

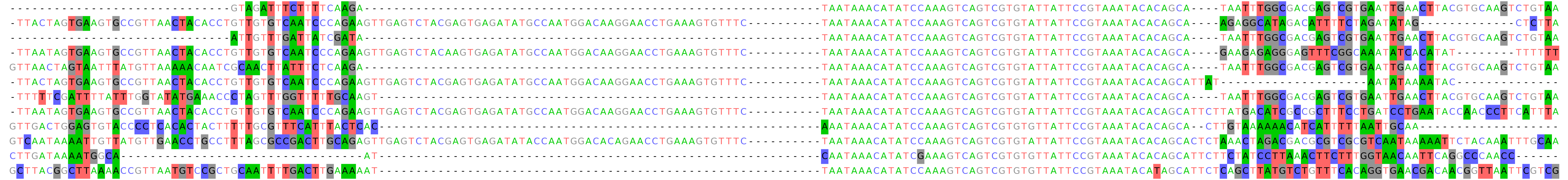
Start crop Point

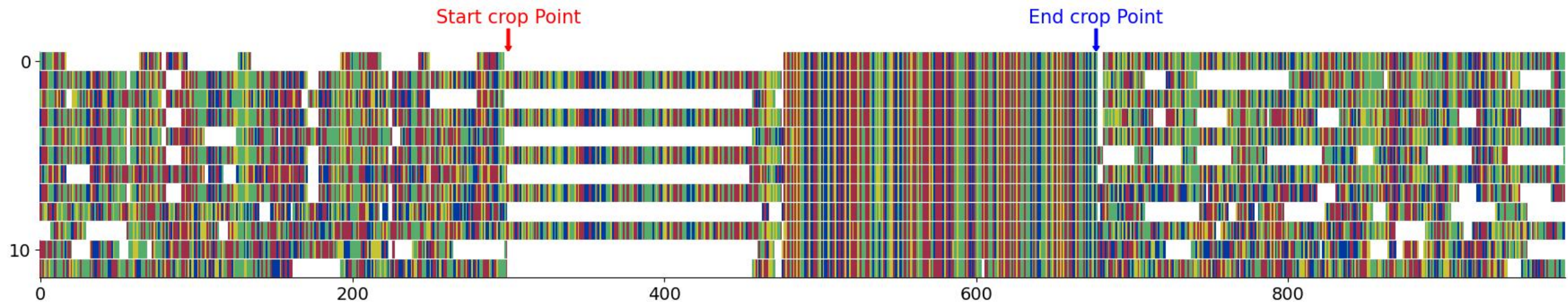


End crop Point



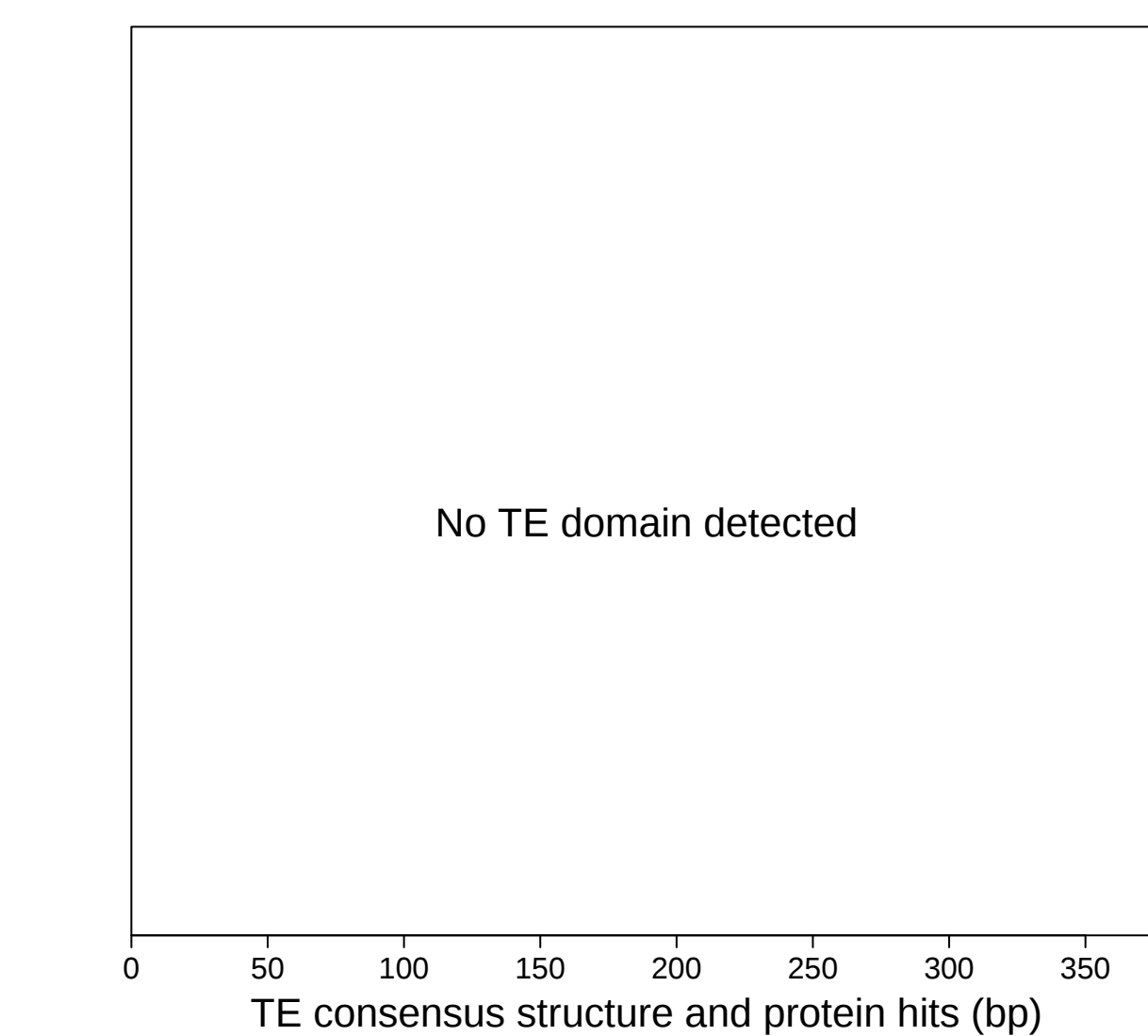
