

Start crop Point

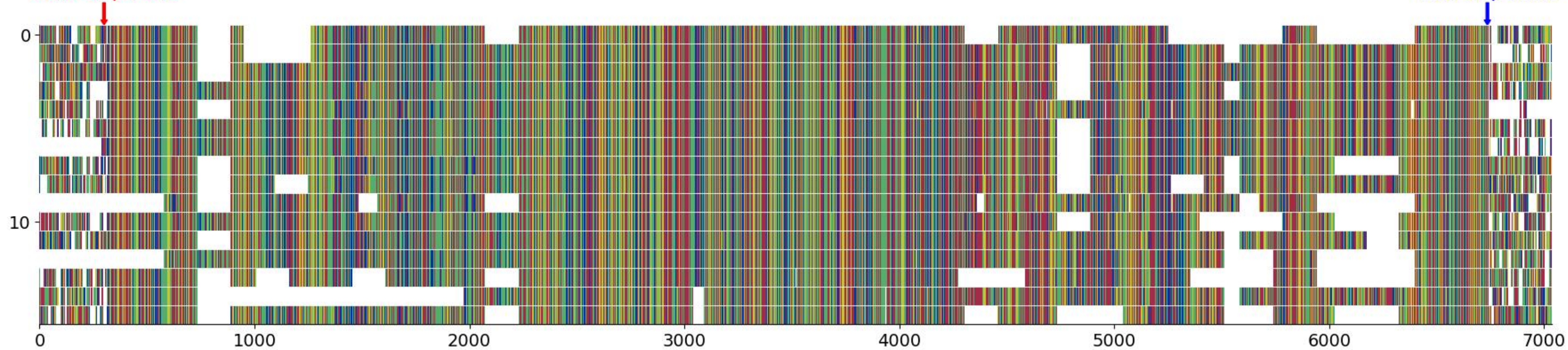
End crop Point

MSA length = 6437



