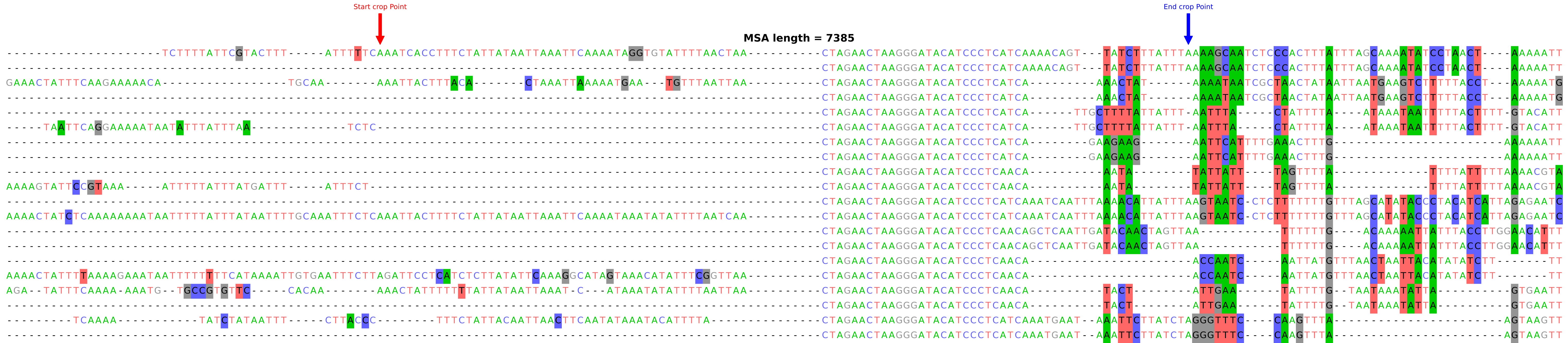
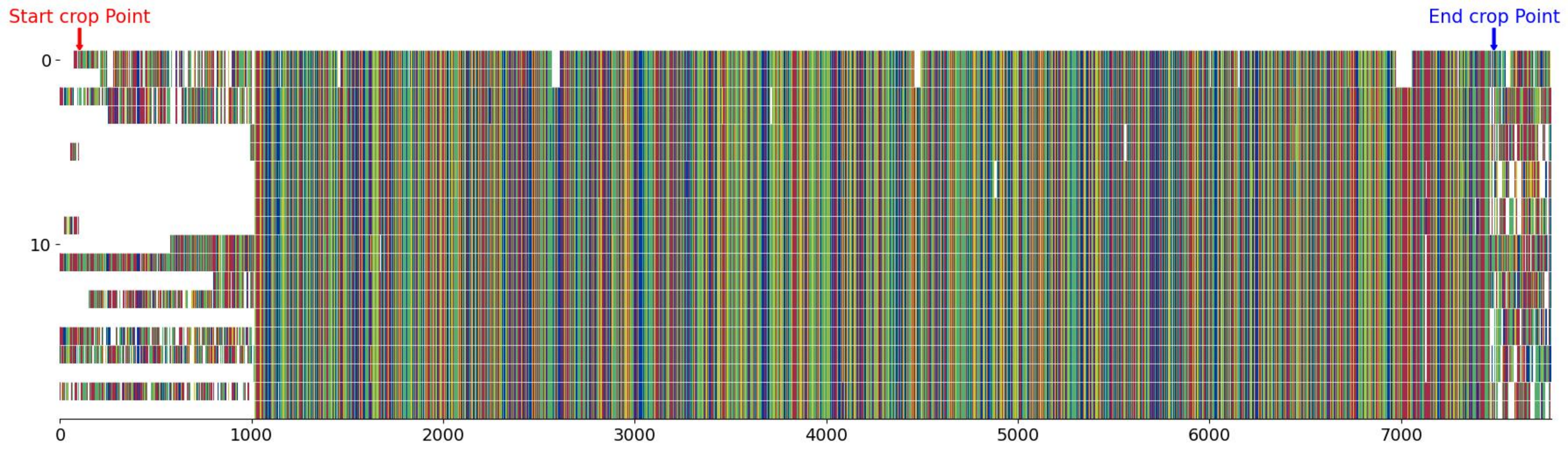
