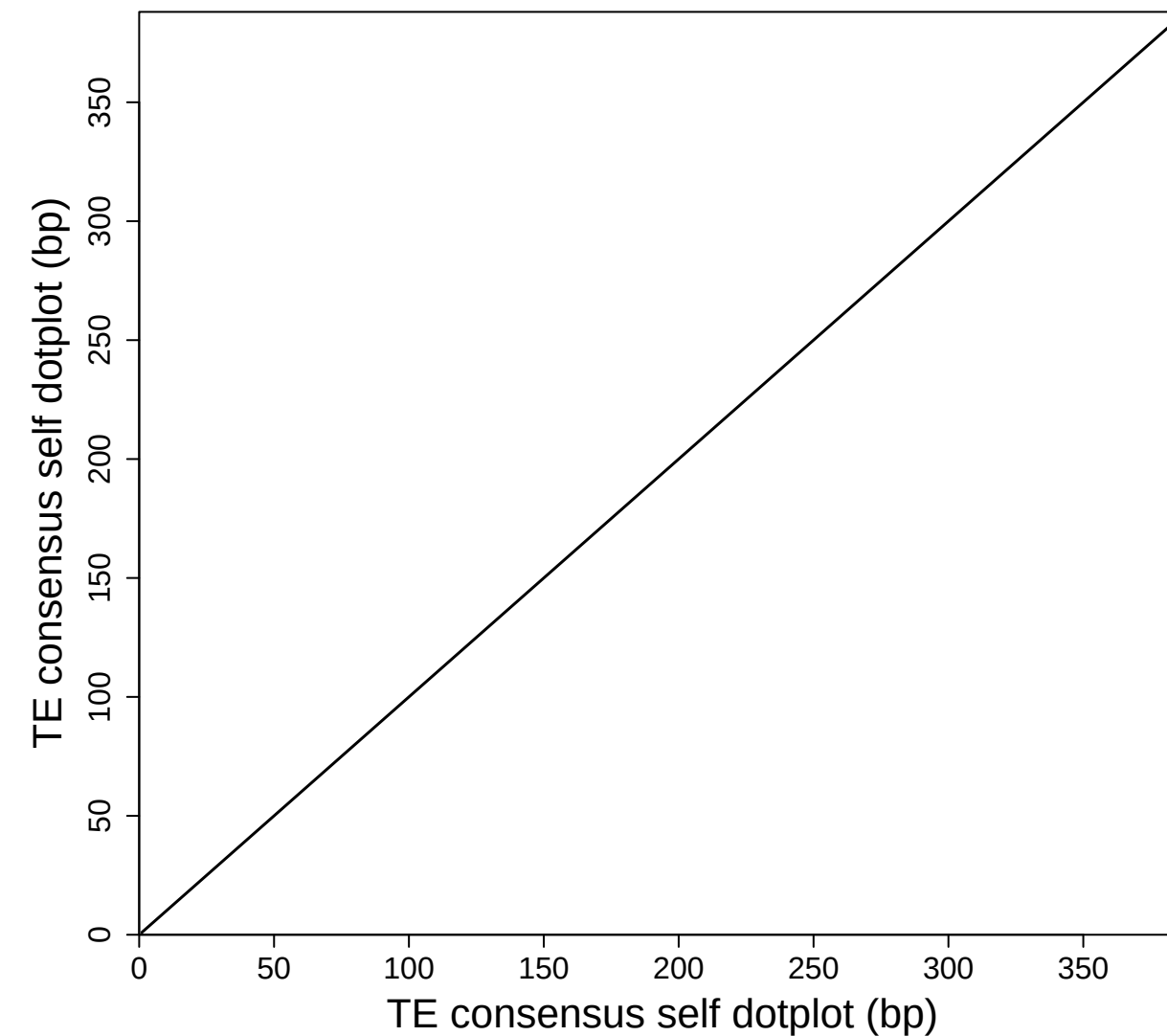
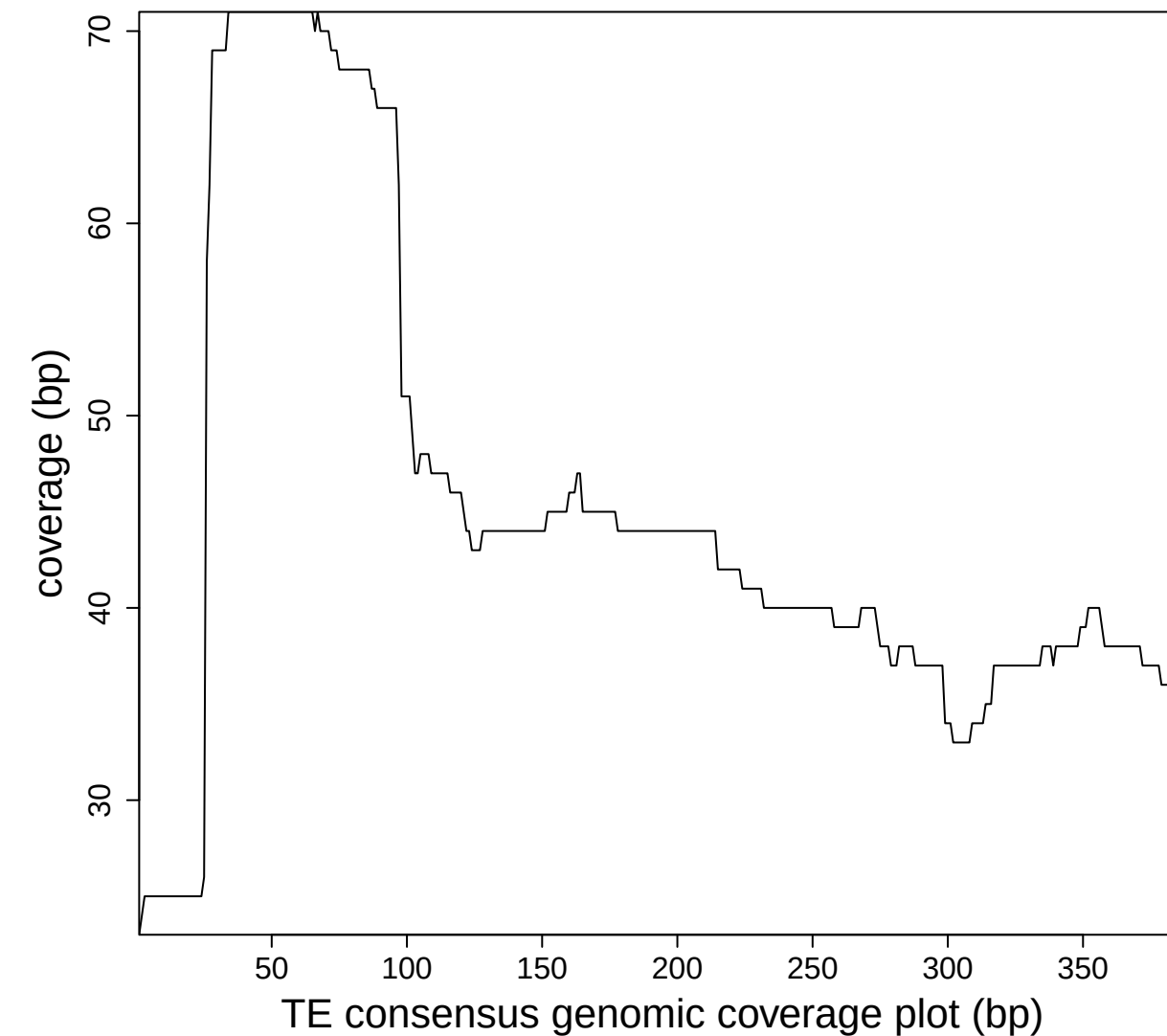
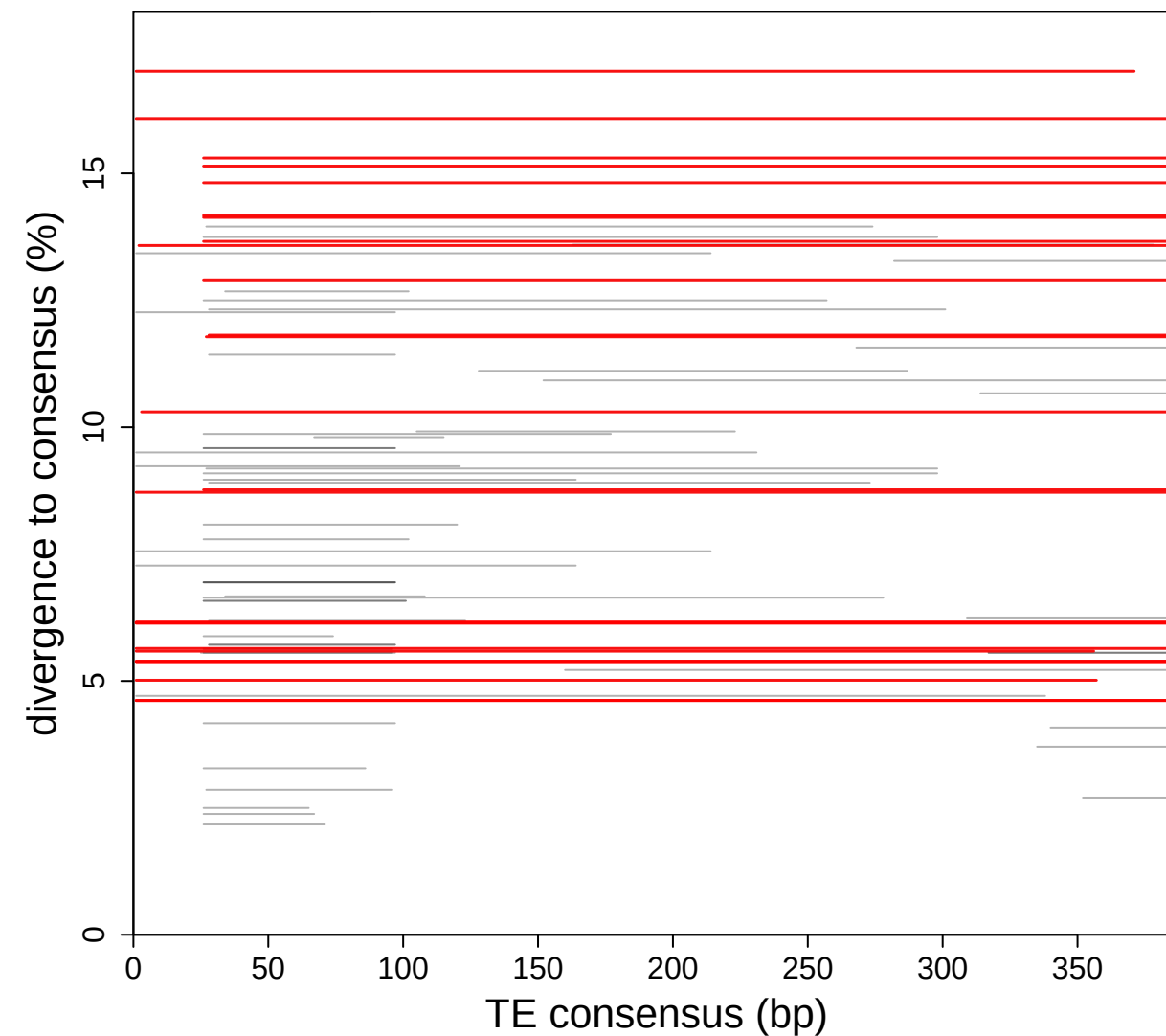


