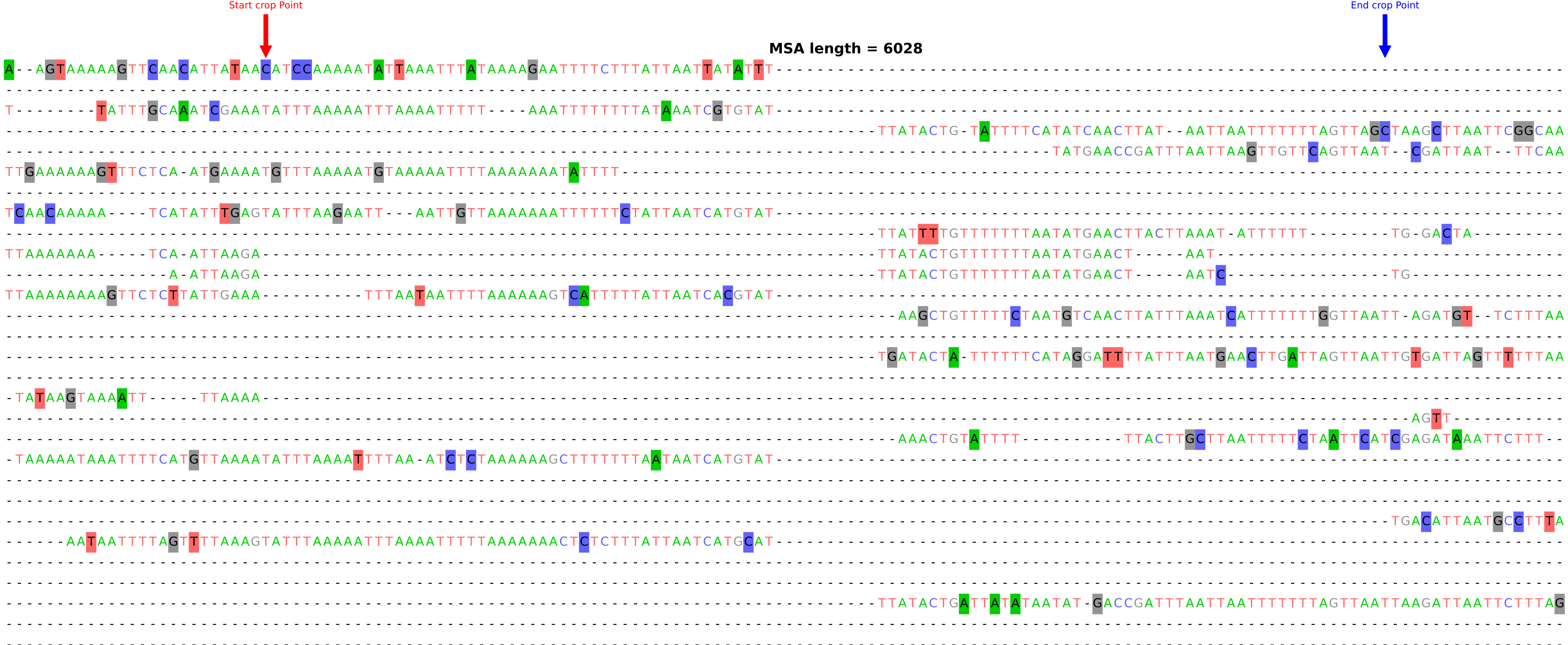
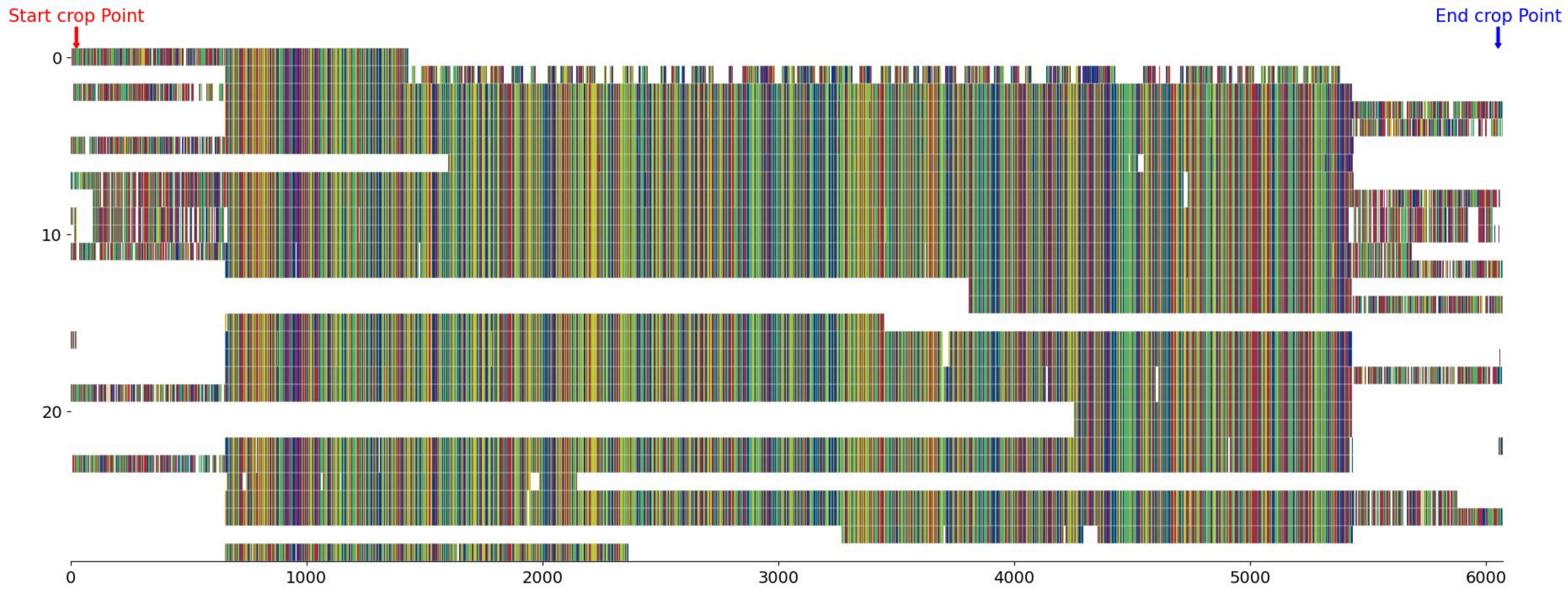
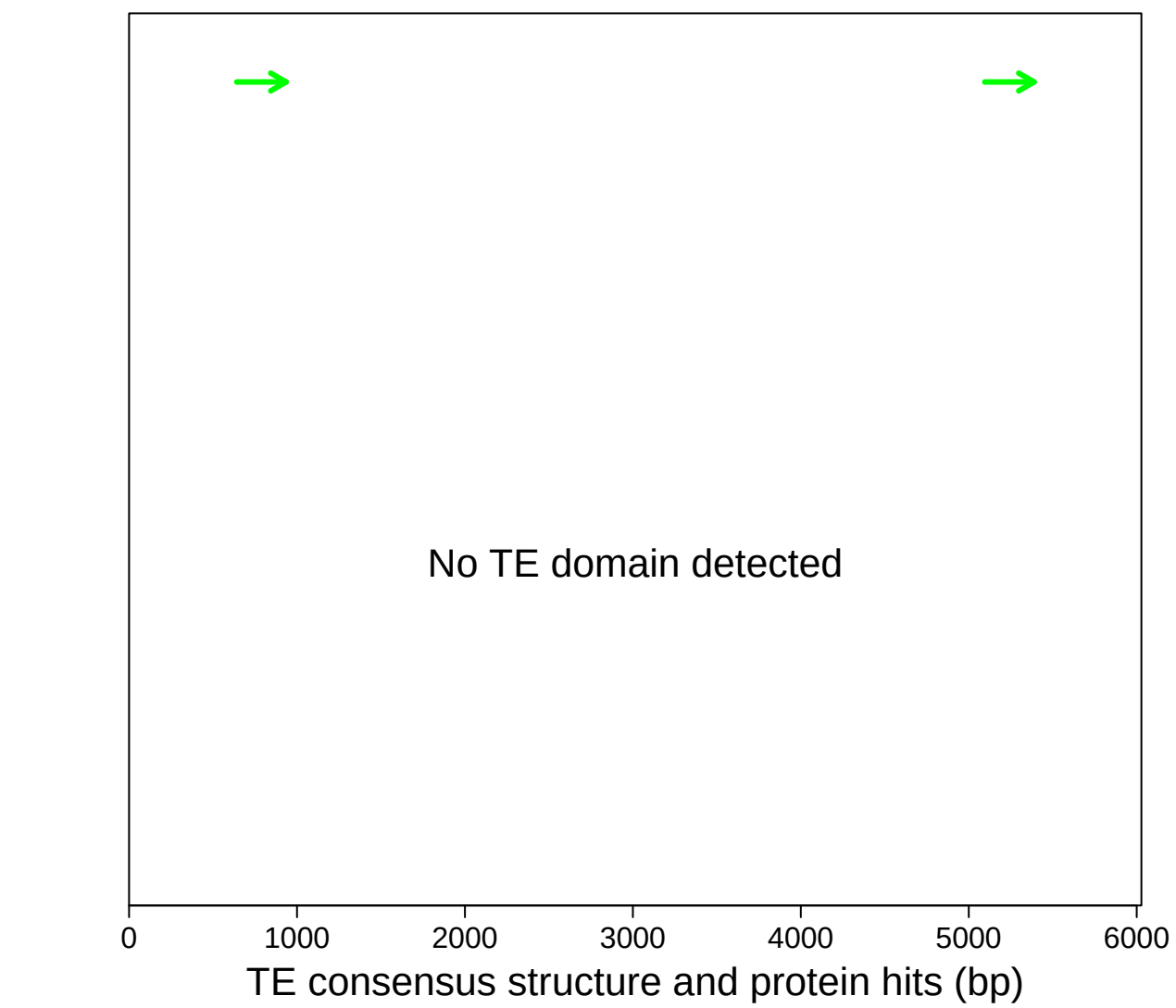