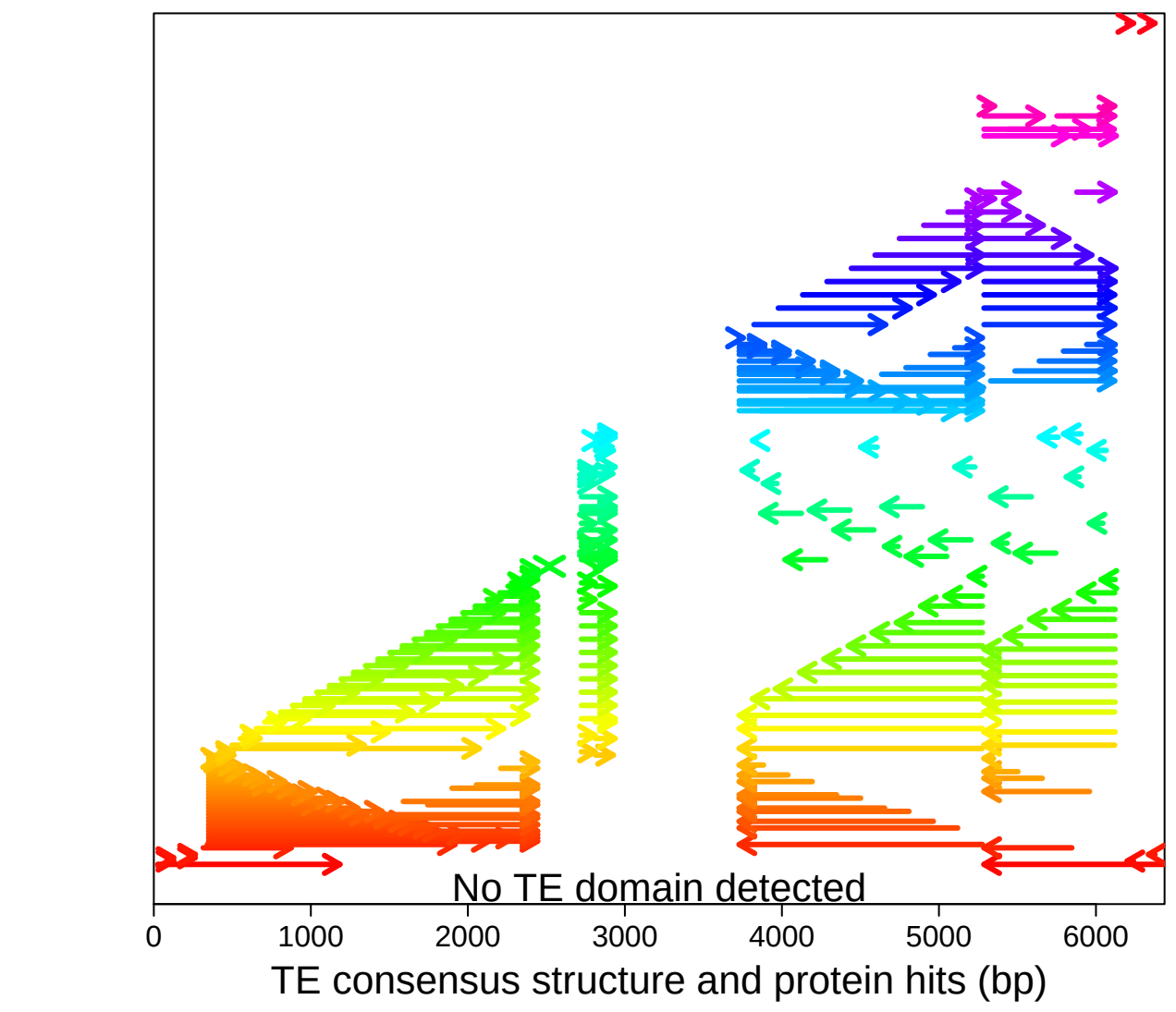
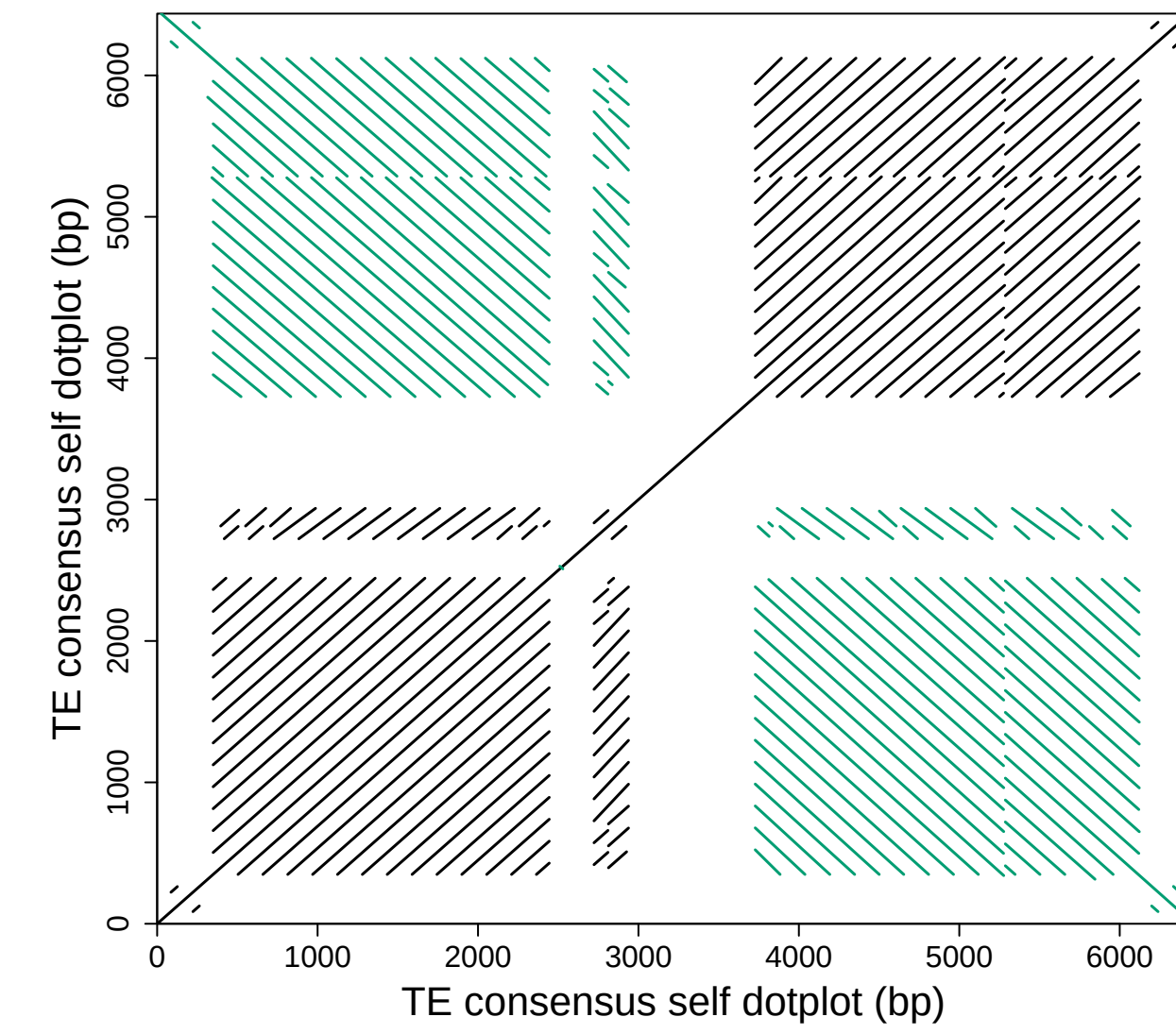
