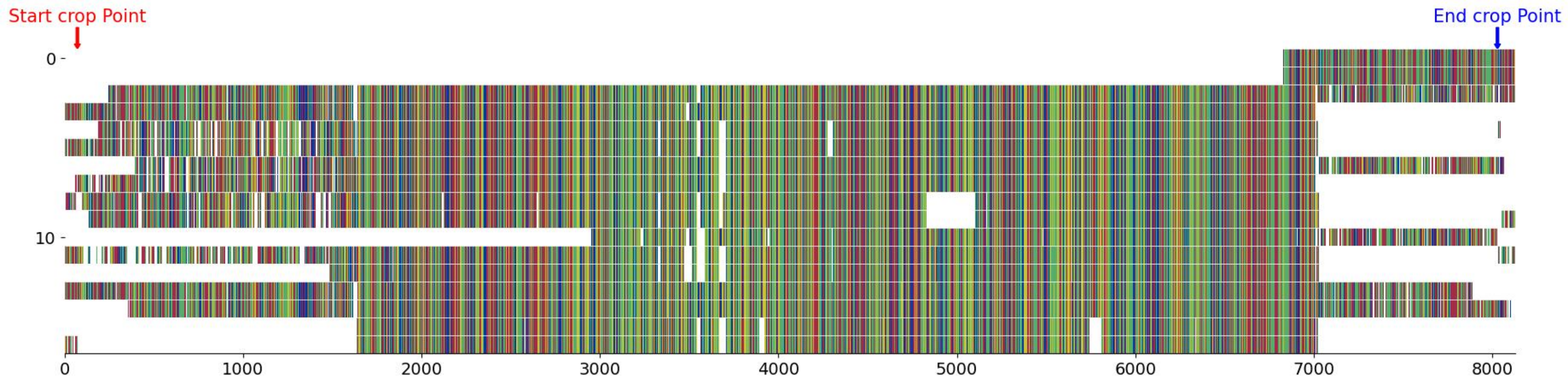


Start crop Point

End crop Point

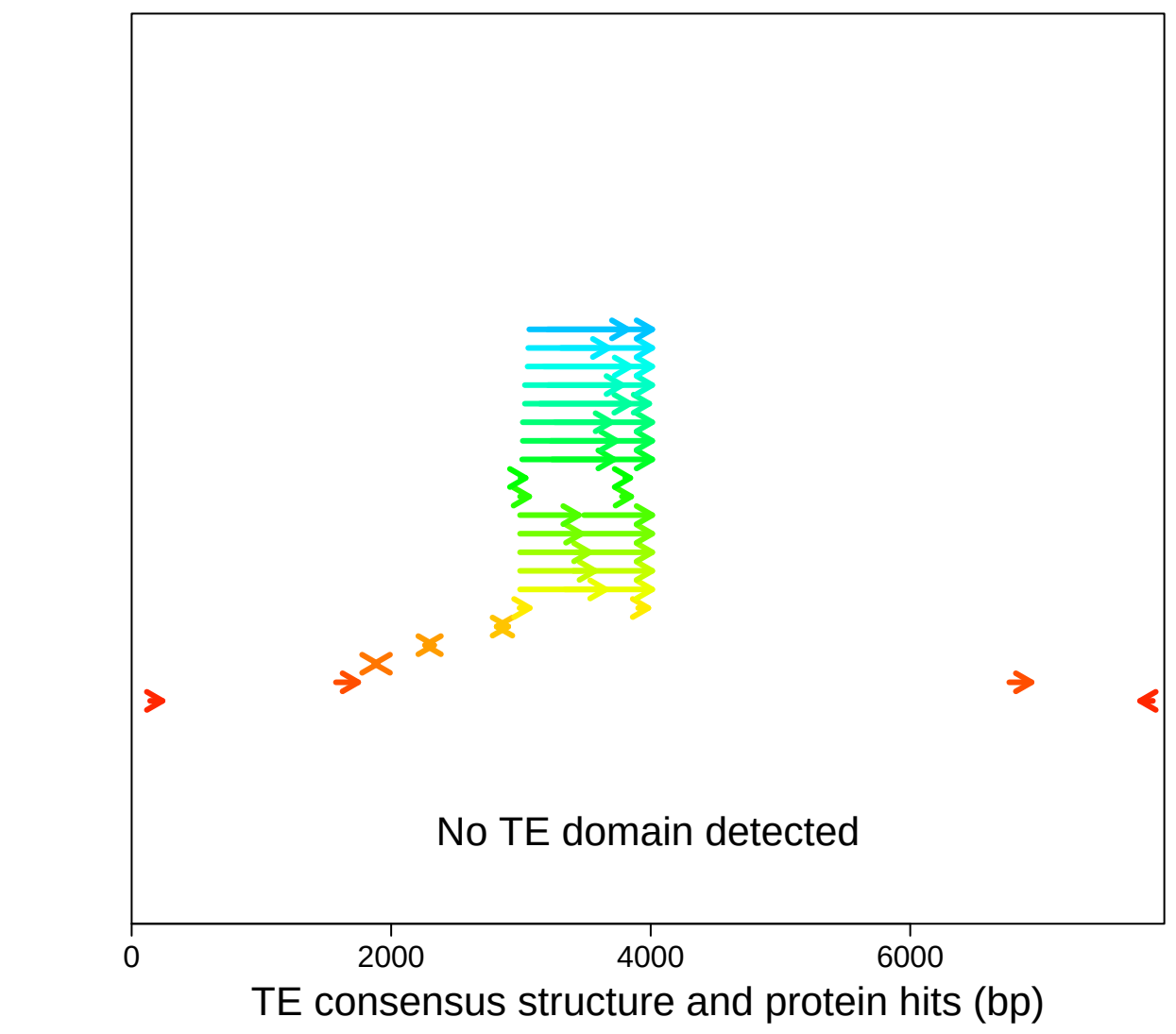
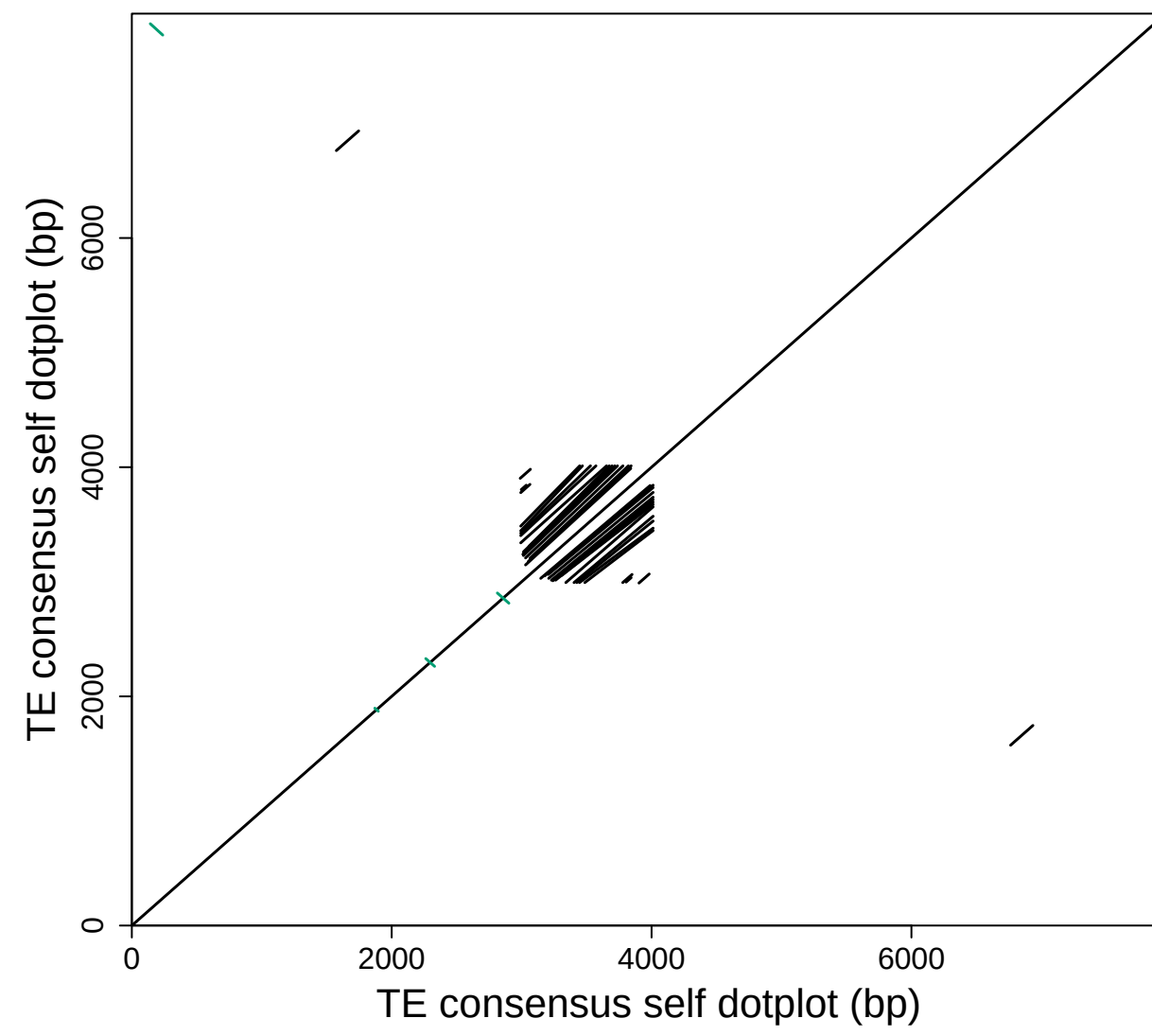