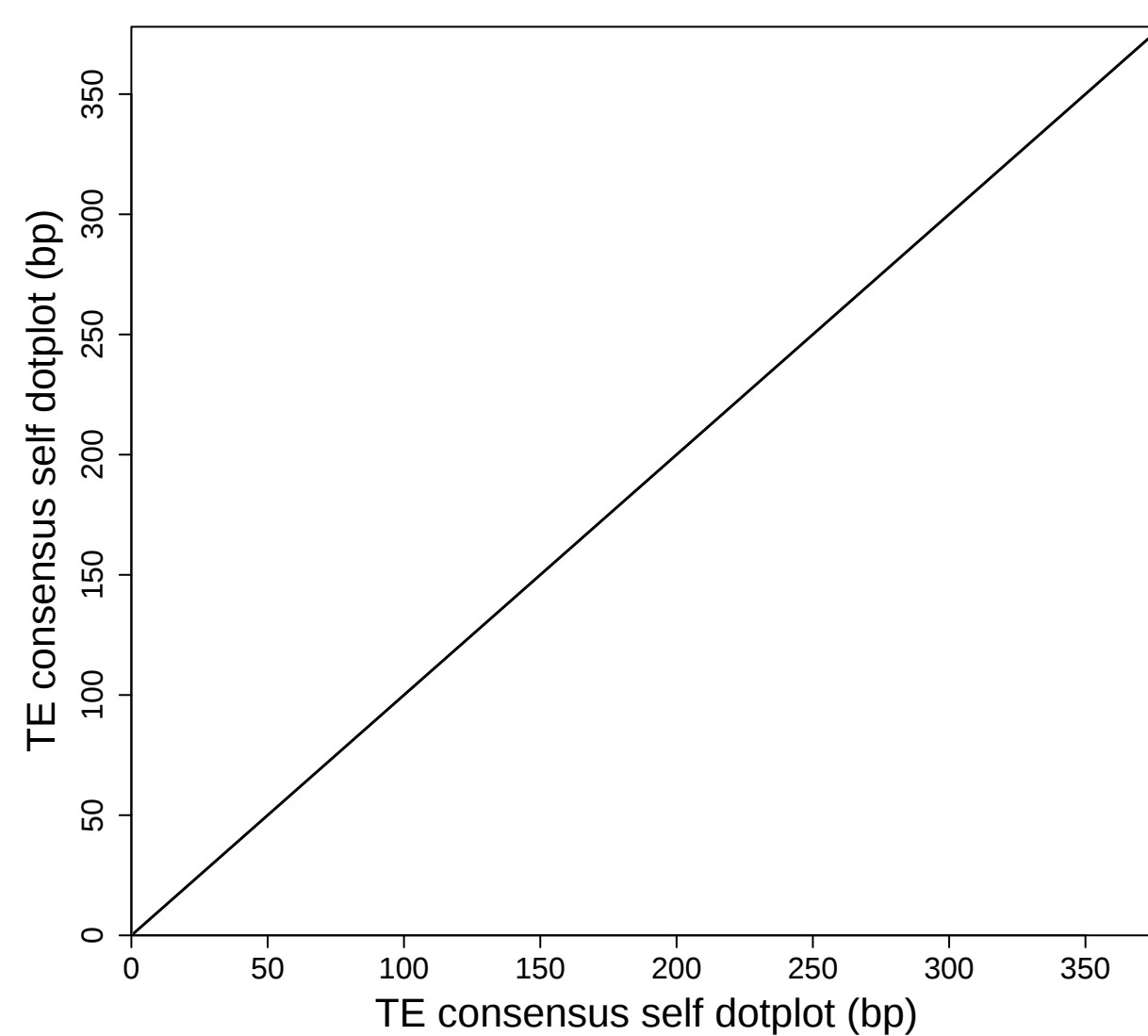
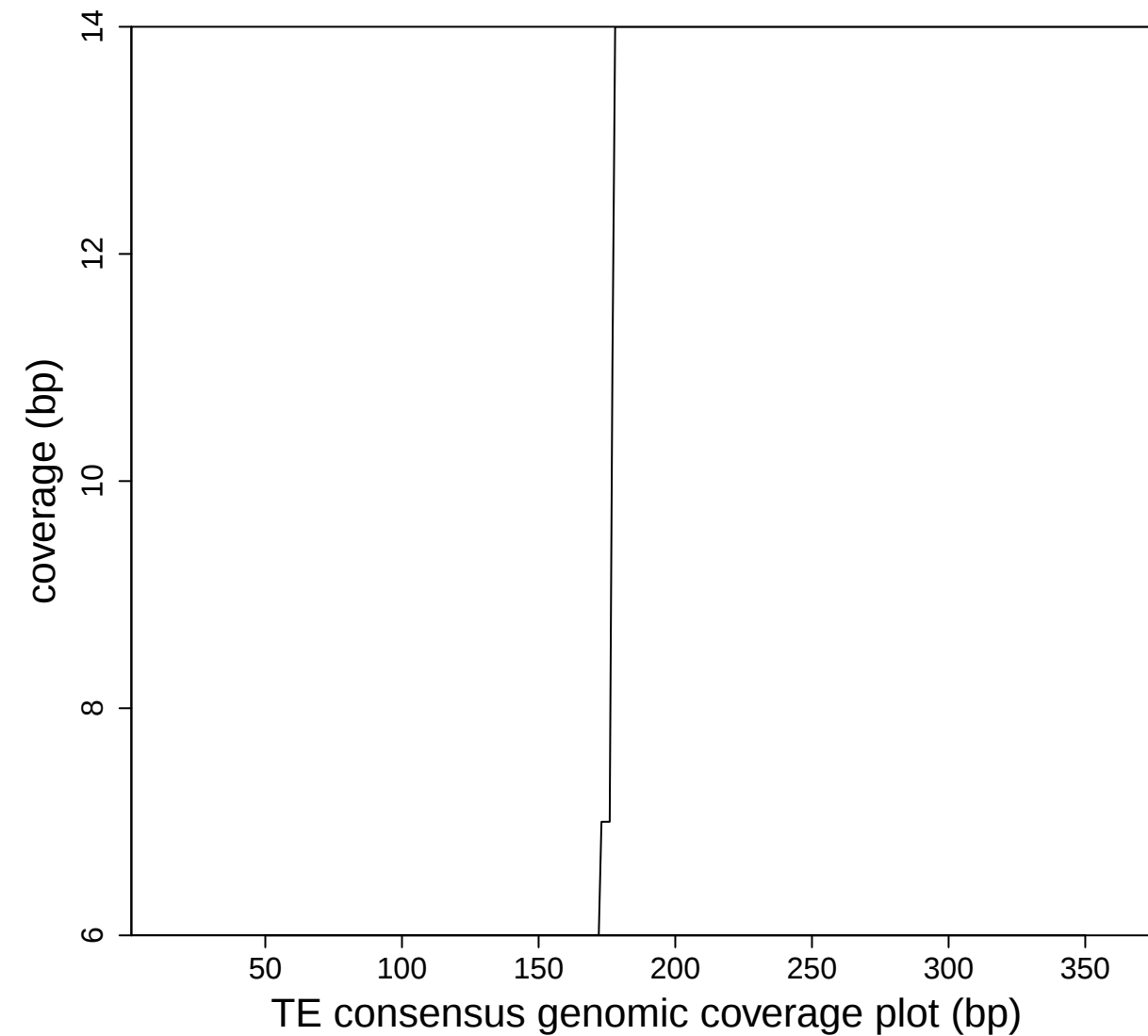
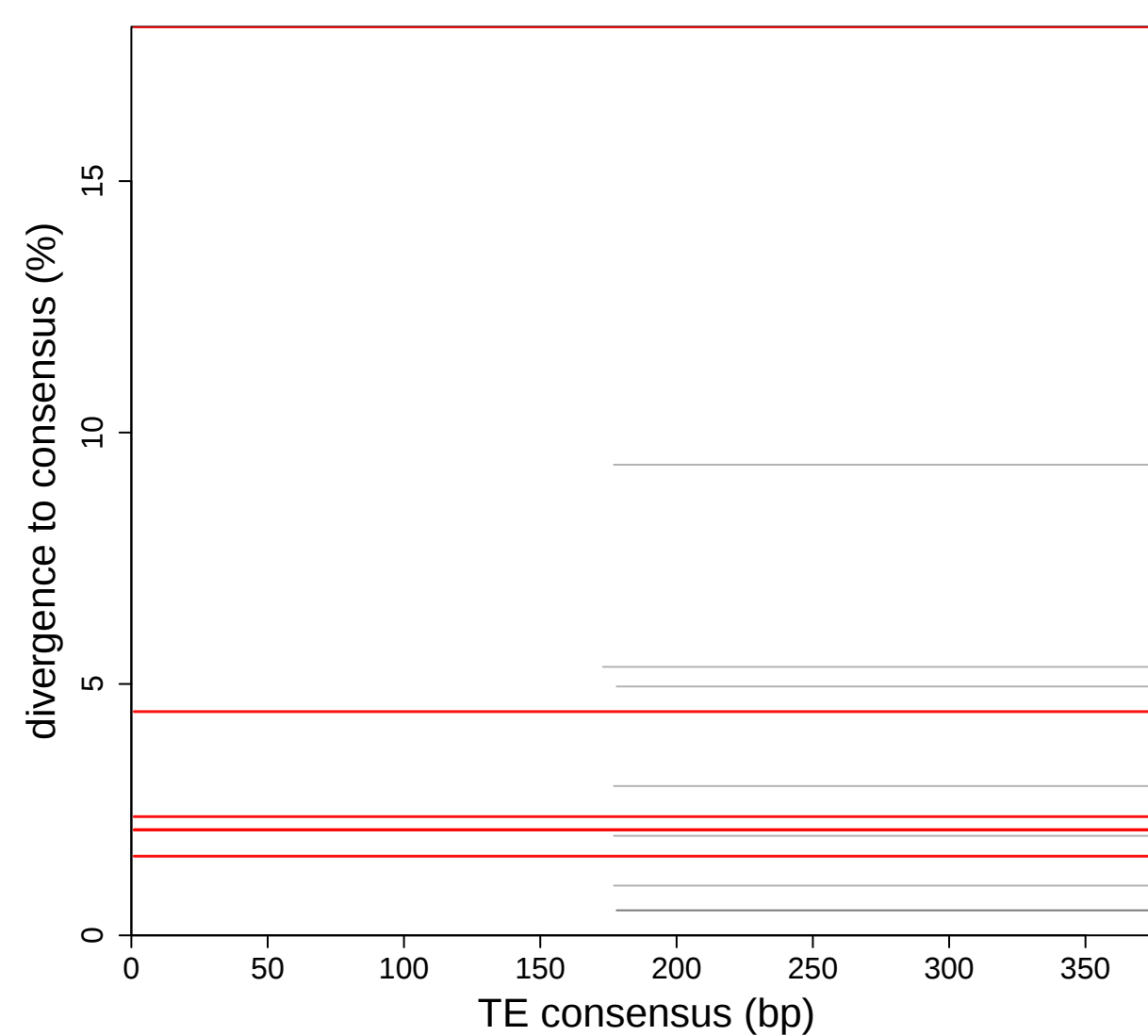