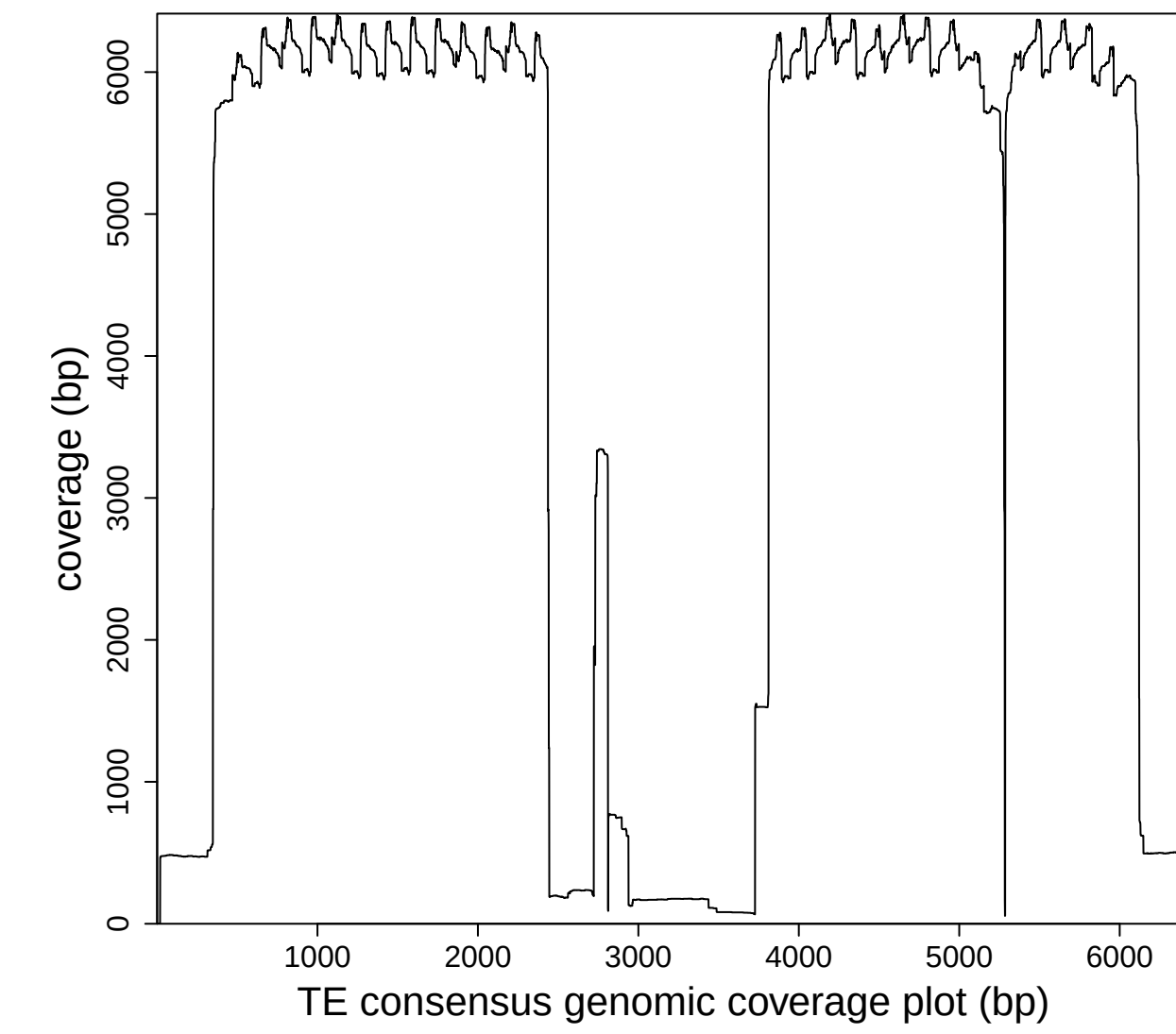
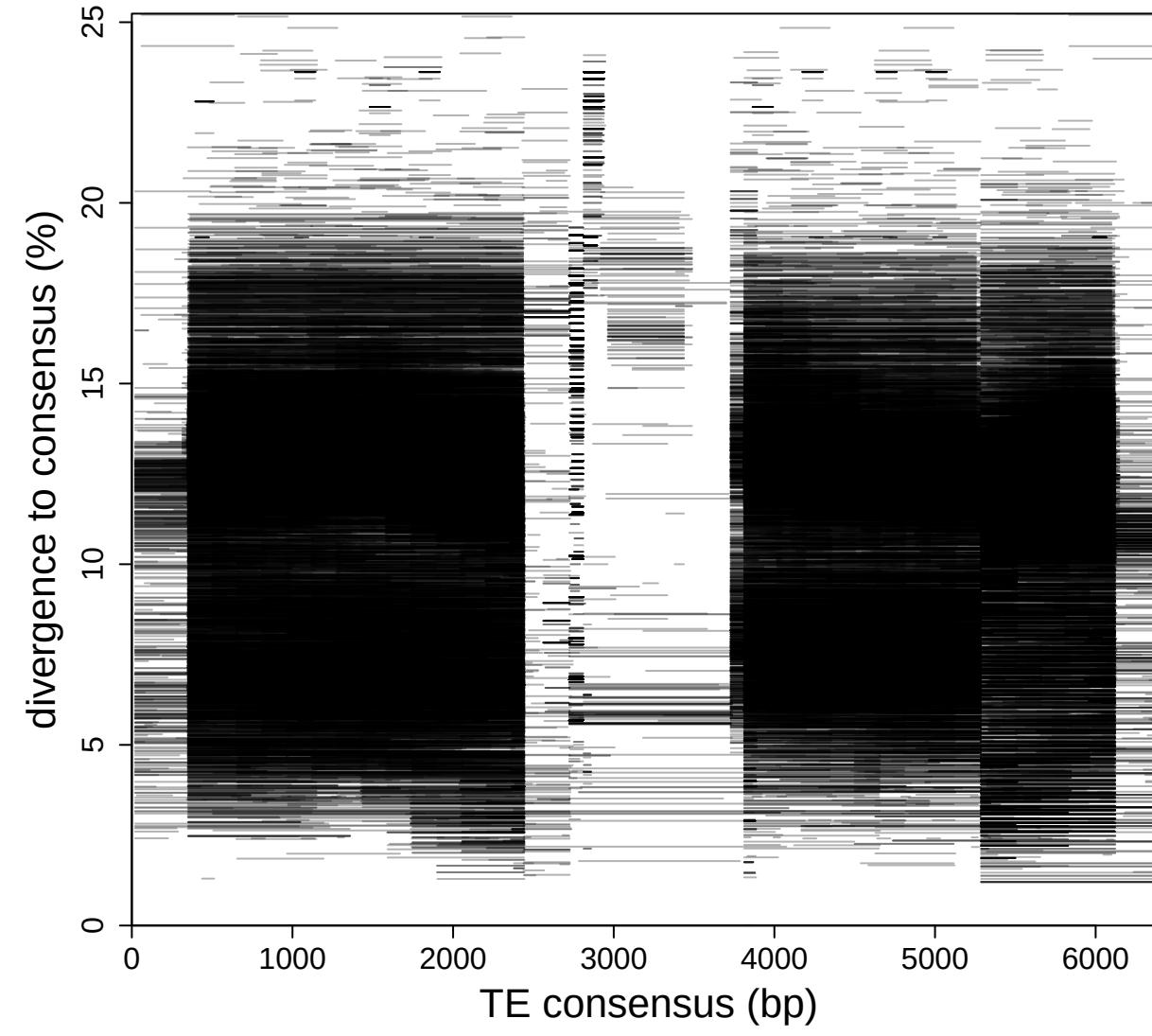
Start crop Point