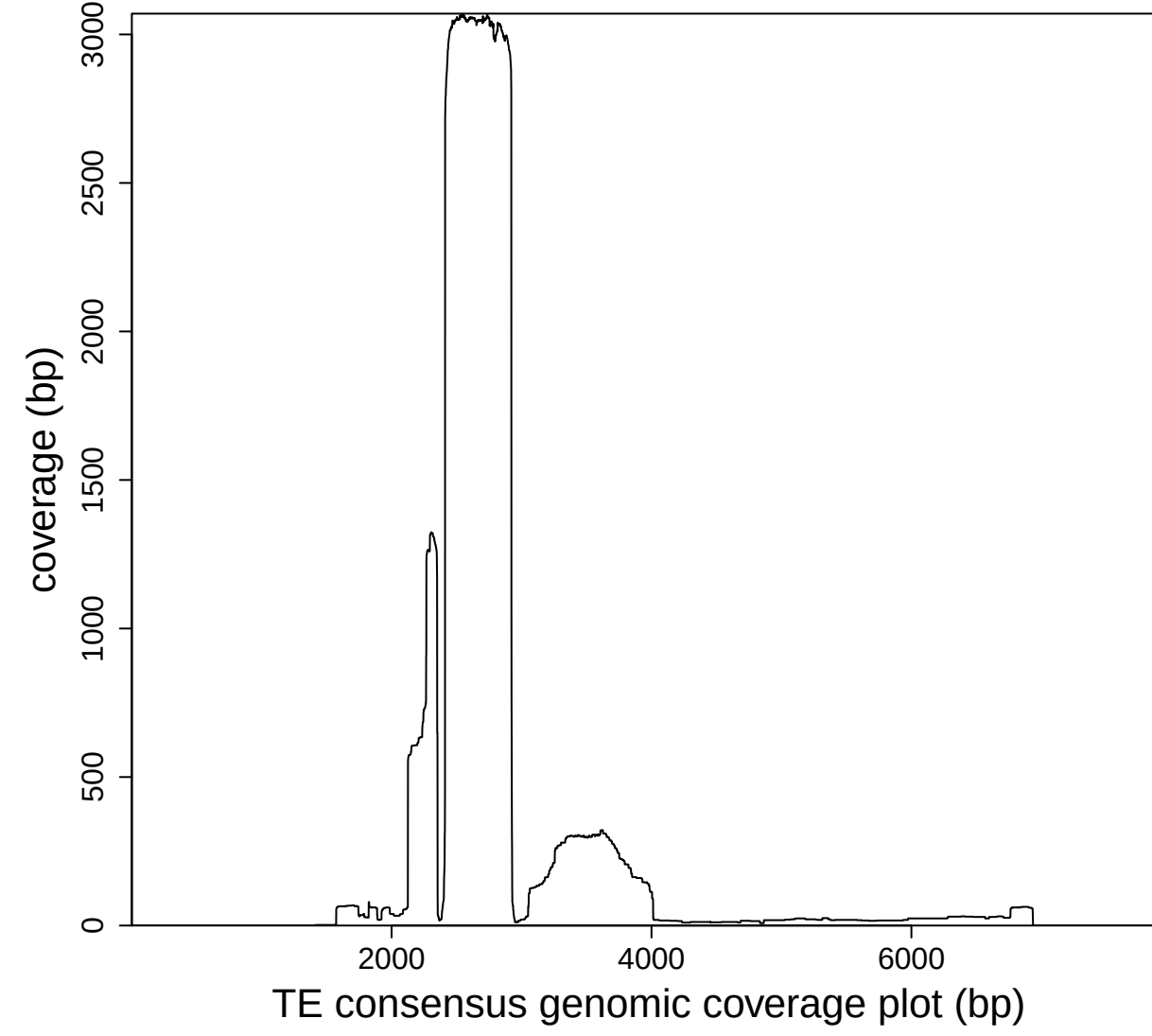
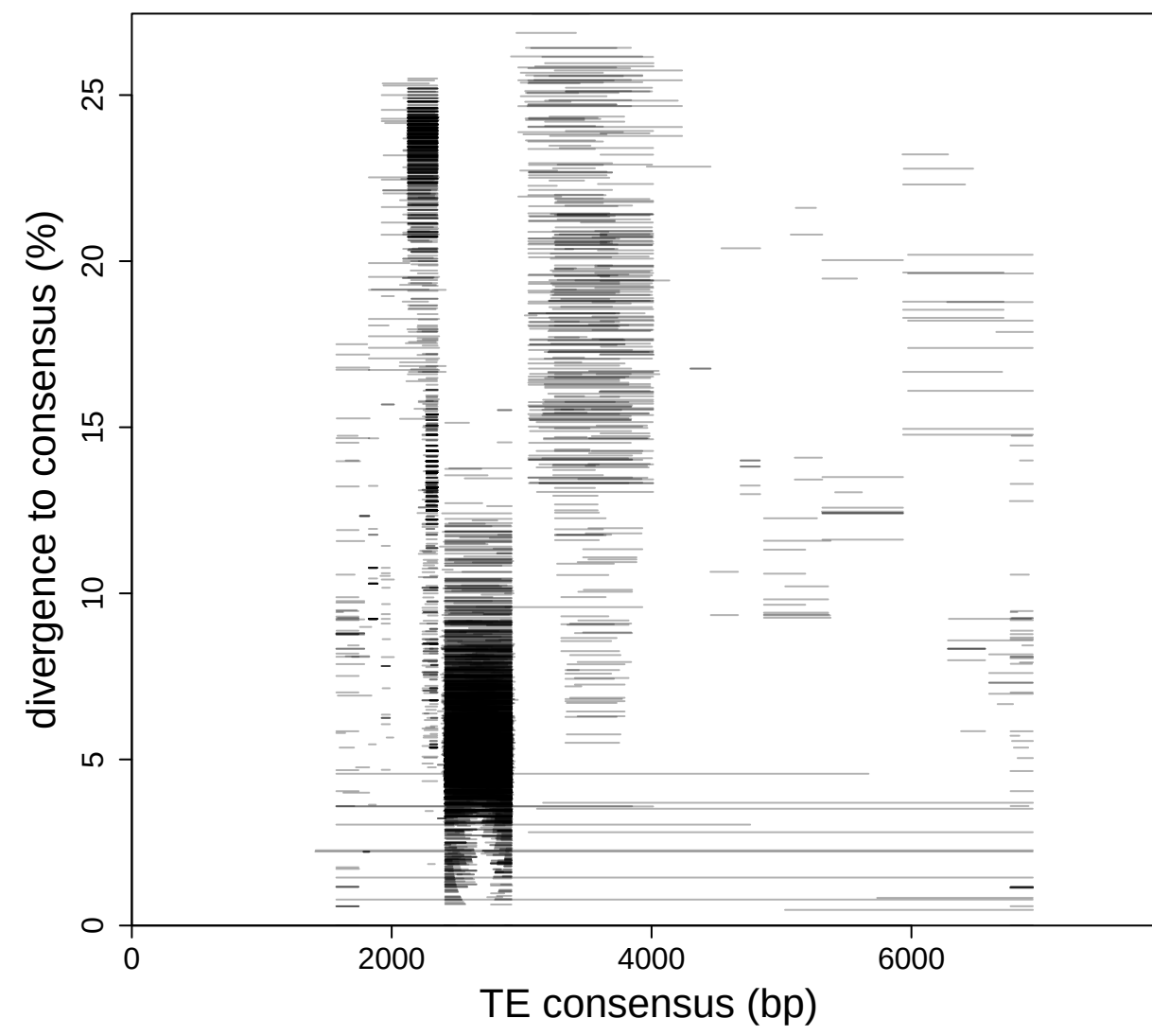
MSA length = 7958





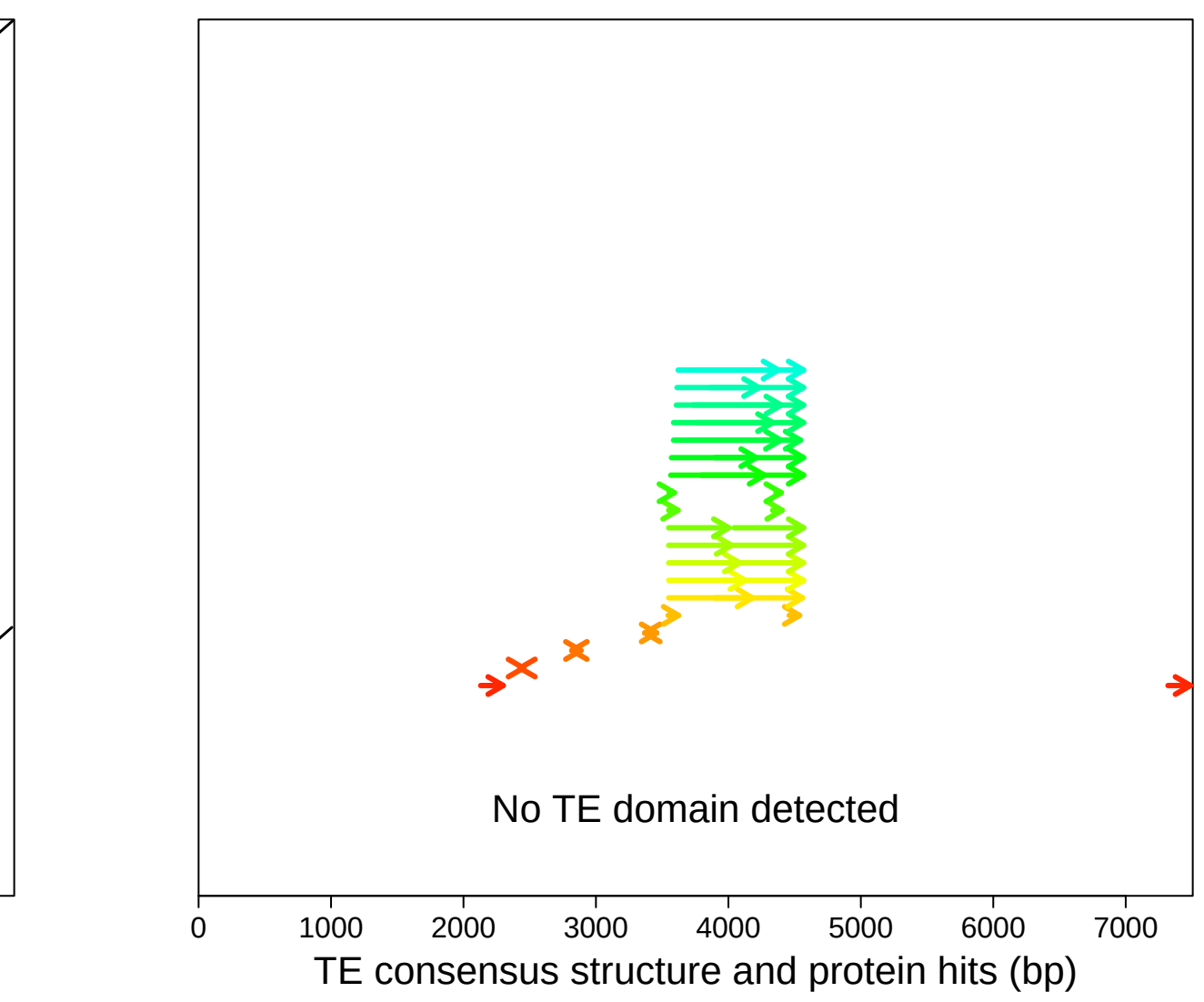