MSA length = 378





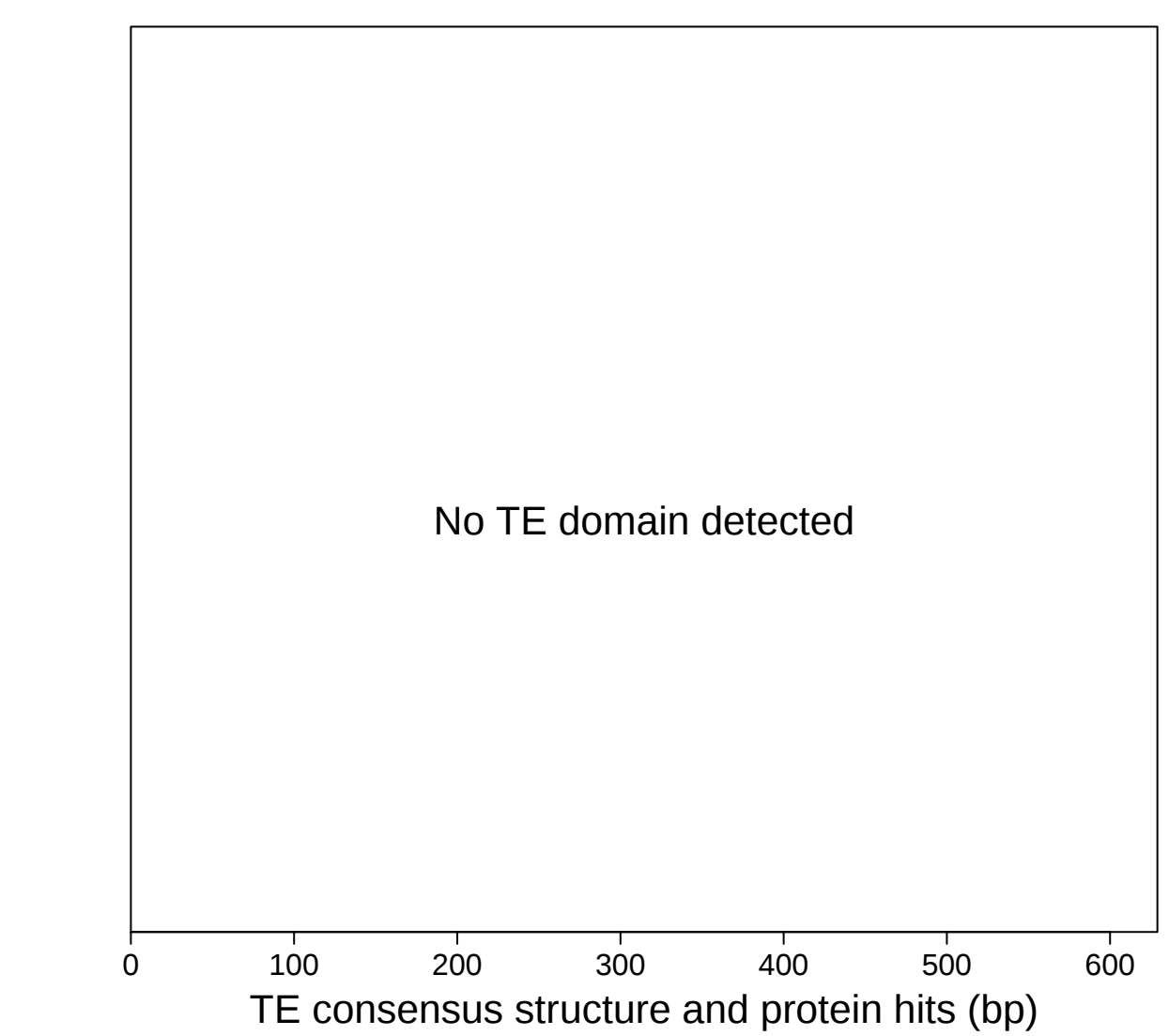
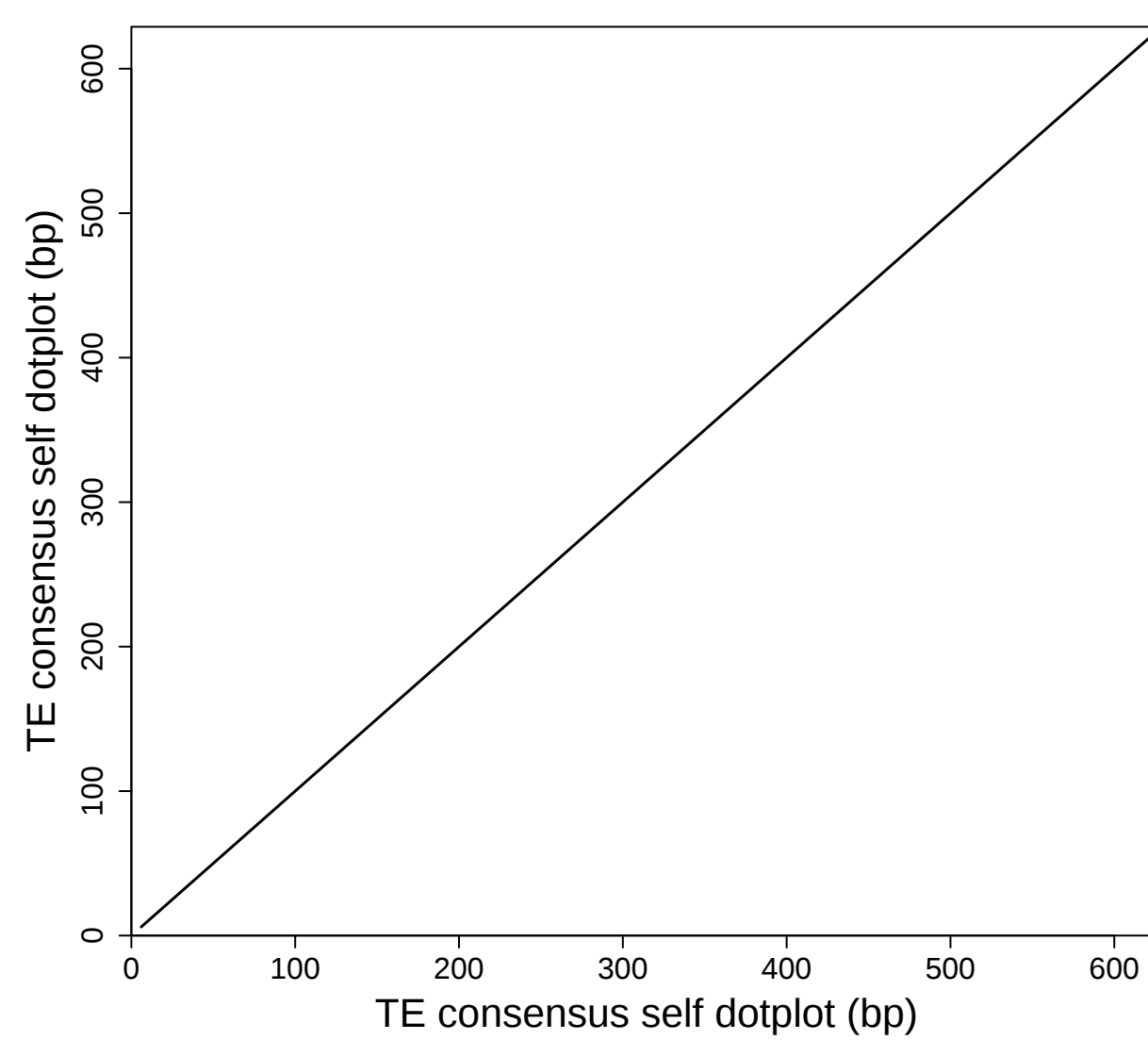
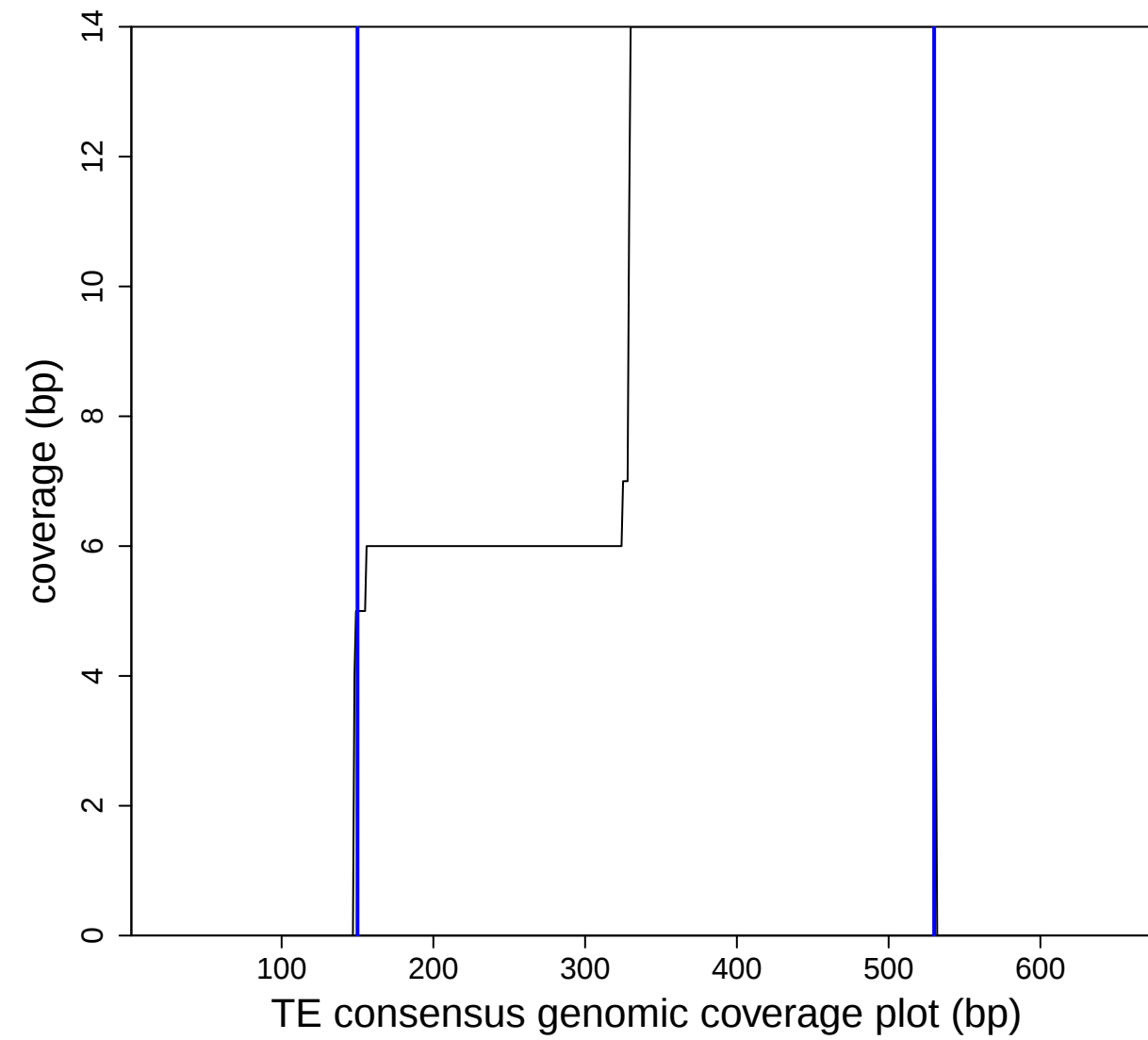