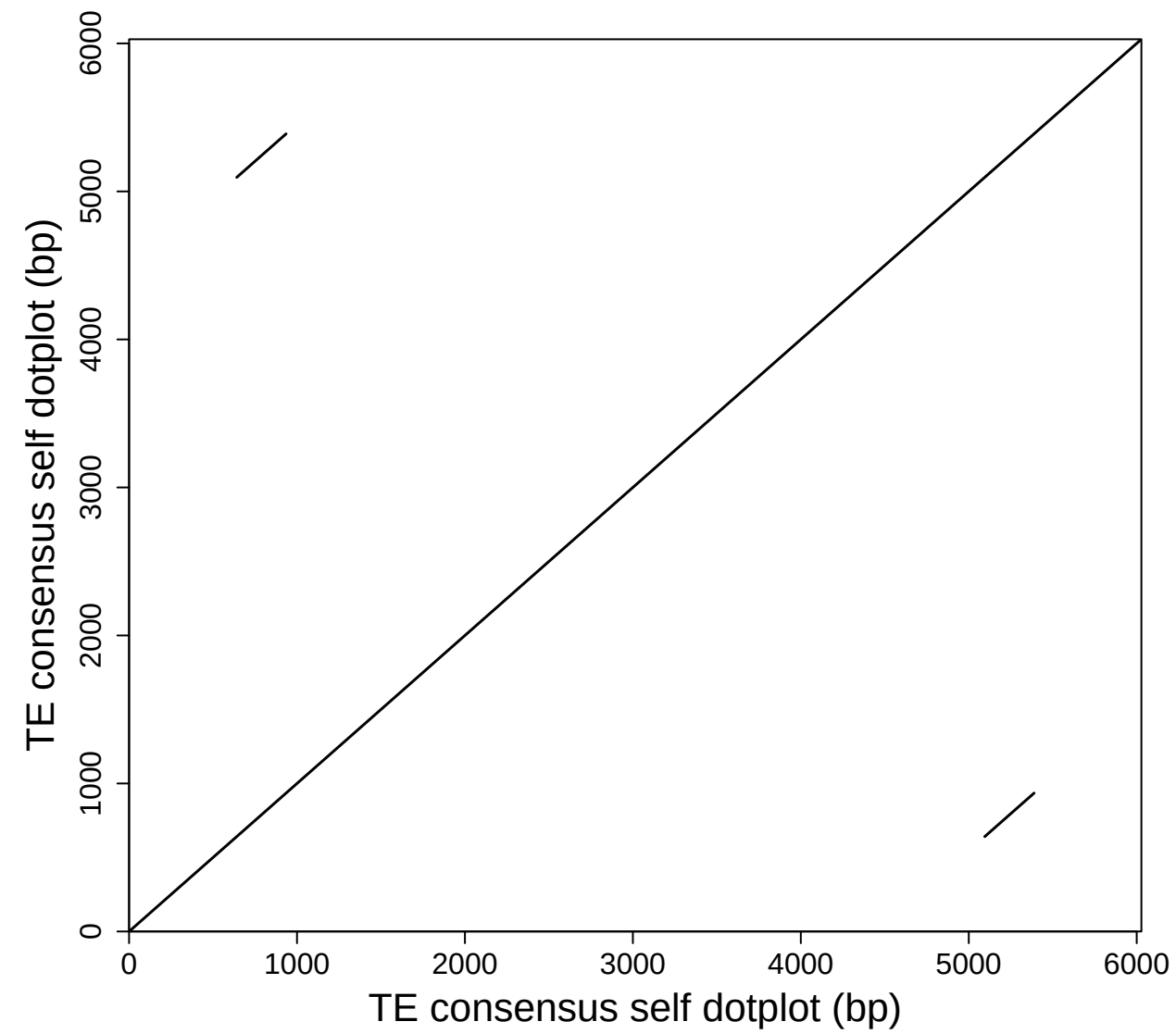
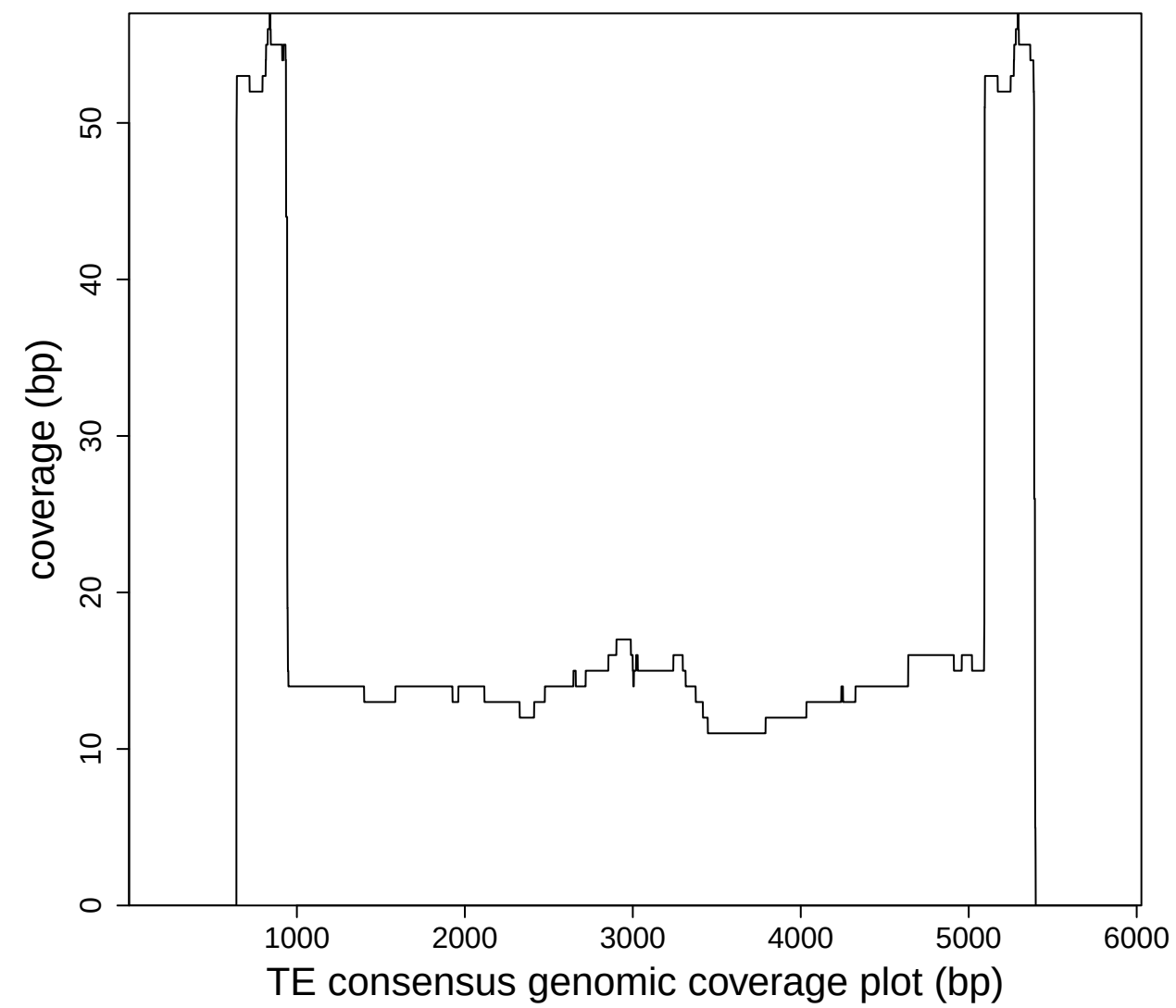