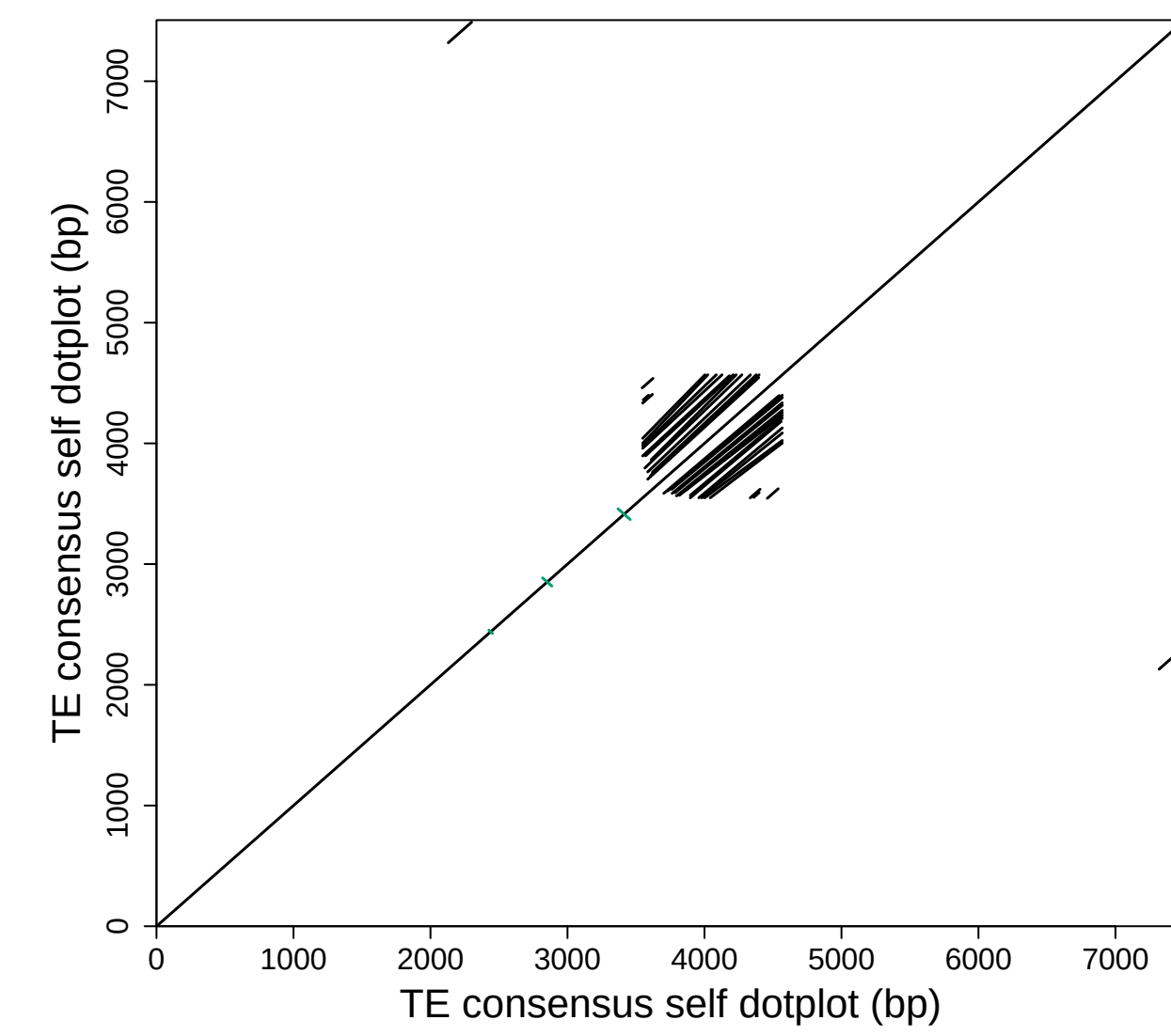
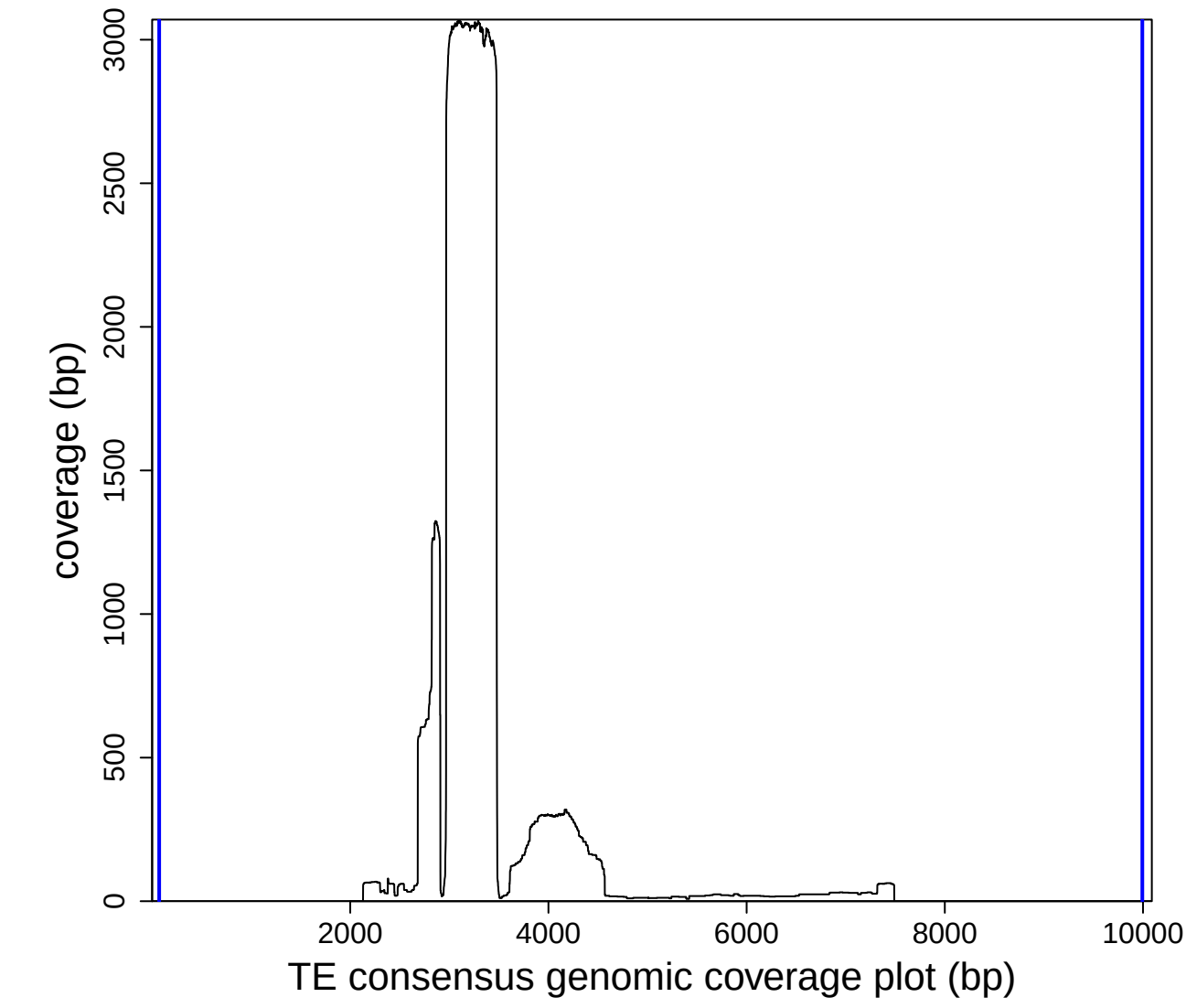
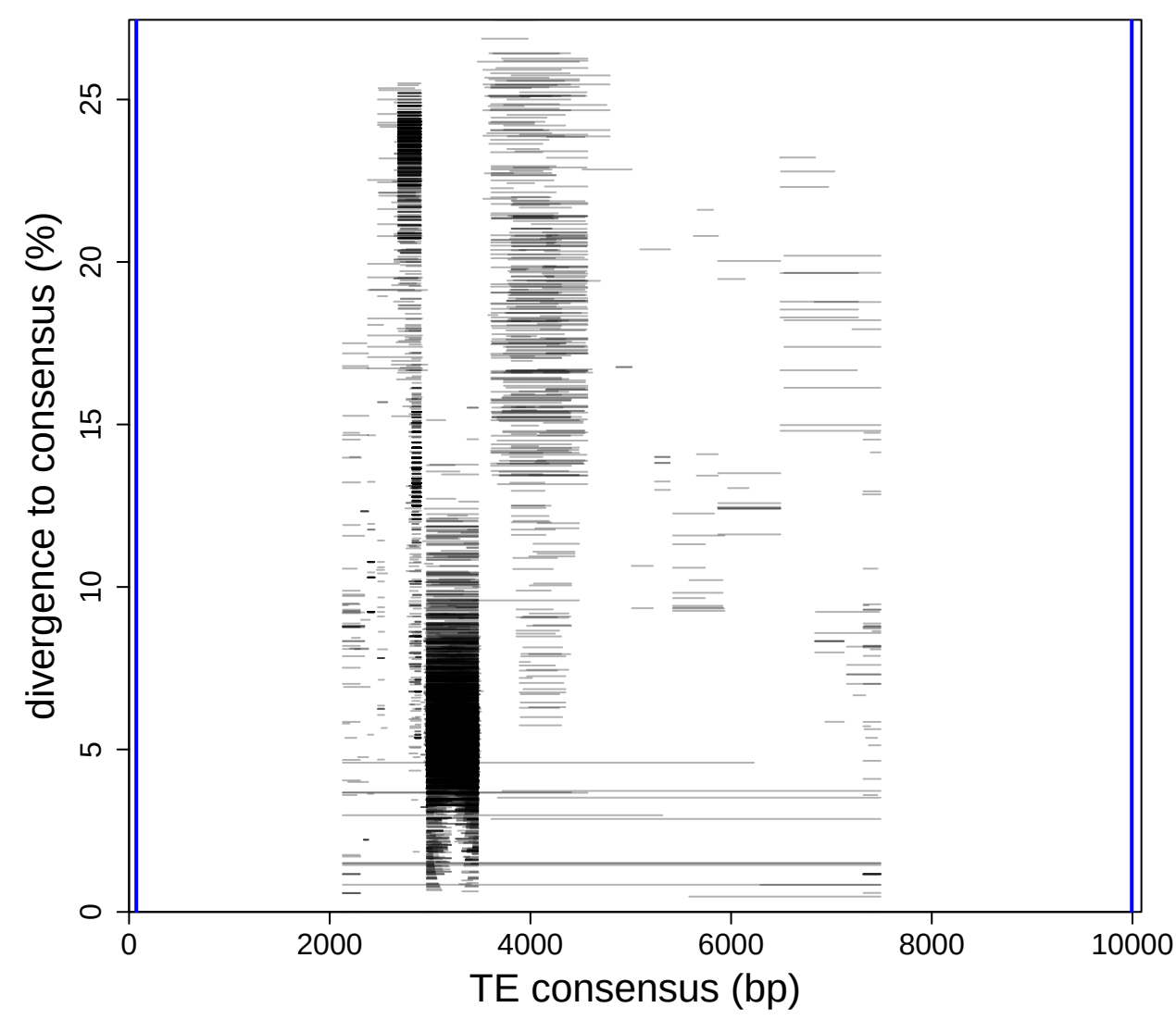
1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcln.fa_alr