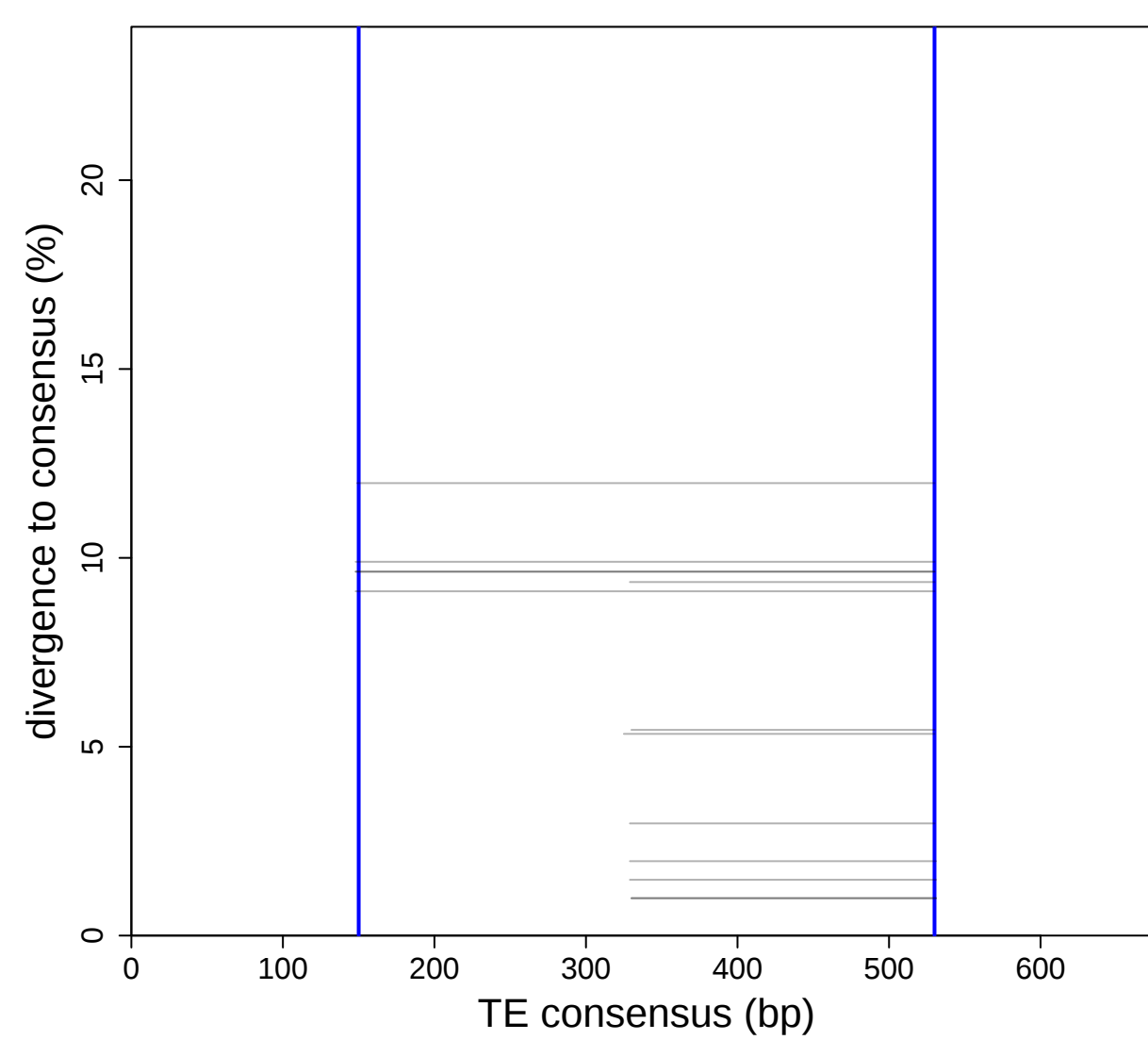
_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcln.fa_al
size: 378bp; fragments: 14; full length: 6 (>=340.2bp)