End crop Point



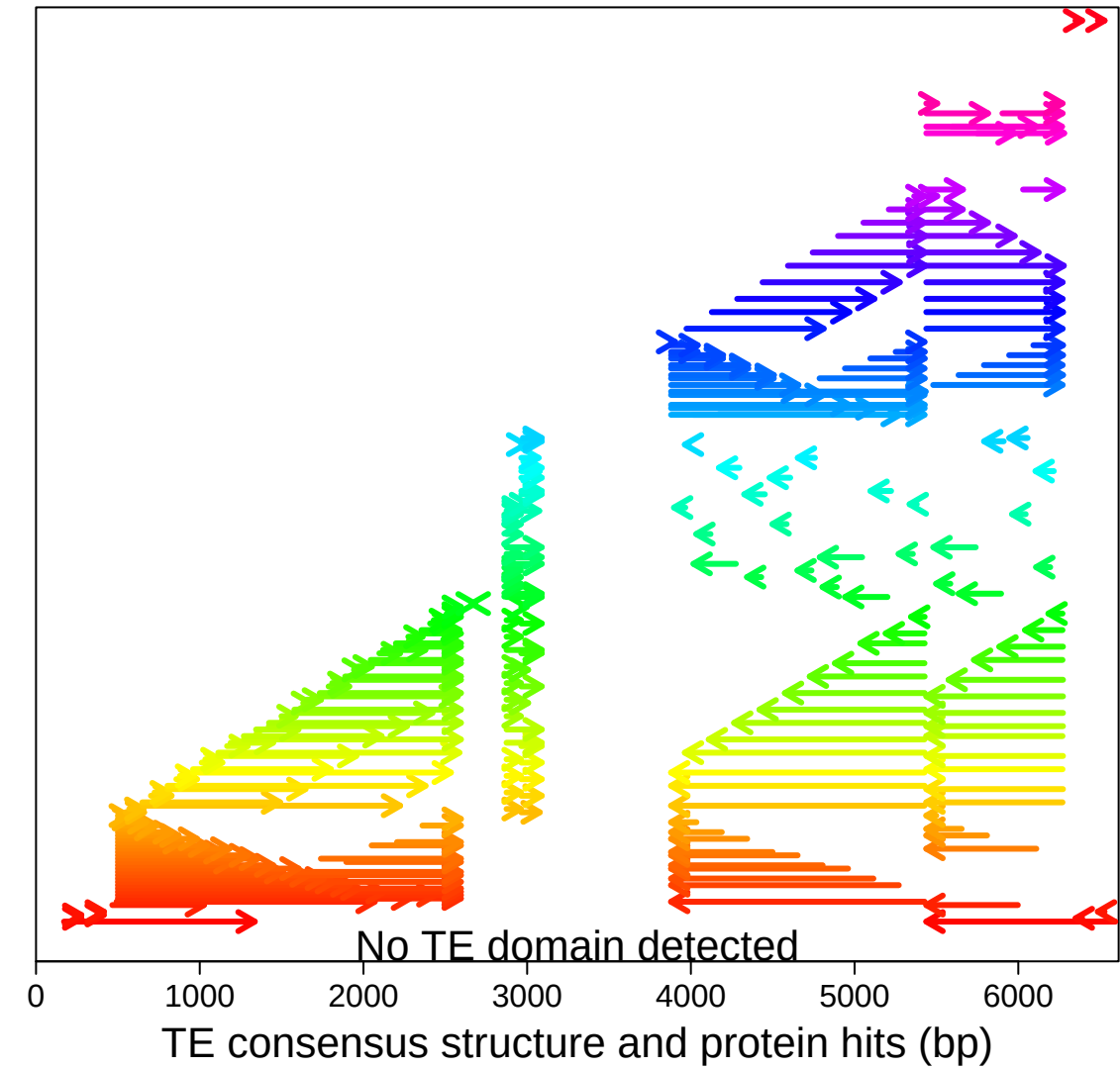
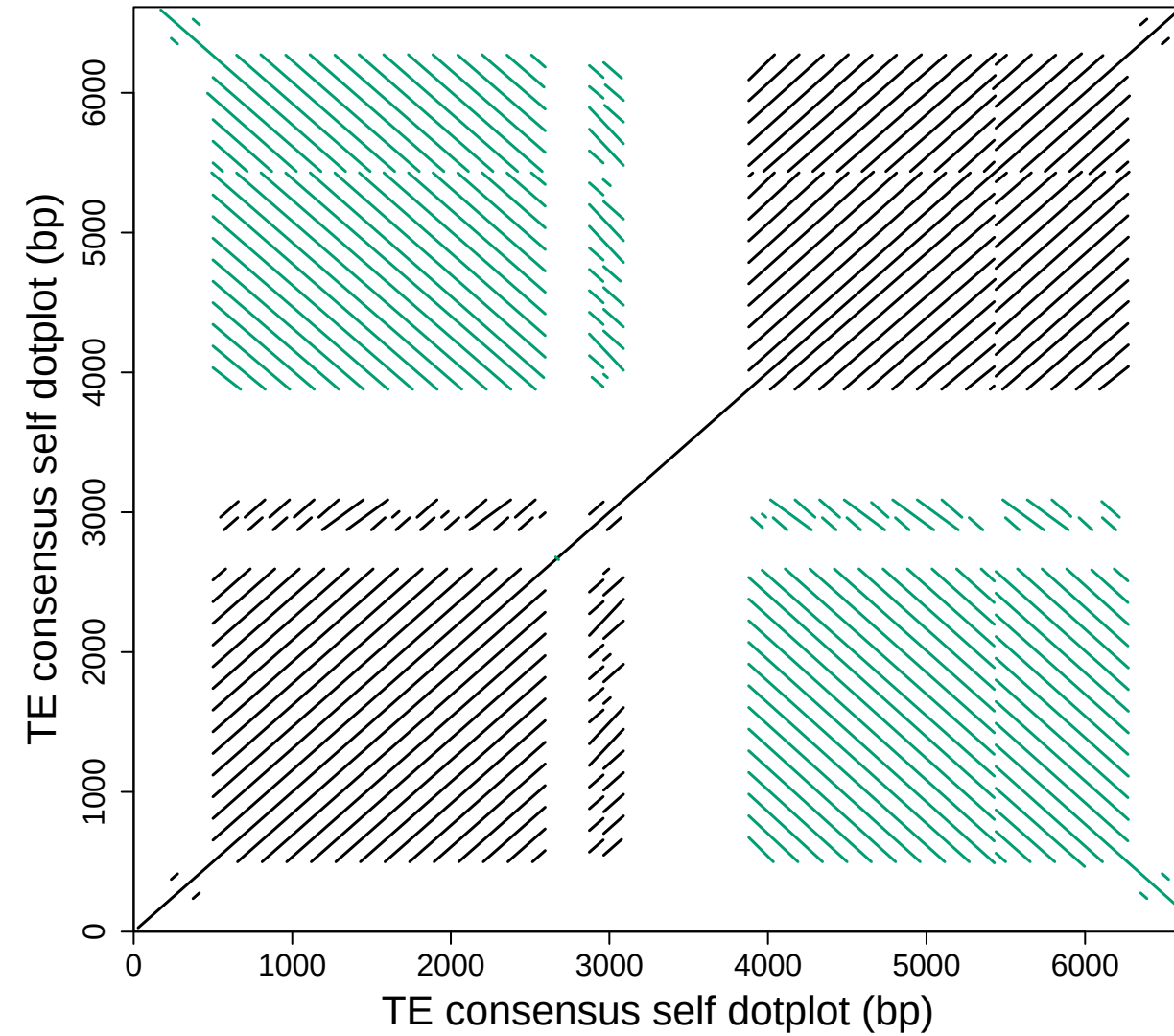
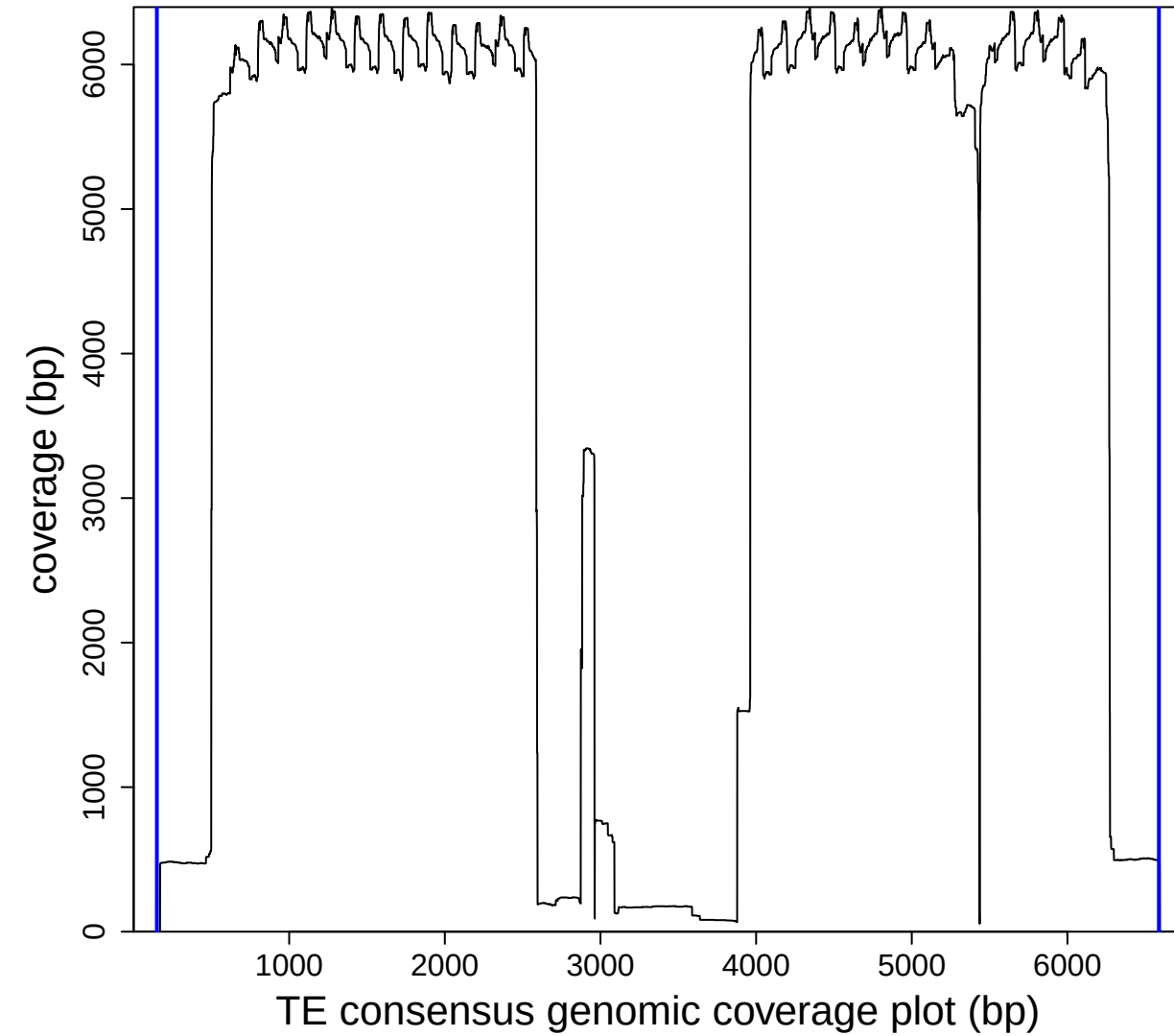