ed_g_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcln.f
size: 13824bp; fragments: 41779; full length: 2 (>=12441.6bp)

After TEtrimmer Extended plot Blue lines are boundaries

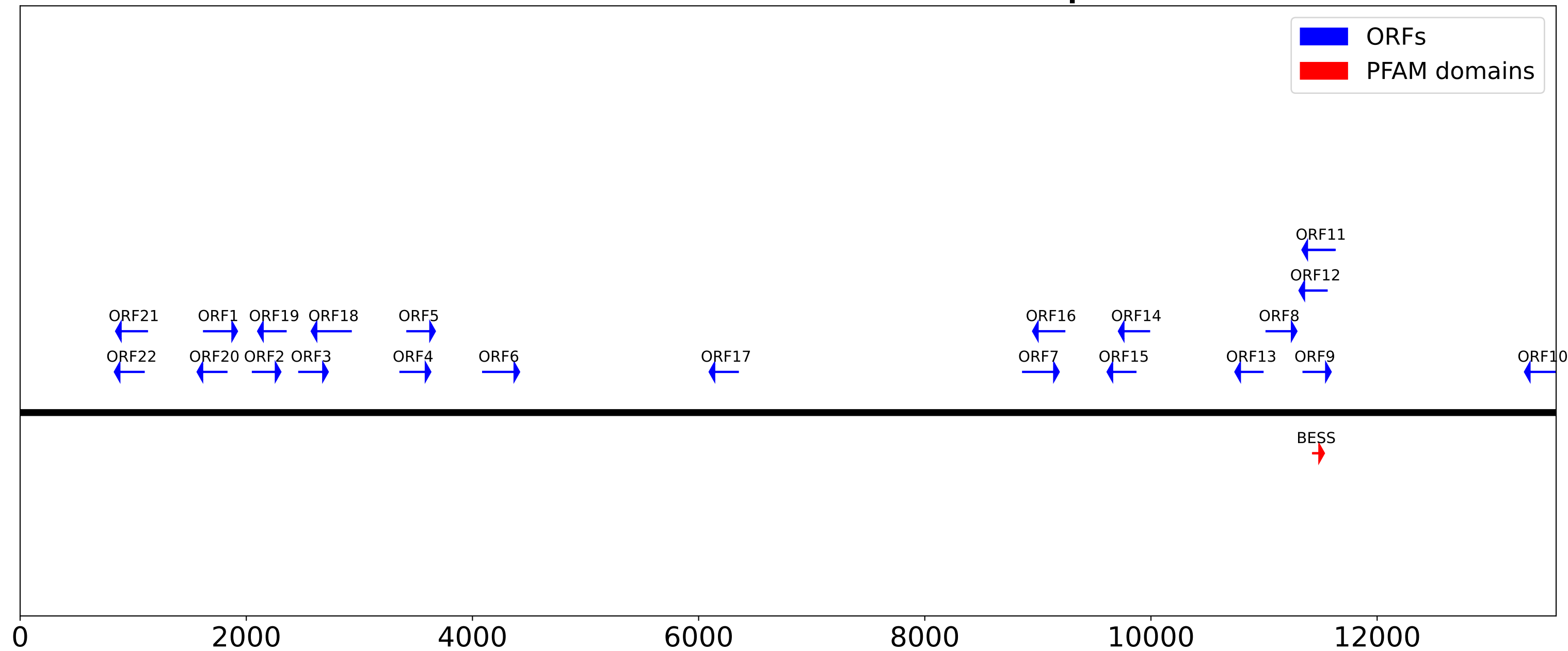


TE: rnd_5_family_2443
size: 388bp; fragments: 87; full length: 27 (≥ 349.2 bp)



Before TEtrimmer 388 bp

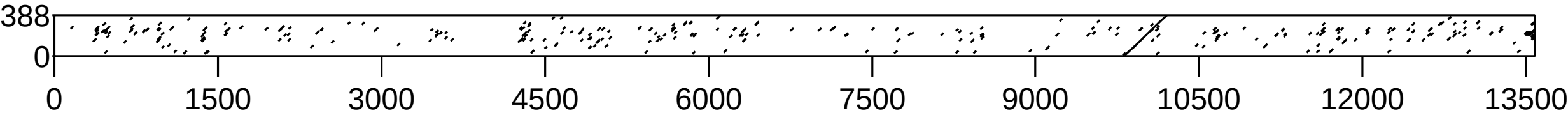
After TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)