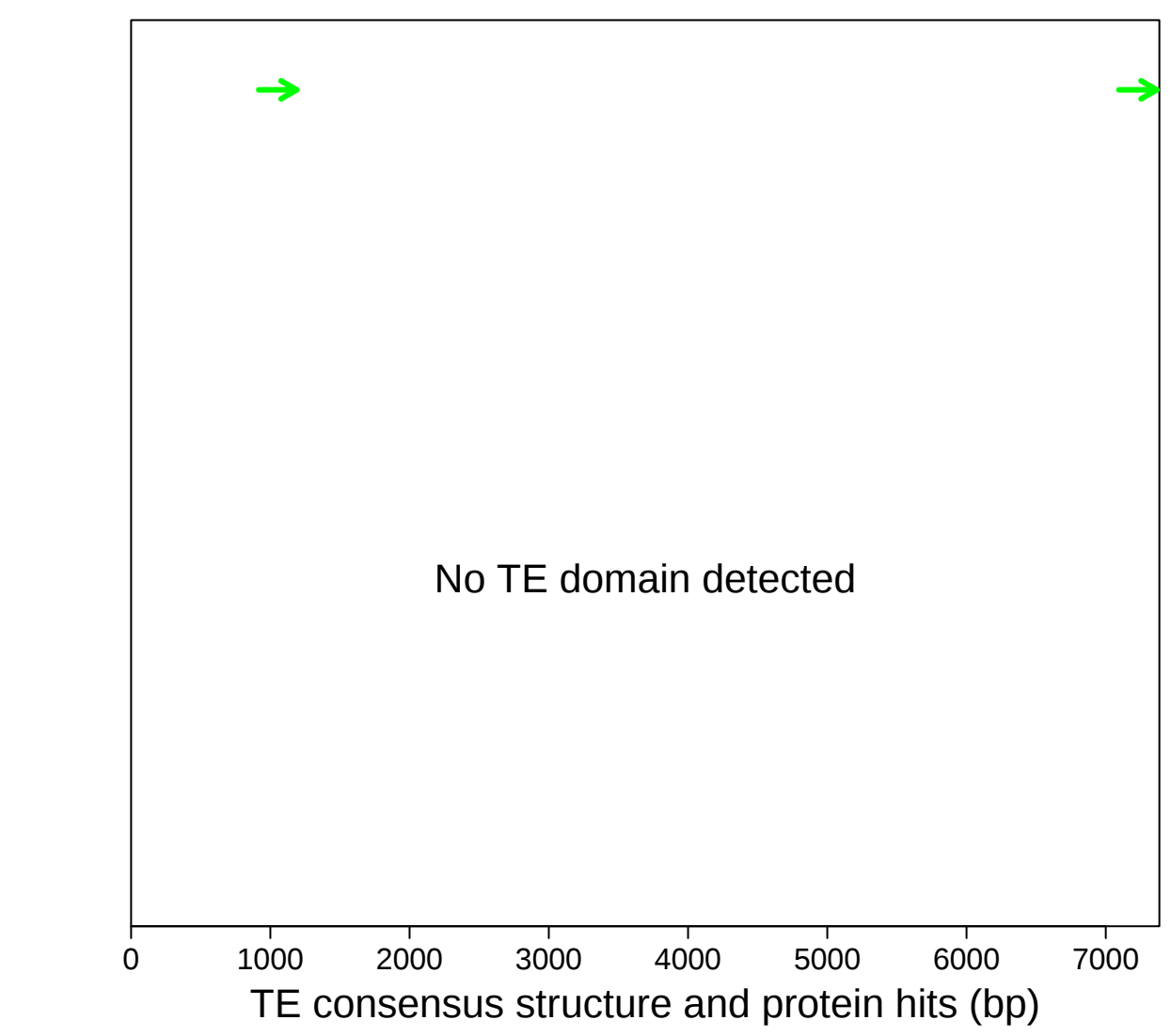
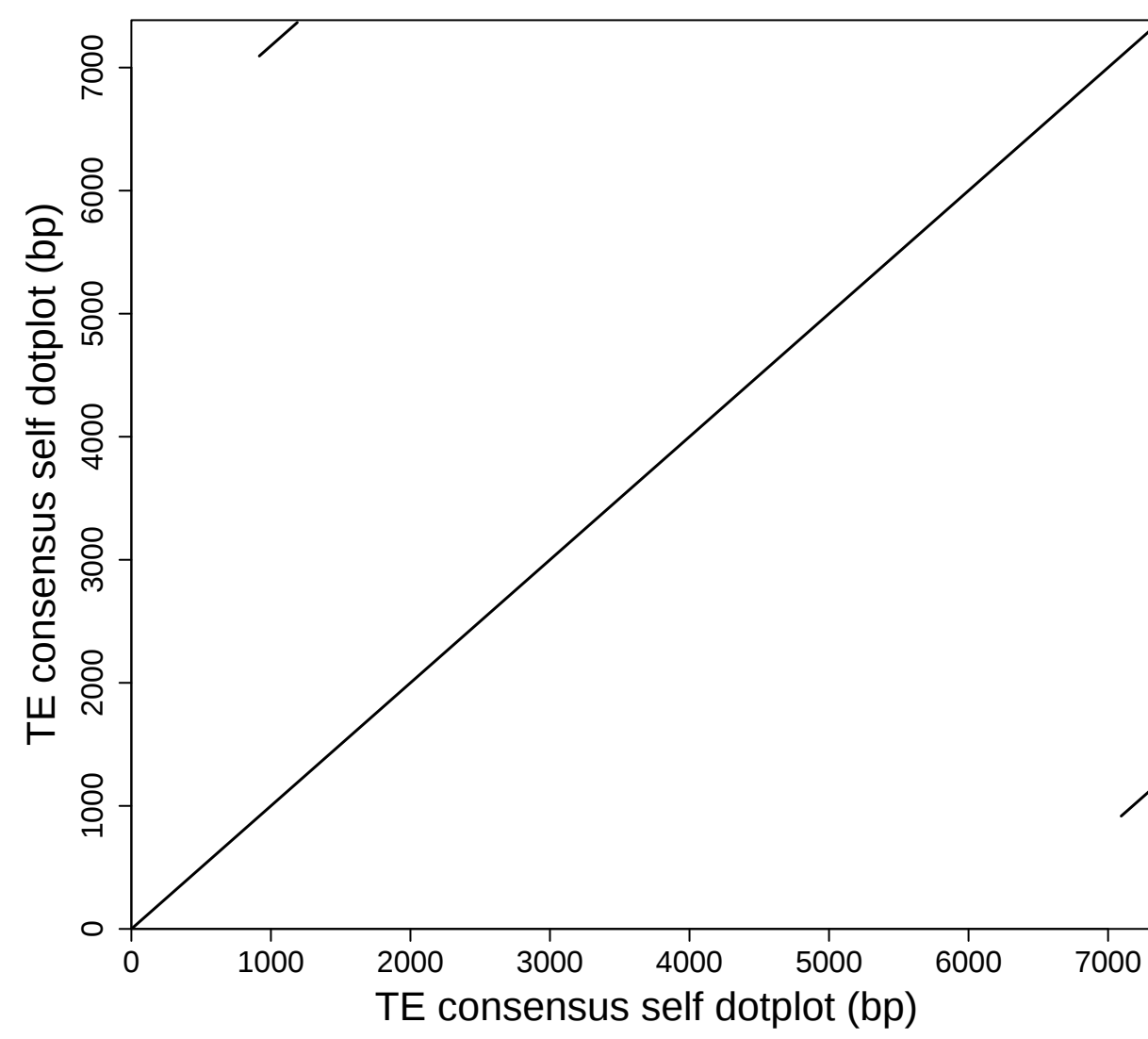