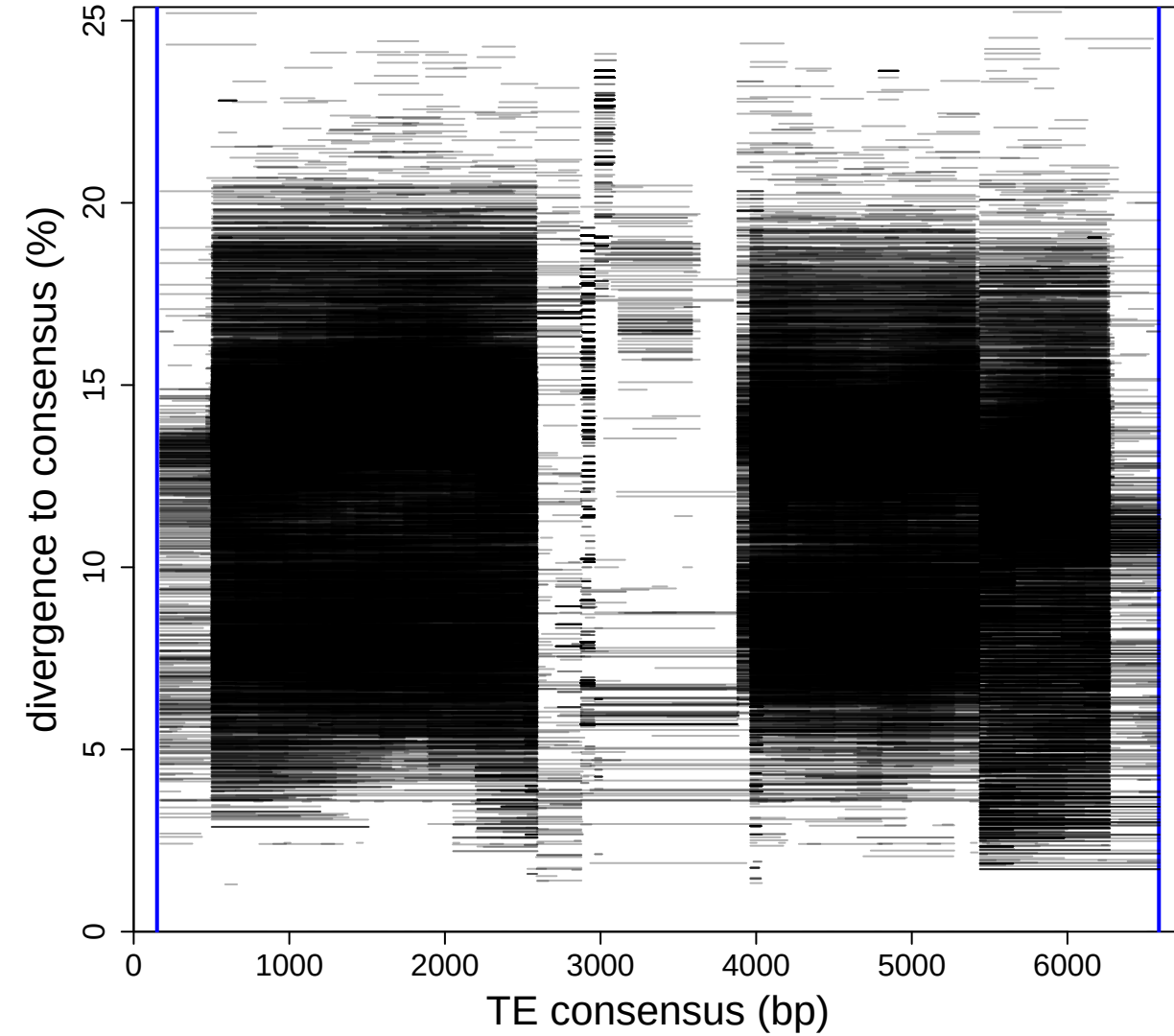
_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcln.fa al
size: 6437bp; fragments: 36832; full length: 0 (>=5793.3bp)