After TEtrimmer 378 bp

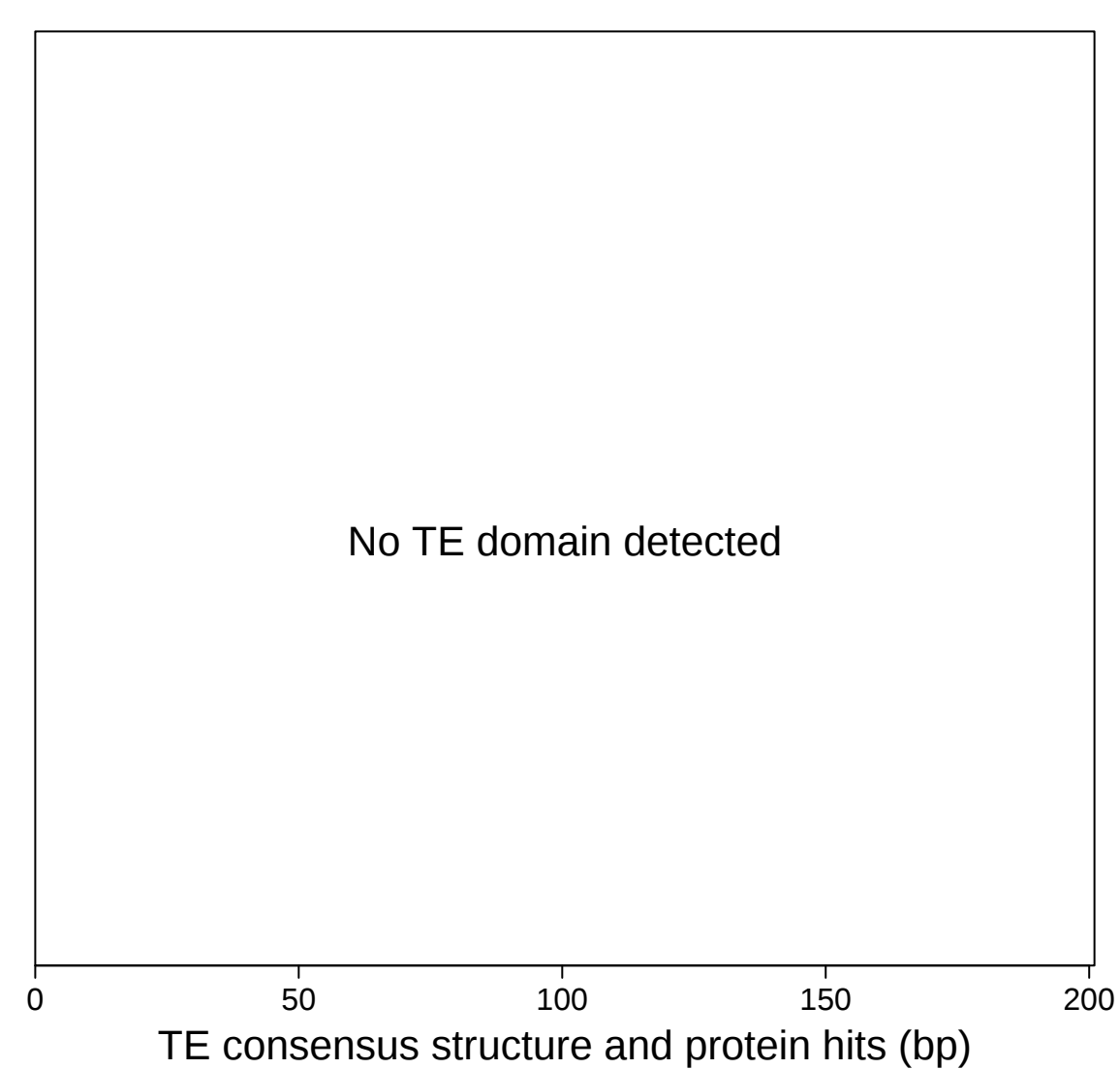
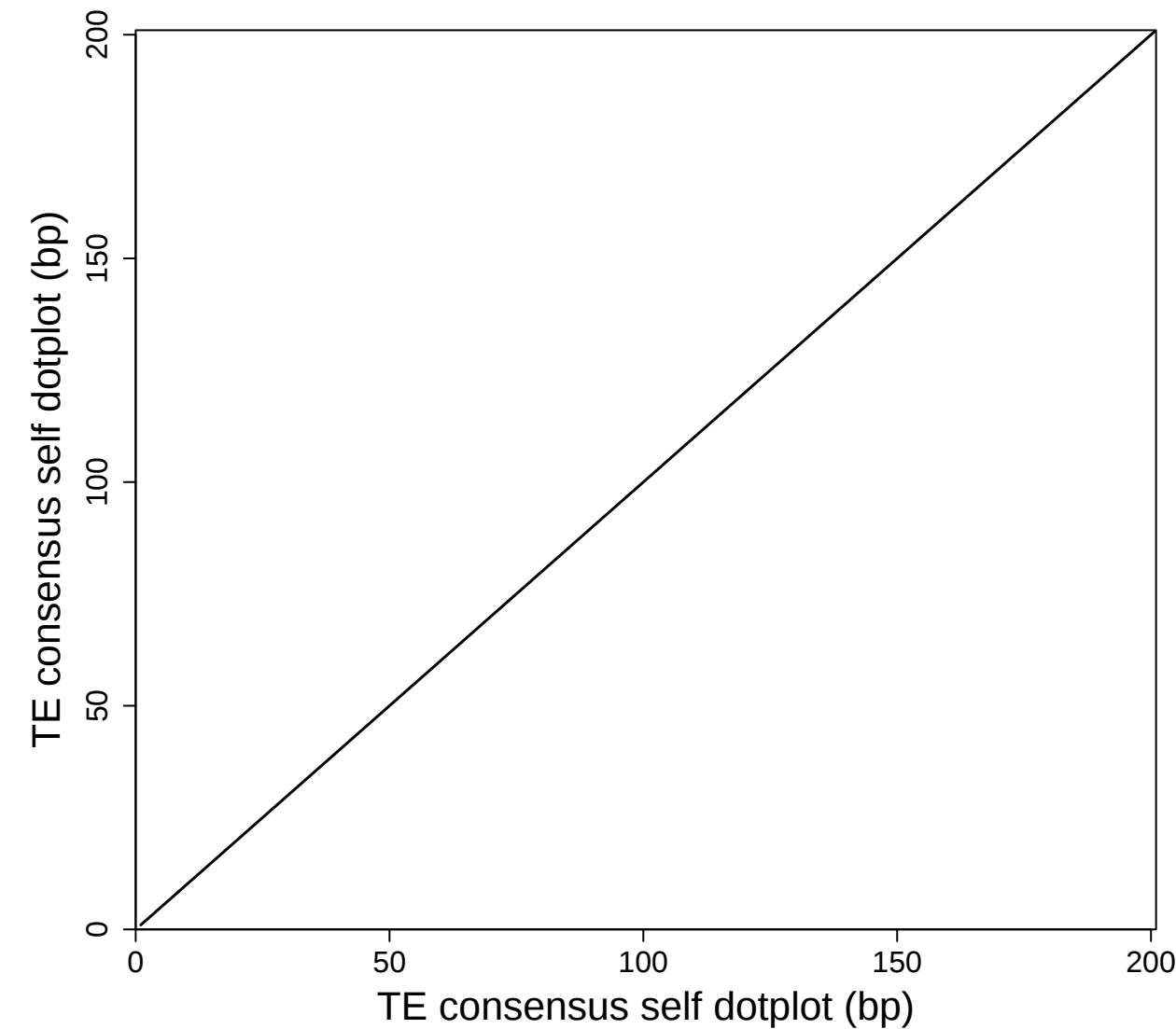
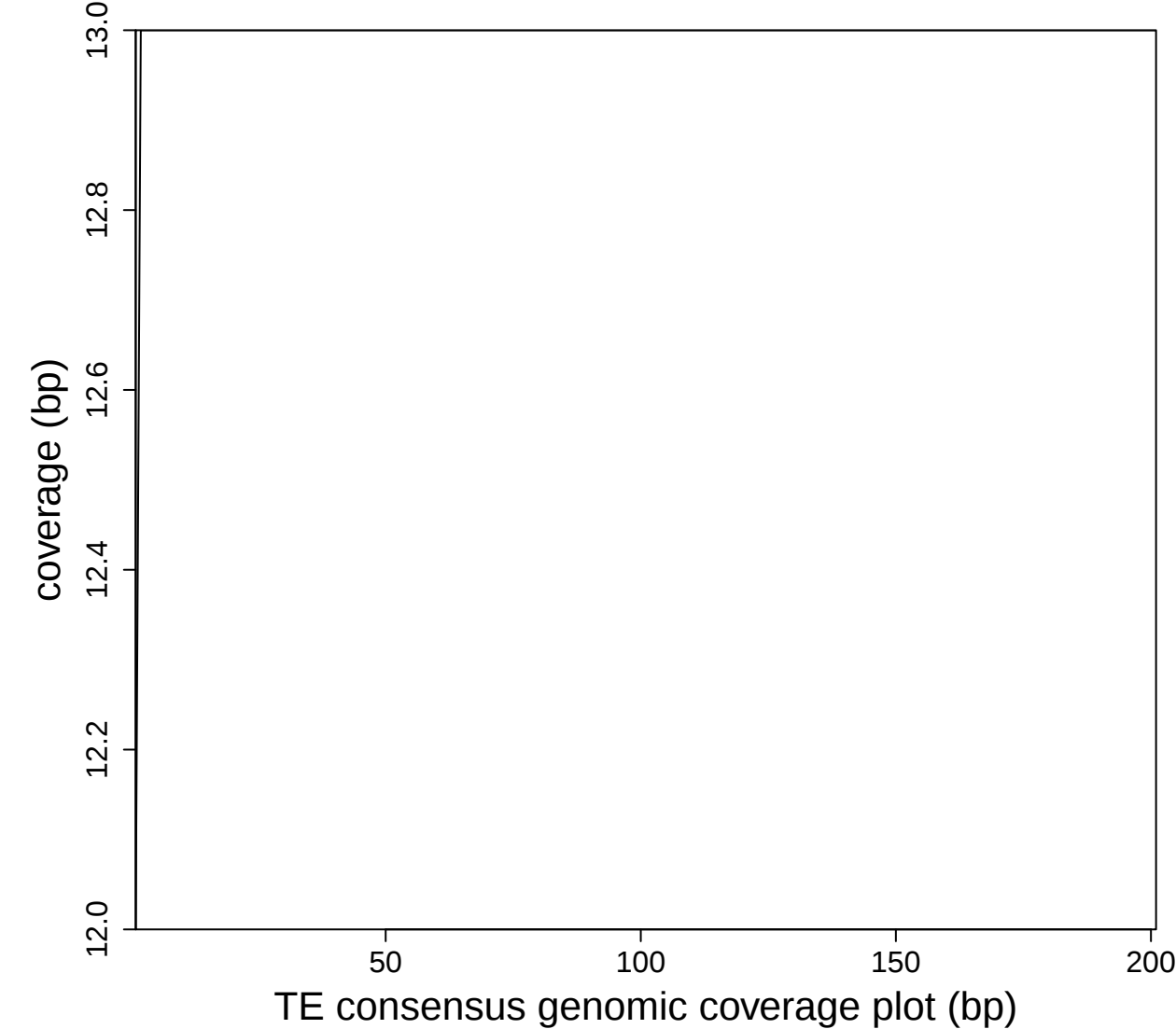