size: 7958bp; fragments: 5661; full length: 0 (>=7162.2bp)

After TEtrimmer 7958 bp



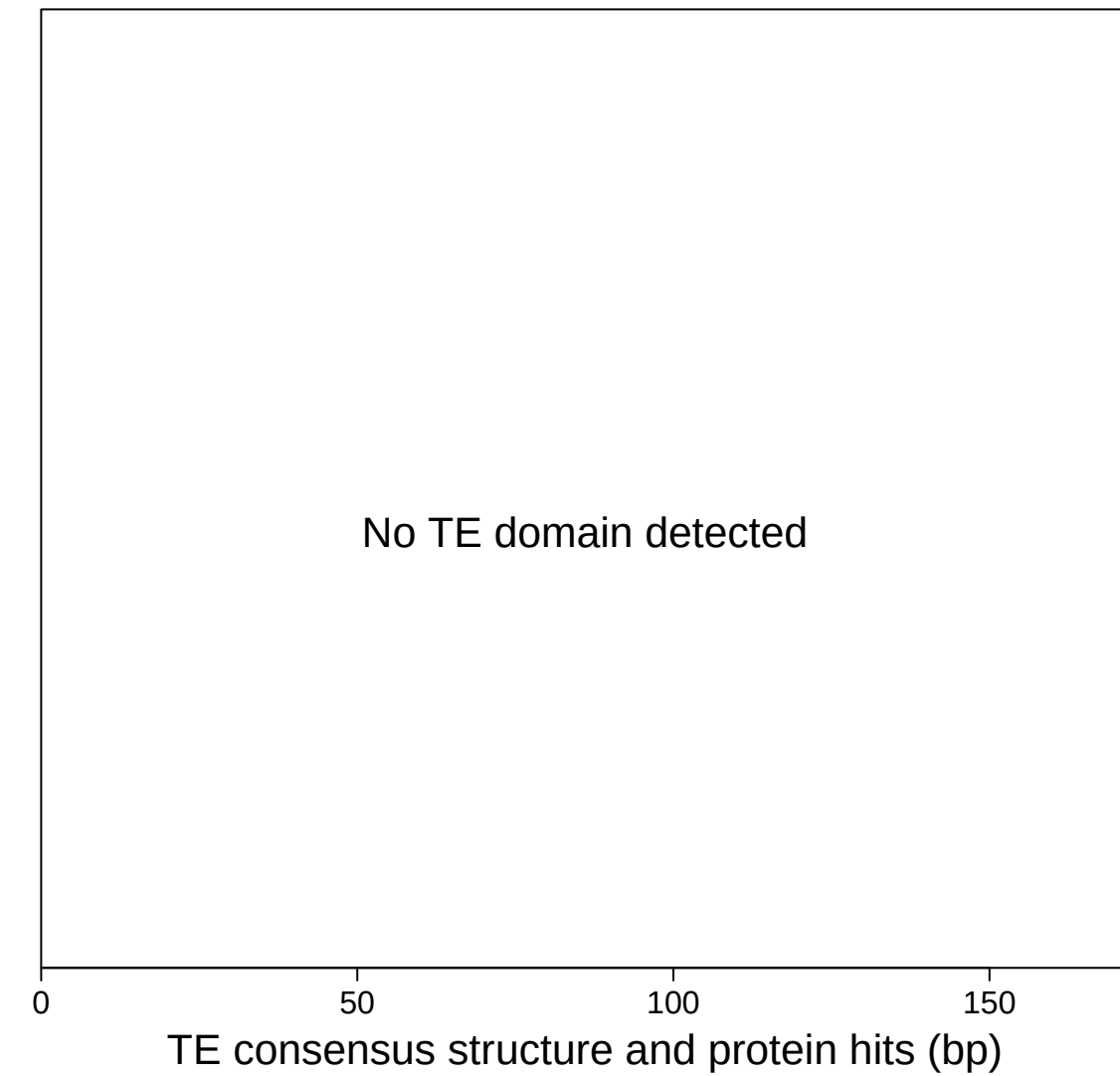