After TEtrimmer 6437 bp

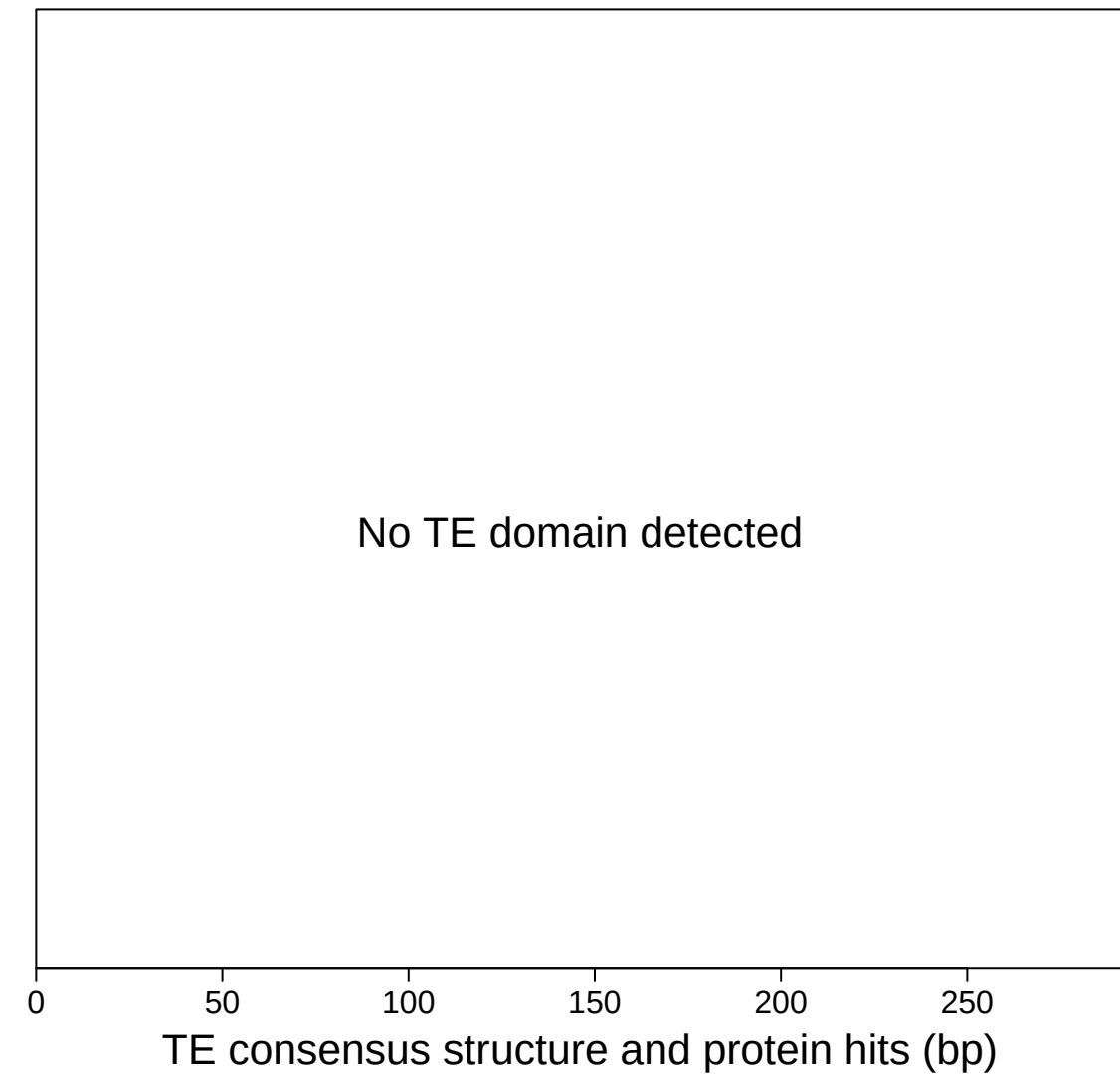
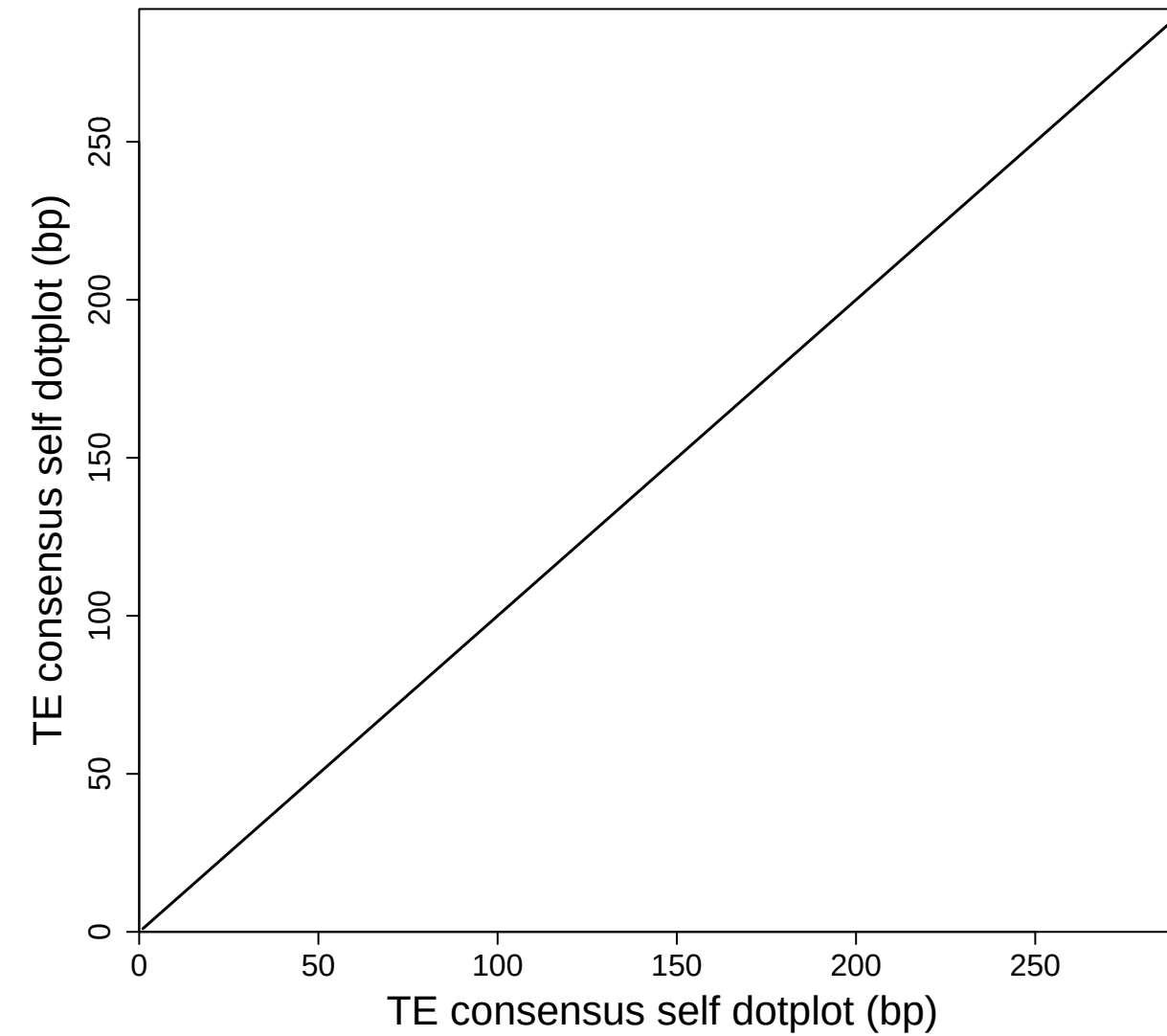