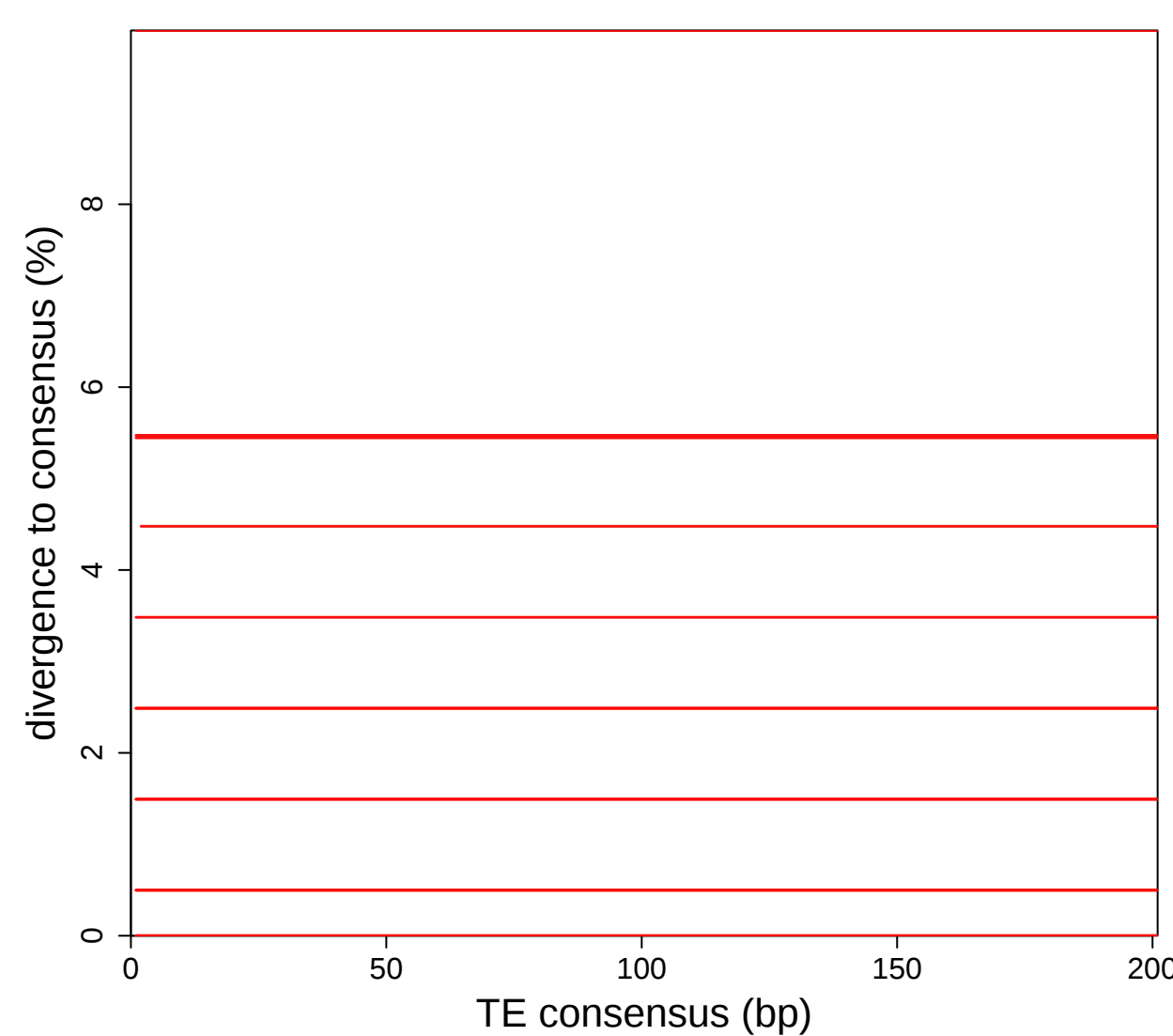


bed_g_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bclr
size: 680bp; fragments: 