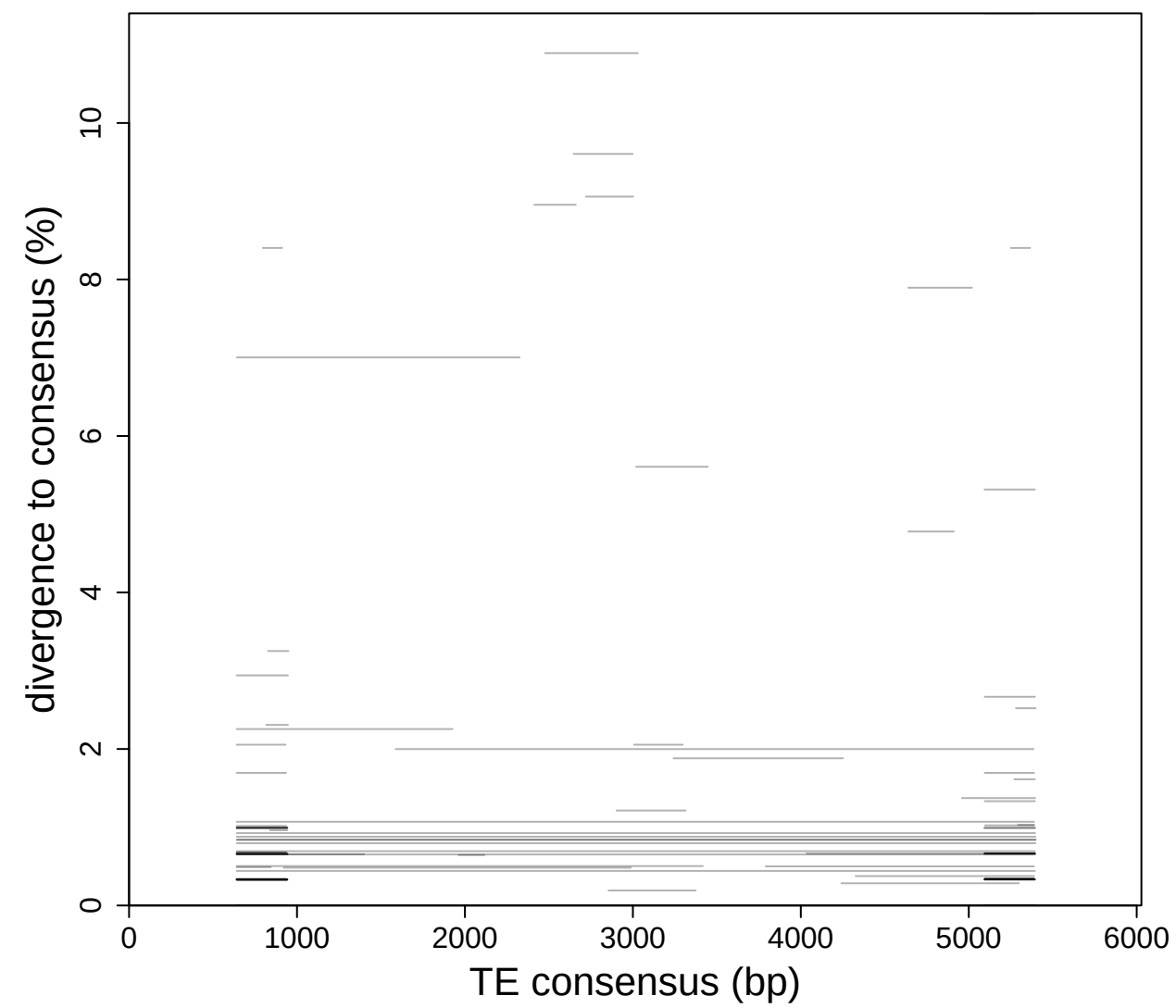


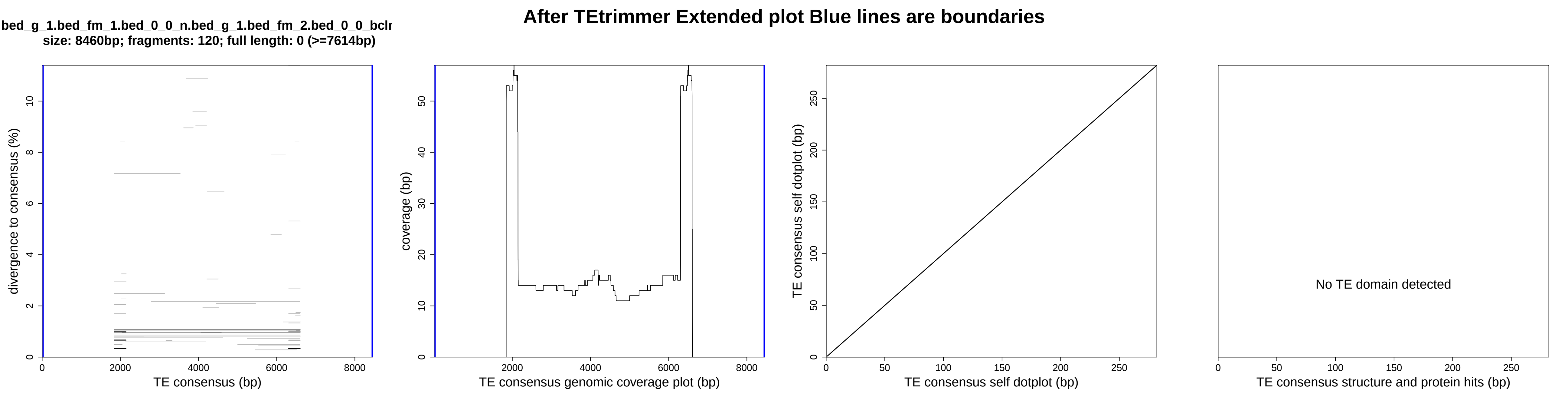