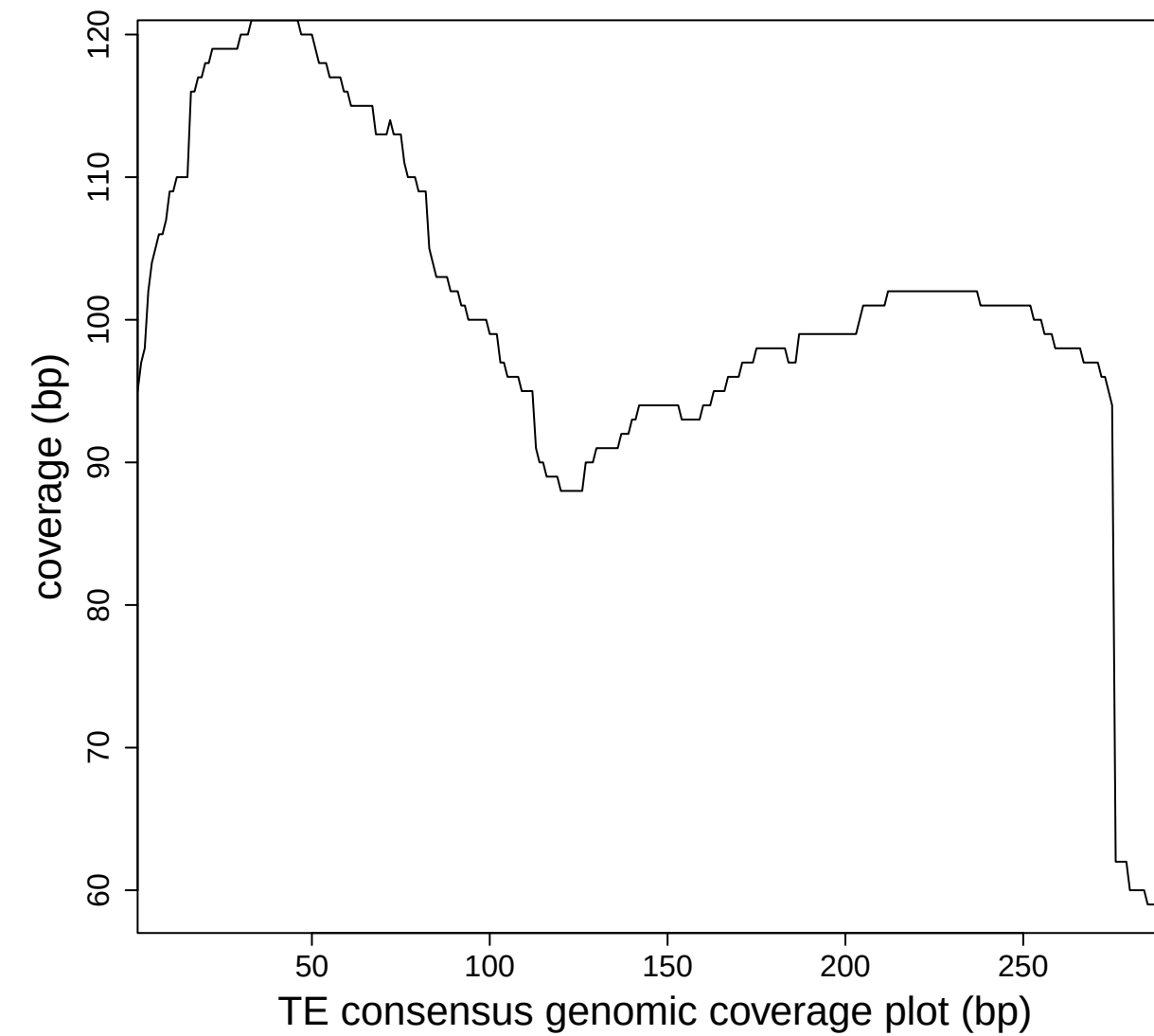
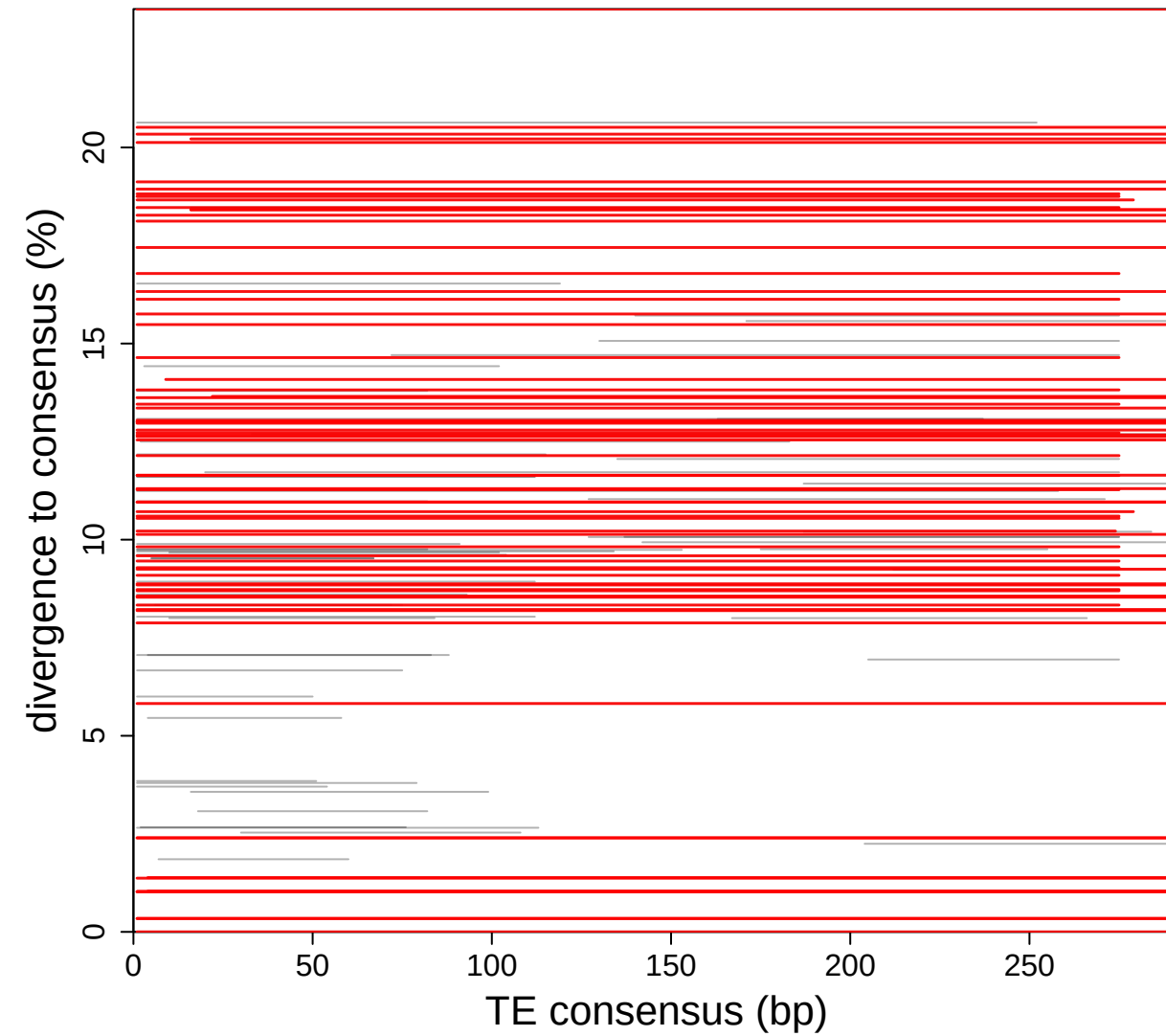