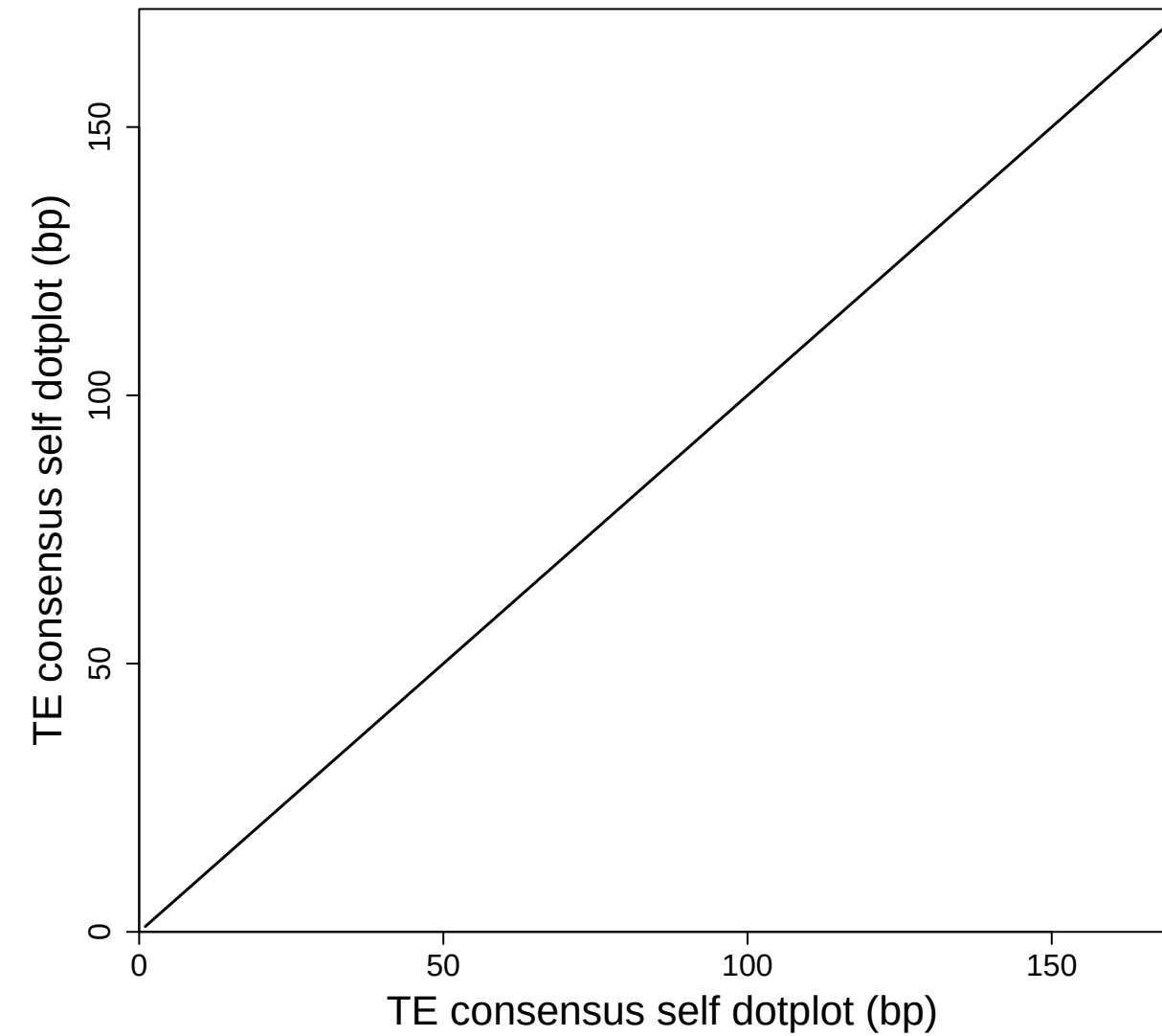
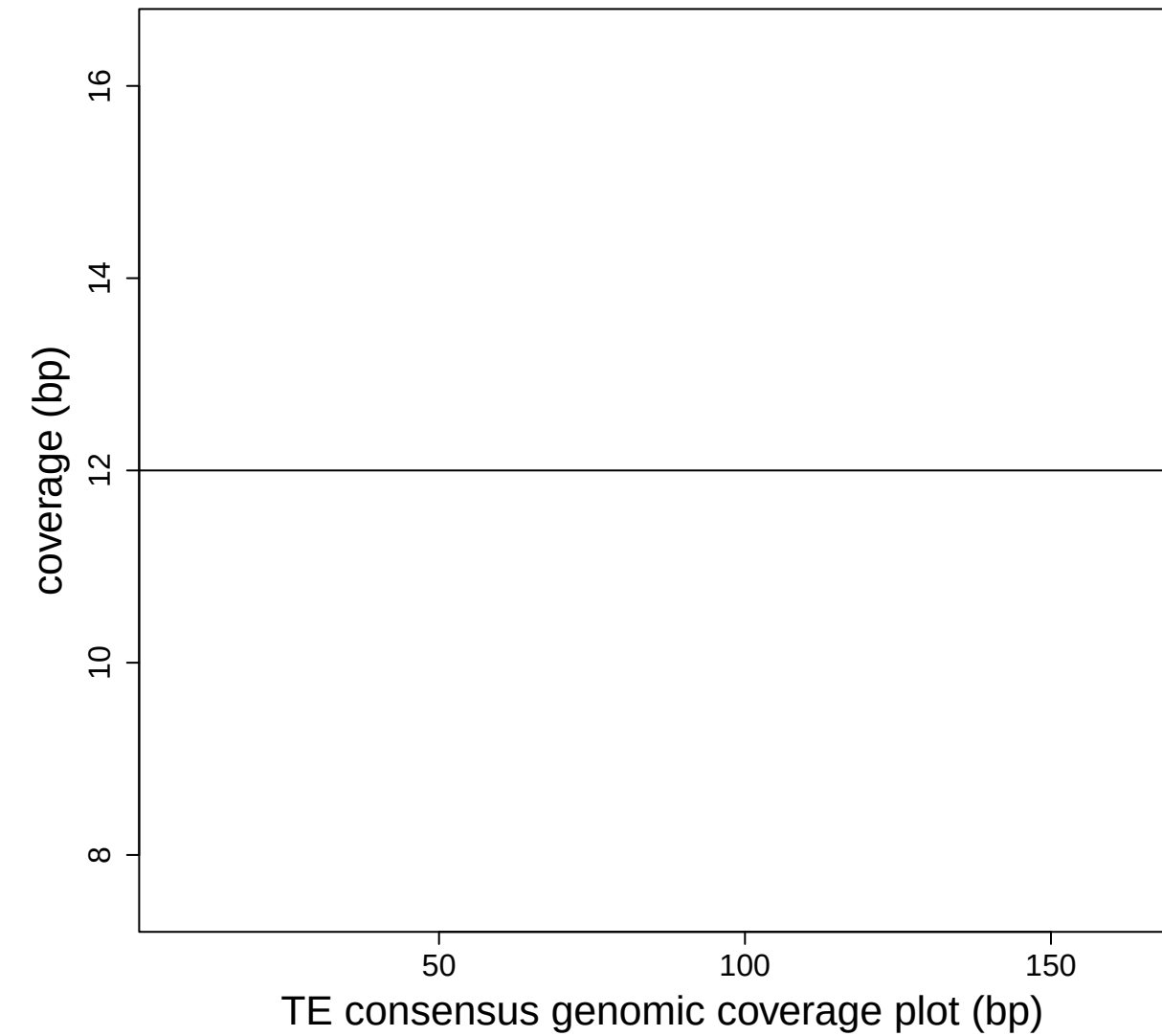
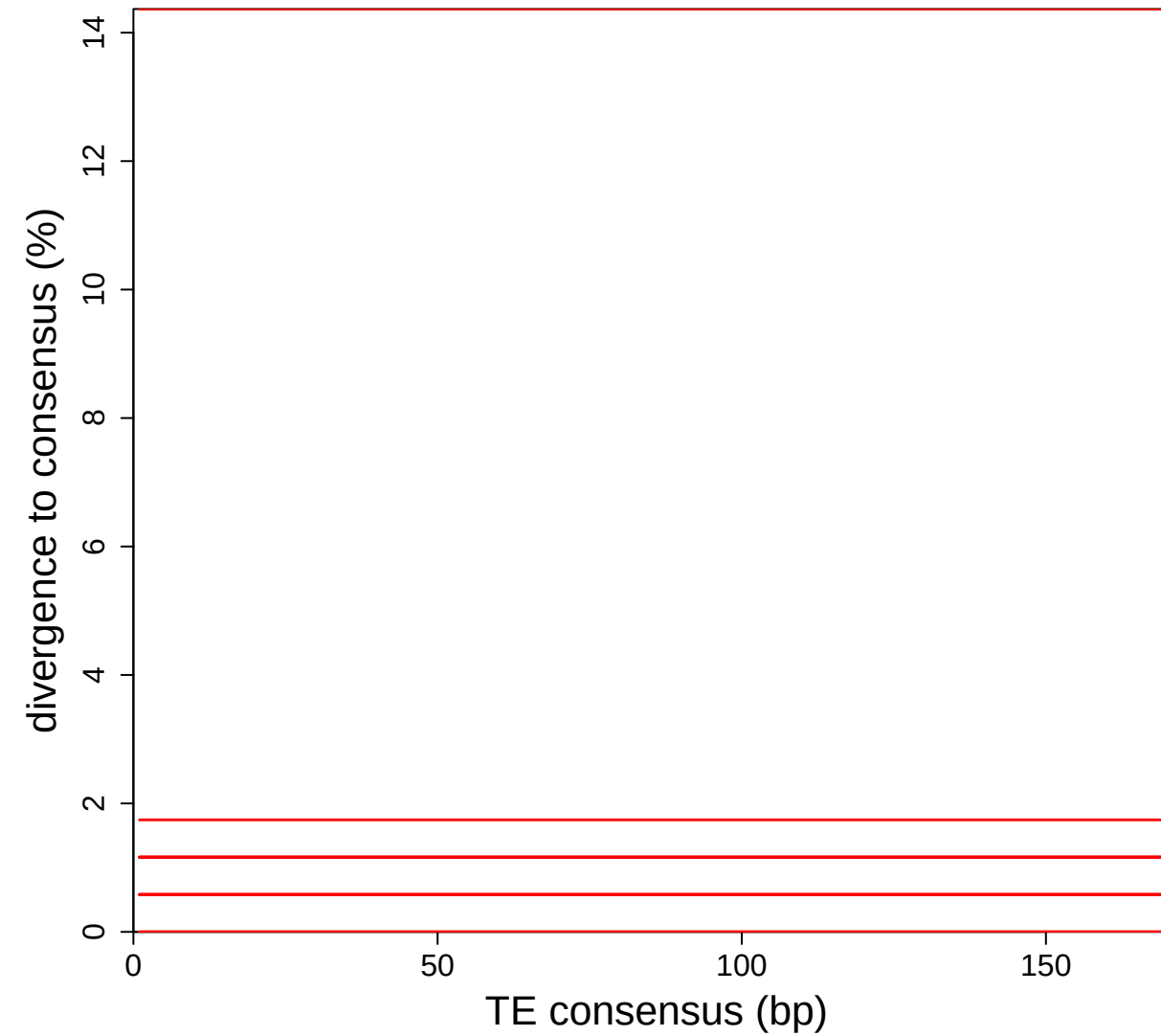
bed_g_1.bed fm_1.bed 0 0 n.bed_g_1.bed fm_2.bed 0 0 bcln