14; full length: 0 (>=612bp)

After TEtrimmer Extended plot Blue lines are boundaries



TE: ltr_1_family_366
size: 201bp; fragments: 13; full length: 13 (≥ 180.9 bp)

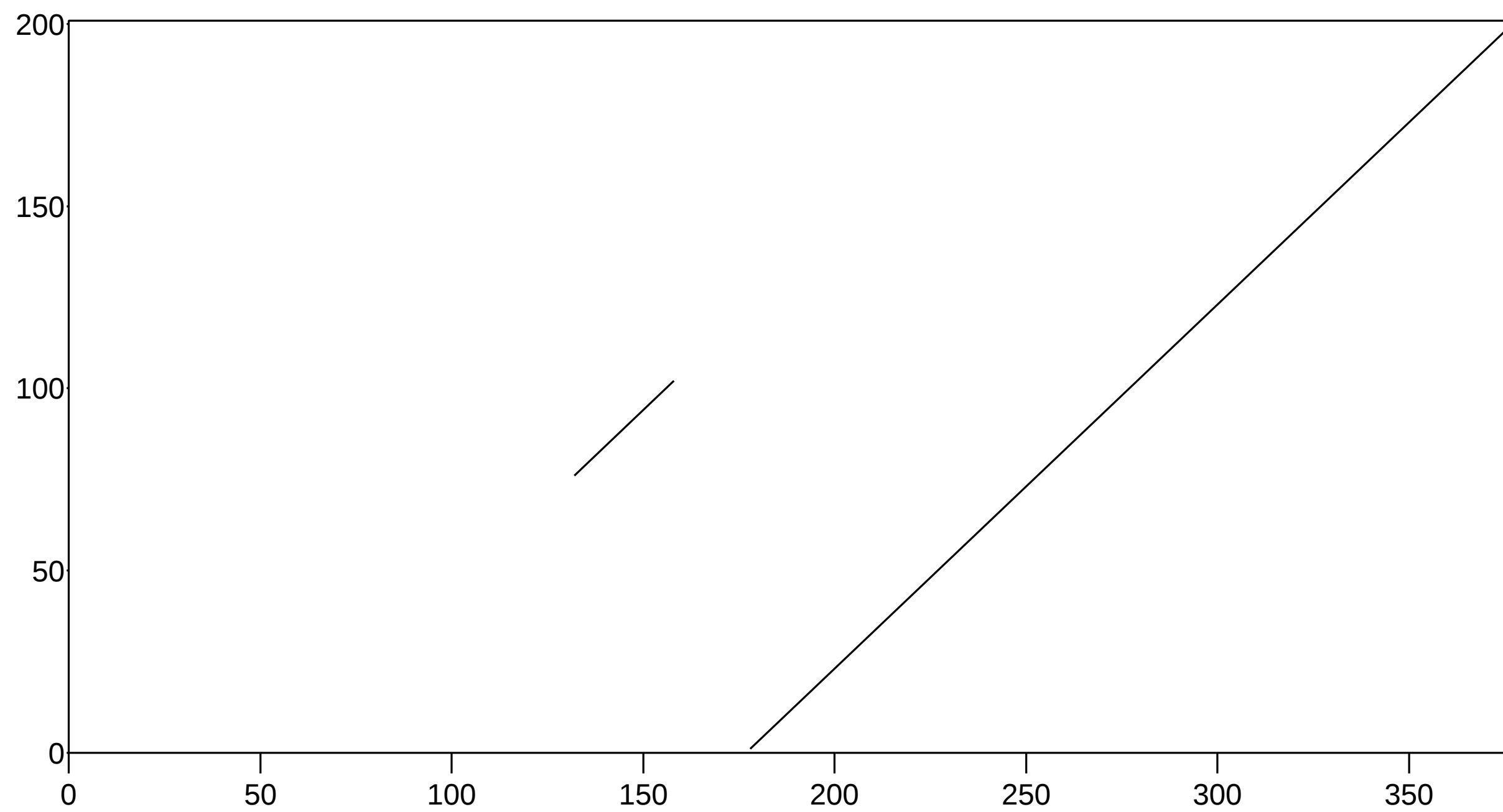


Before TEtrimmer 201 bp

Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)