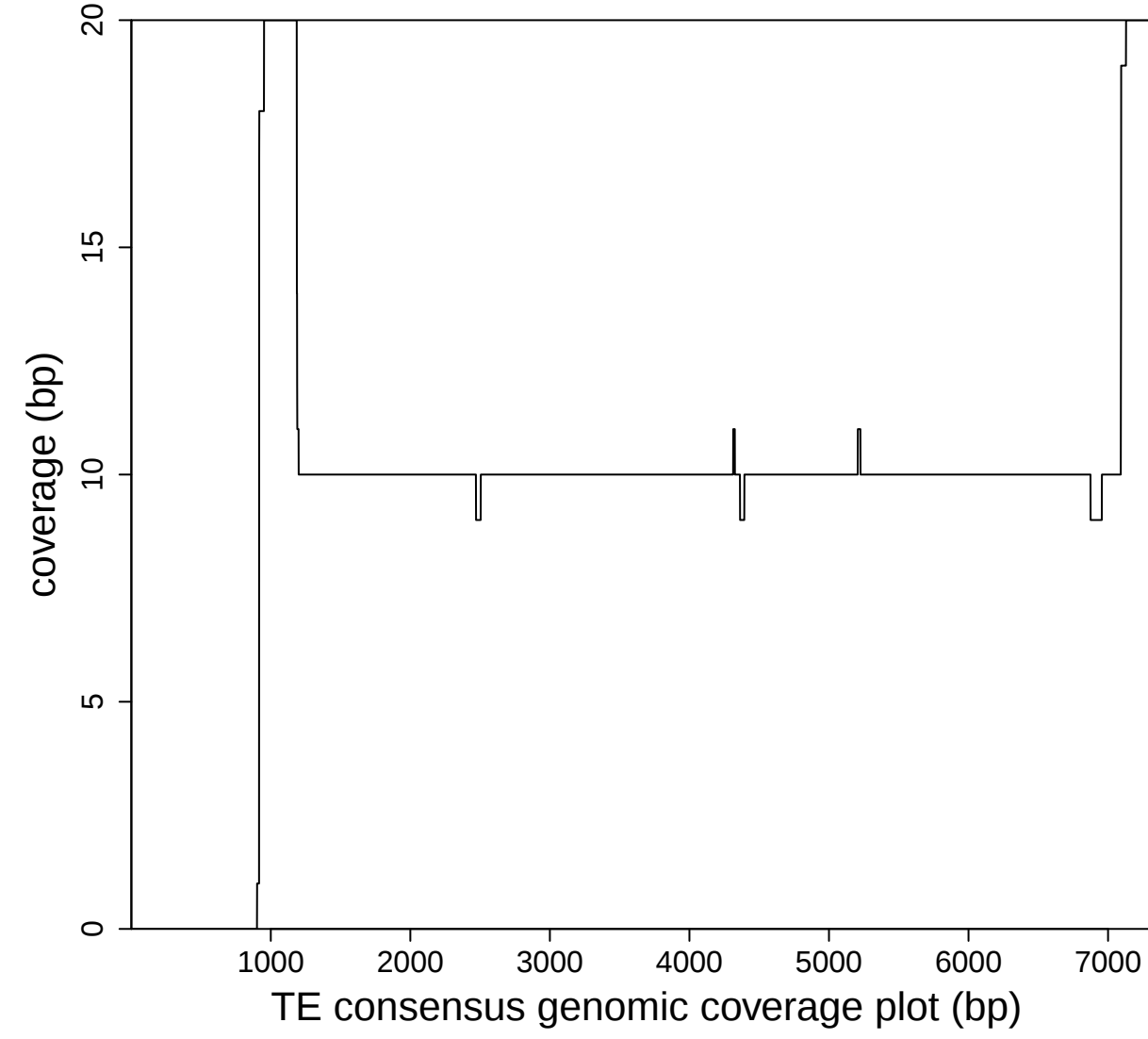
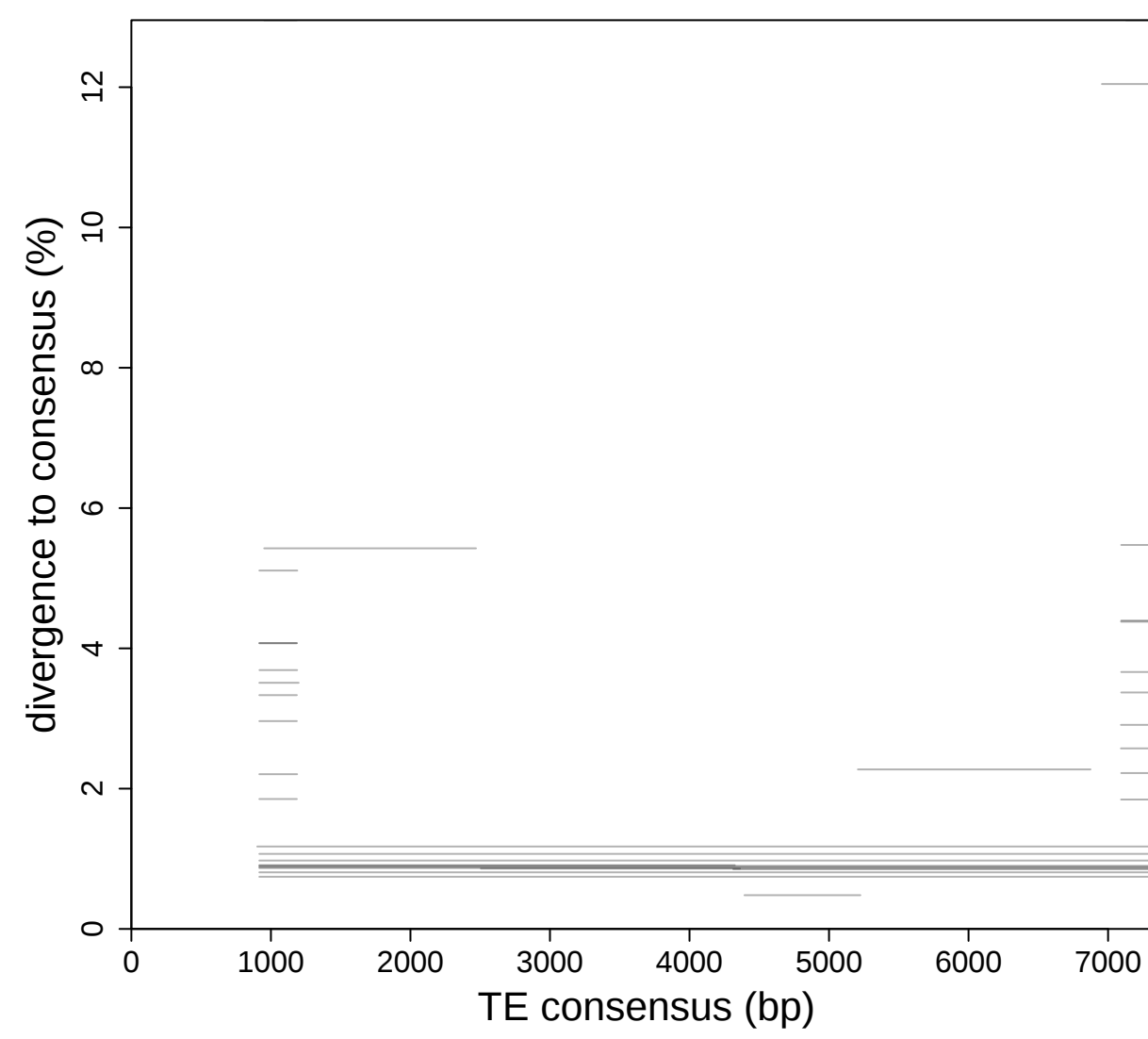