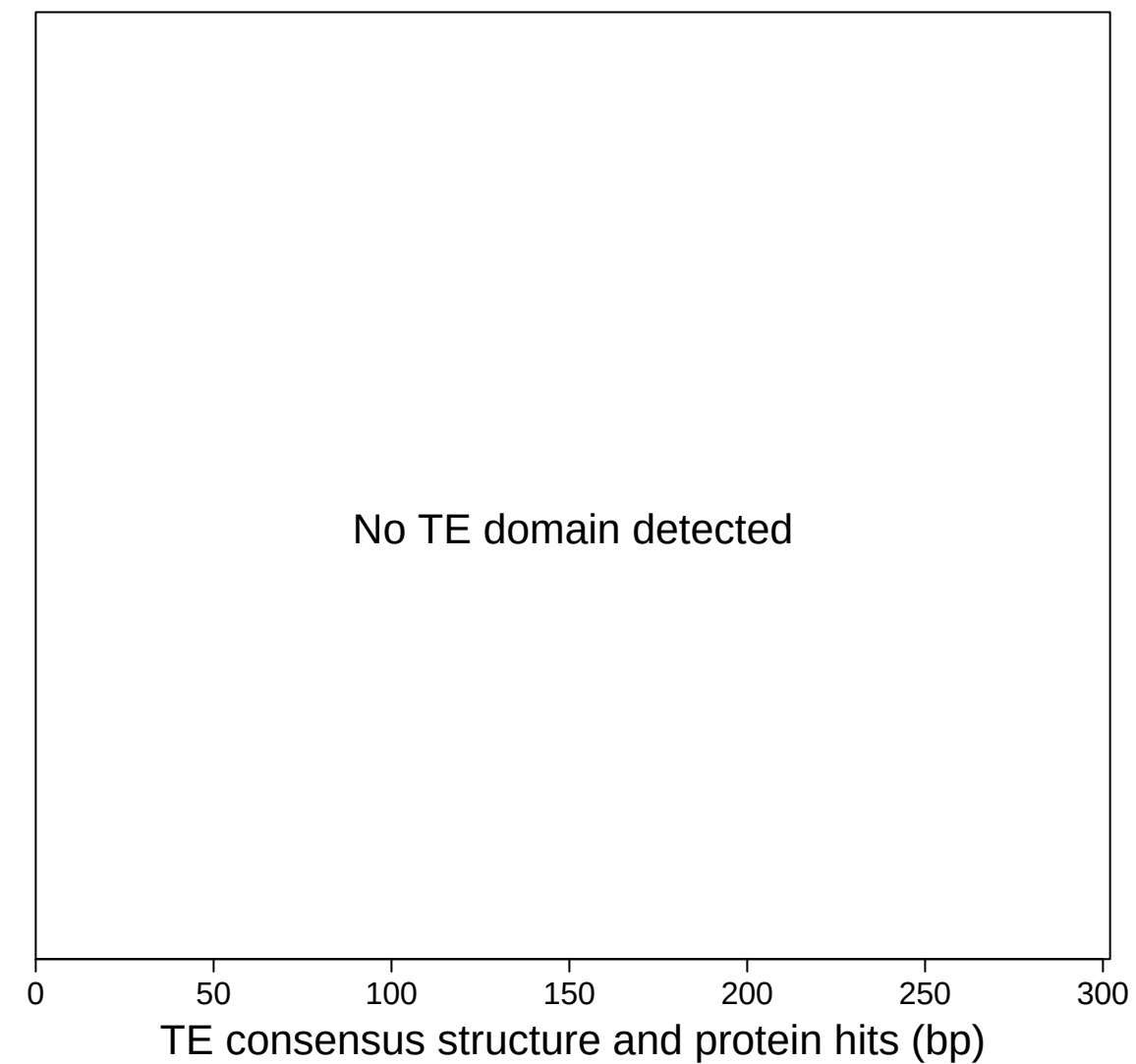
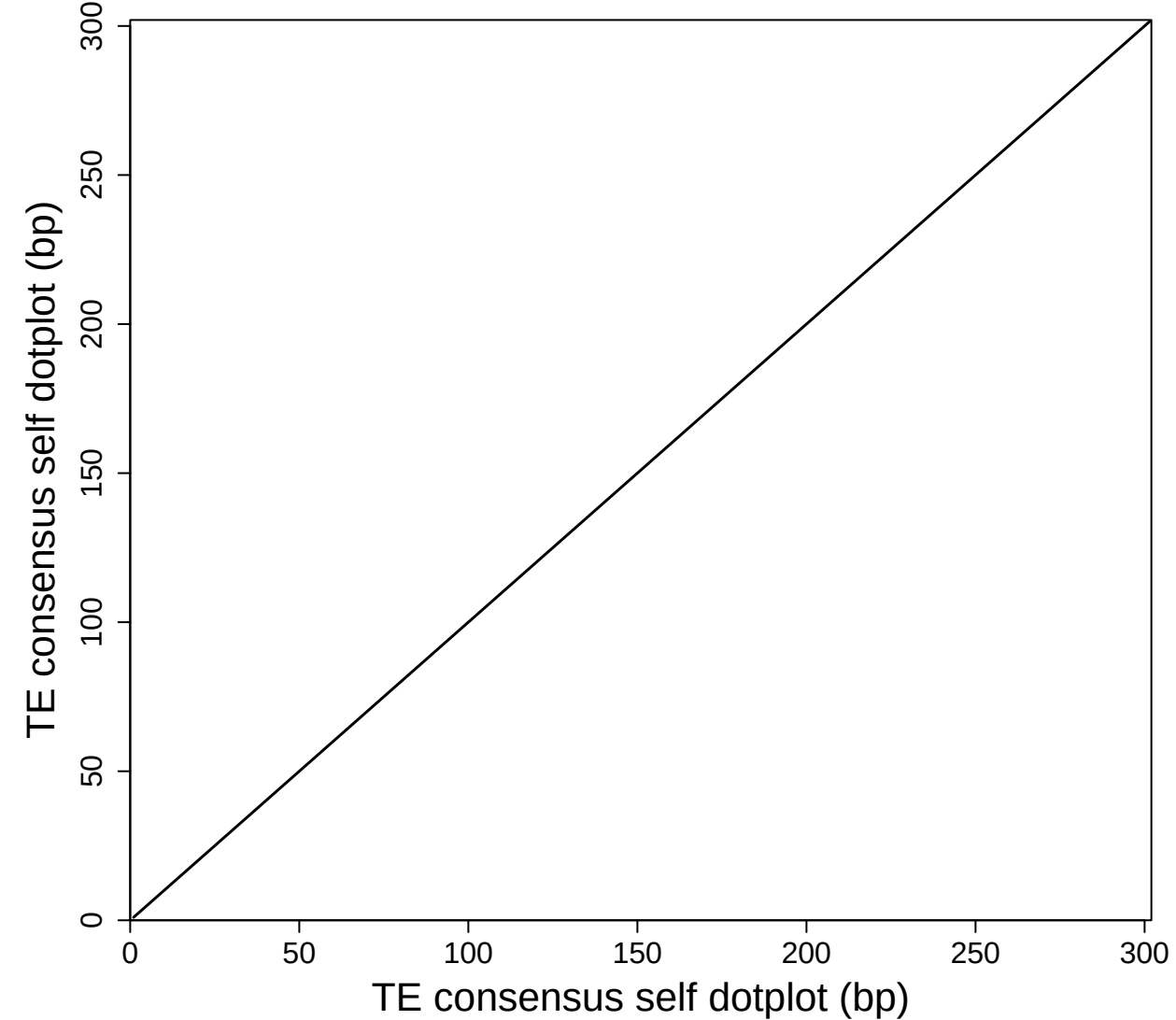
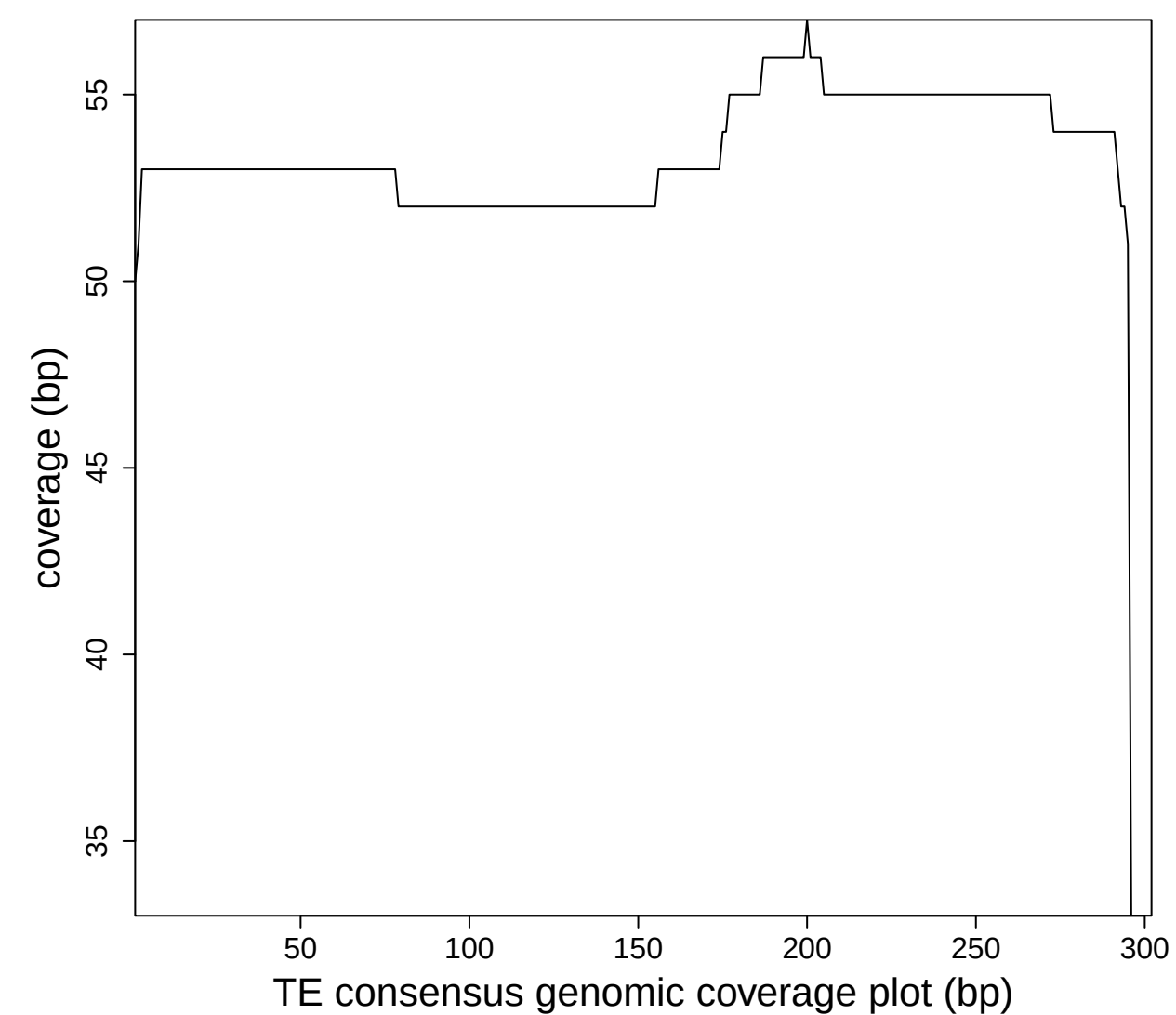