size: 10089bp; fragments: 5660; full length: 0 (>=9080.1bp)

After TEtrimmer Extended plot Blue lines are boundaries

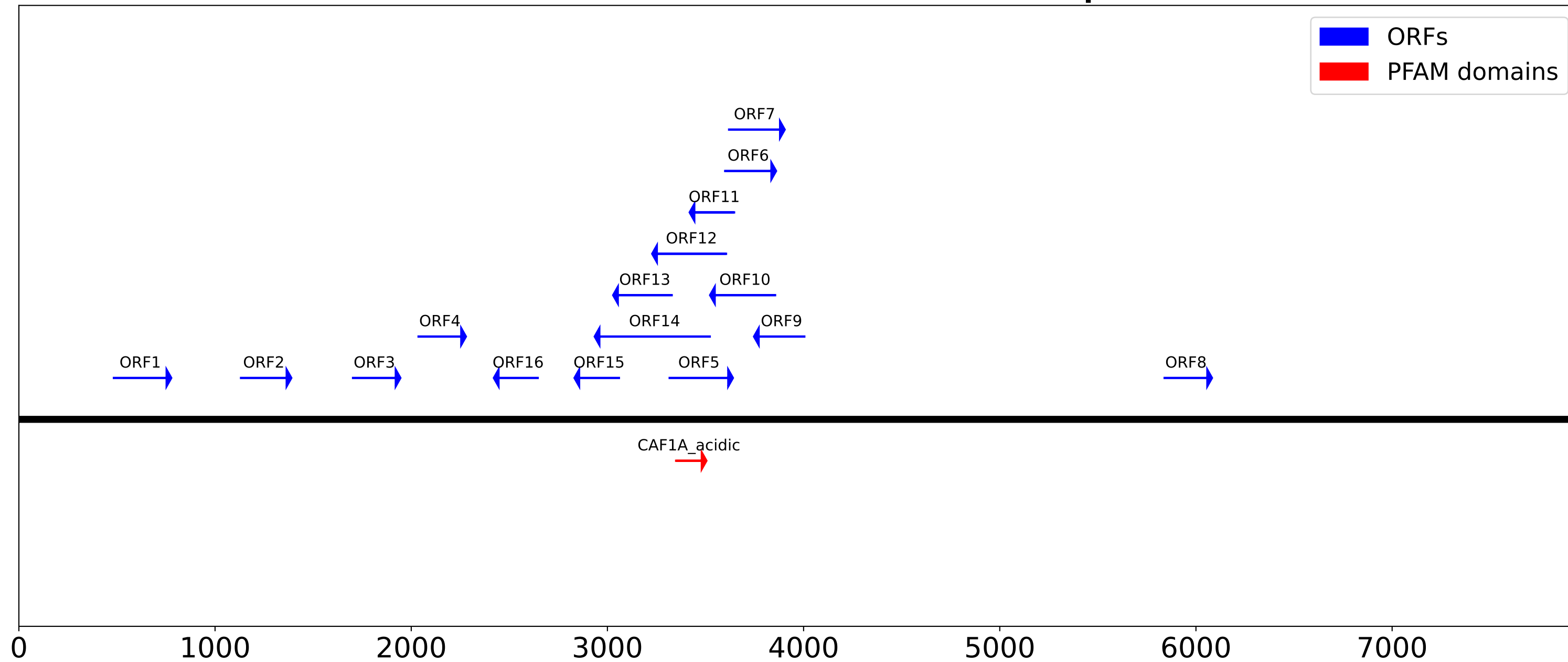


Before TEtrimmer 172 bp

TE: ltr_1_family_24
size: 172bp; fragments: 12; full length: 12 (≥ 154.8 bp)



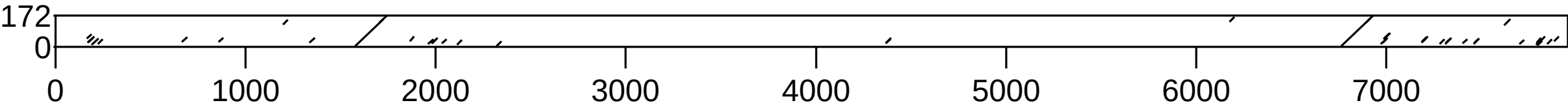
After TEtrimmer ORF and PFAM domain plot



Dotplot

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

1sus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)