bed_g_1.bed fm_1.bed 0 0 n.bed g_1.bed fm_2.bed 0 0 bclr
size: 6738bp; fragments: 36524; full length: 0 (>=6064.2bp)

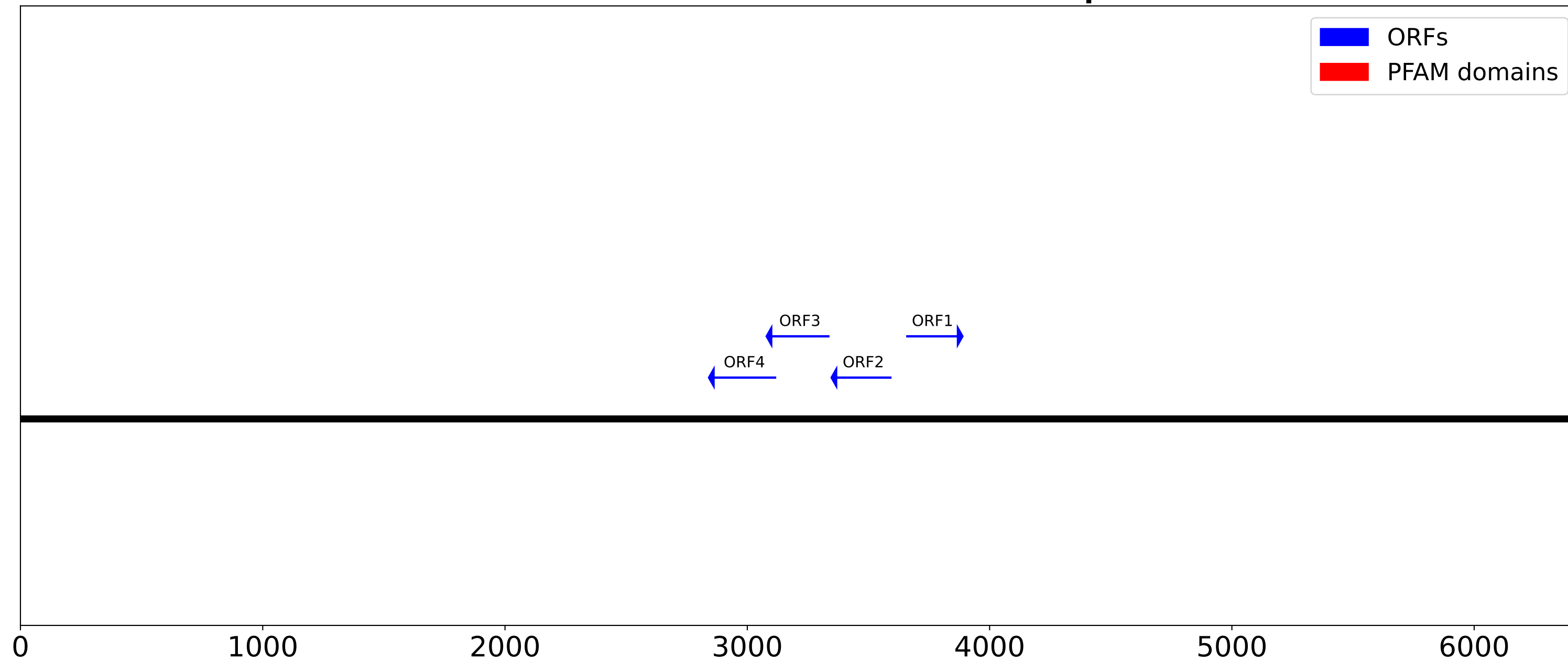
After TEtrimmer Extended plot Blue lines are boundaries



TE: ltr_1_family_217
size: 292bp; fragments: 139; full length: 77 (≥ 262.8 bp)



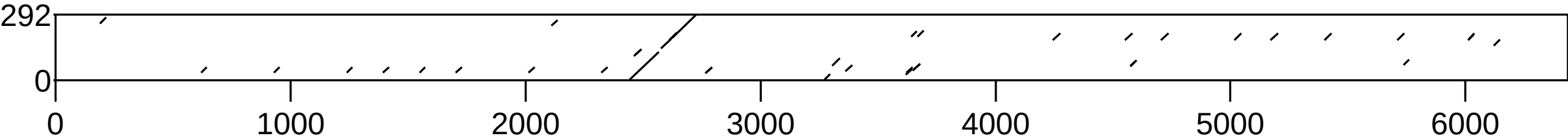
After TEtrimmer ORF and PFAM domain plot



Dotplot

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

nsus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)