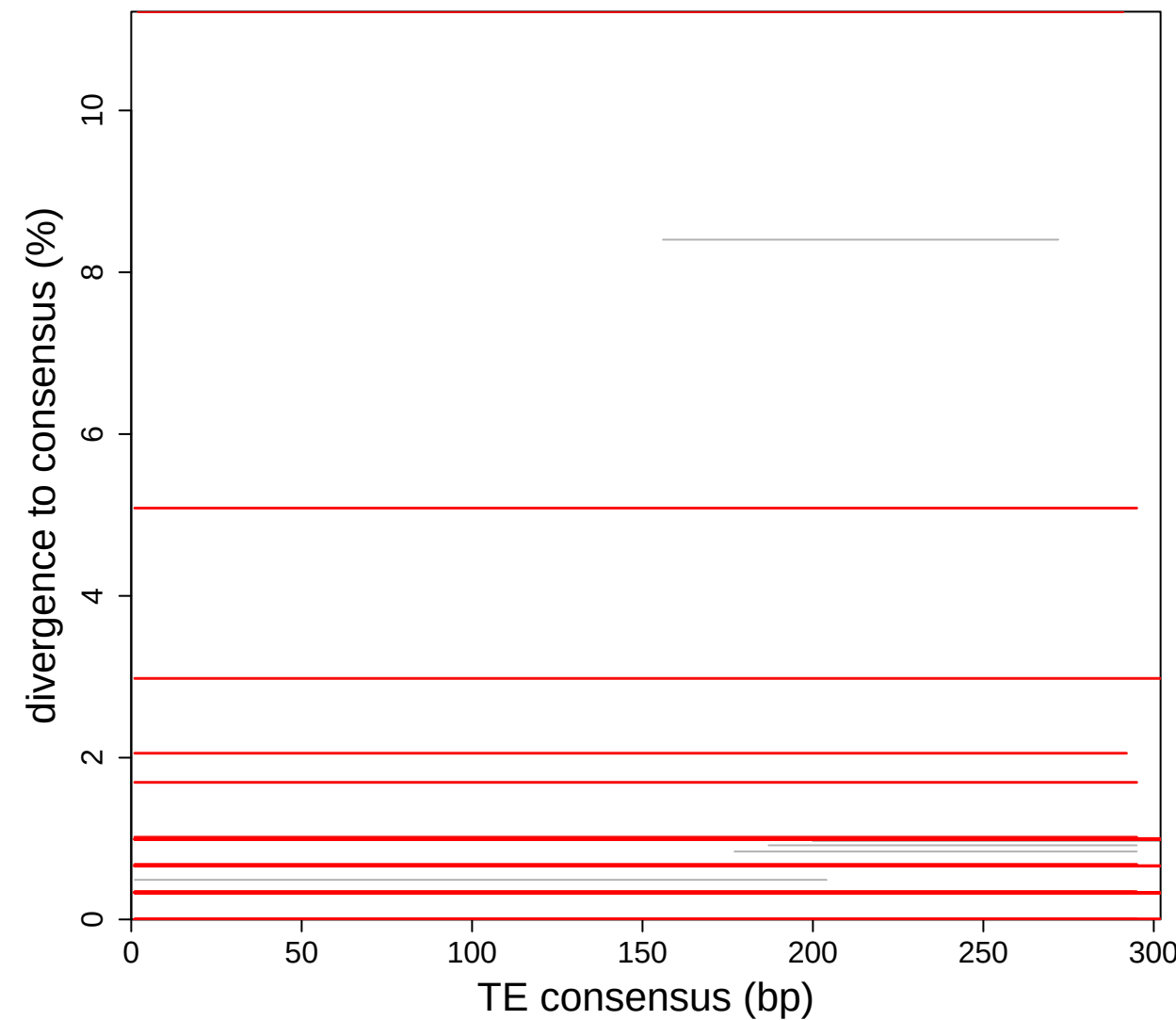
_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcln.fa_al
size: 6028bp; fragments: 120; full length: 0 (>=5425.2bp)

After TEtrimmer 6028 bp

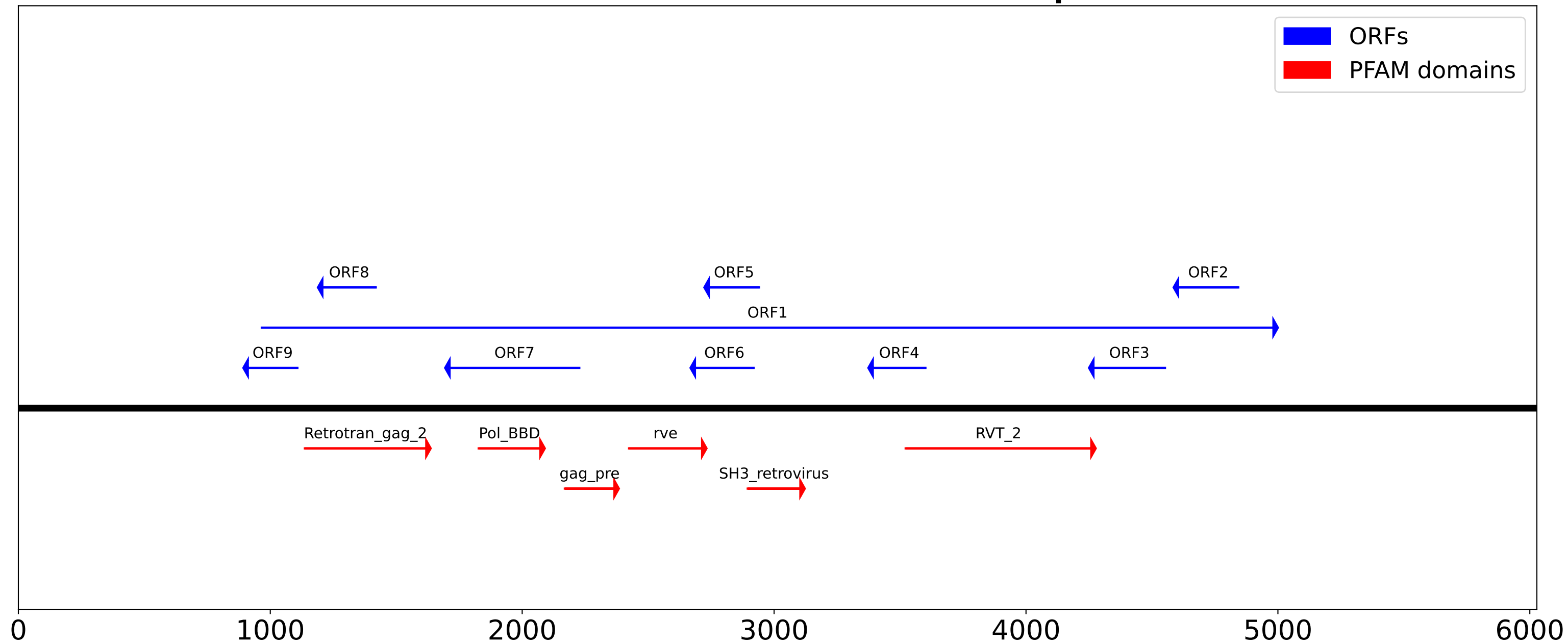




TE: ltr_1_family_103
size: 302bp; fragments: 58; full length: 50 (≥ 271.8 bp)



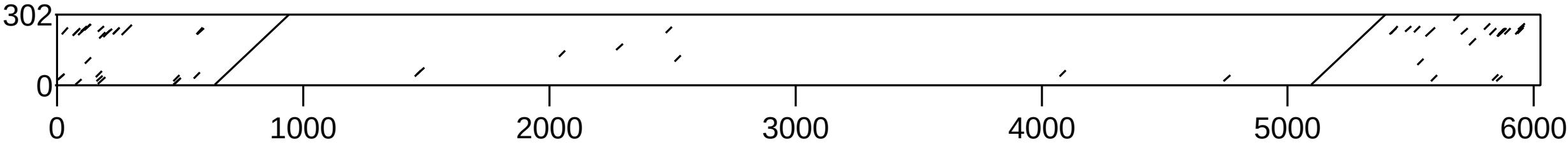
After TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)