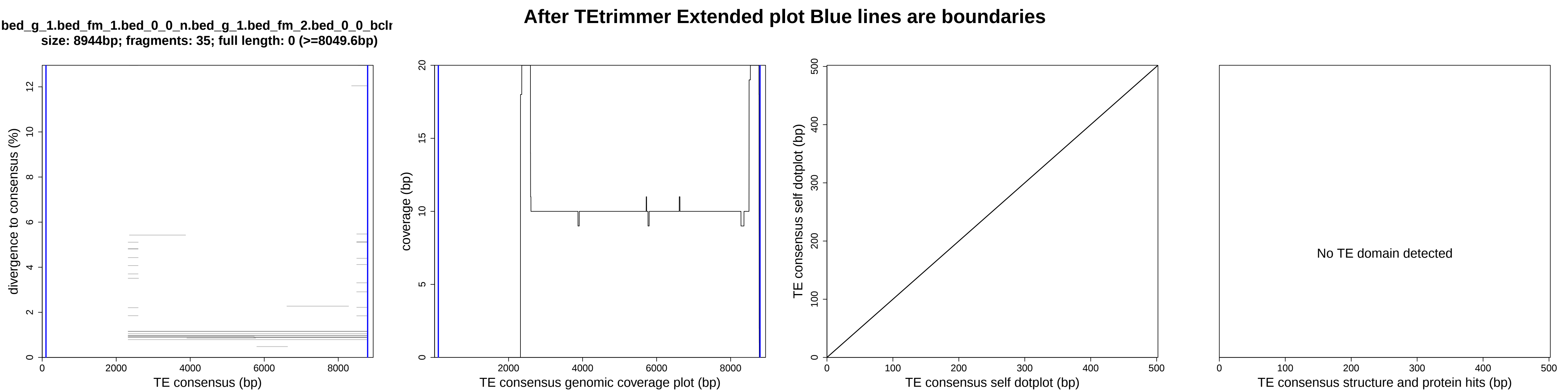




_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcIn.fa_al
size: 7385bp; fragments: 35; full length: 0 (>=6646.5bp)

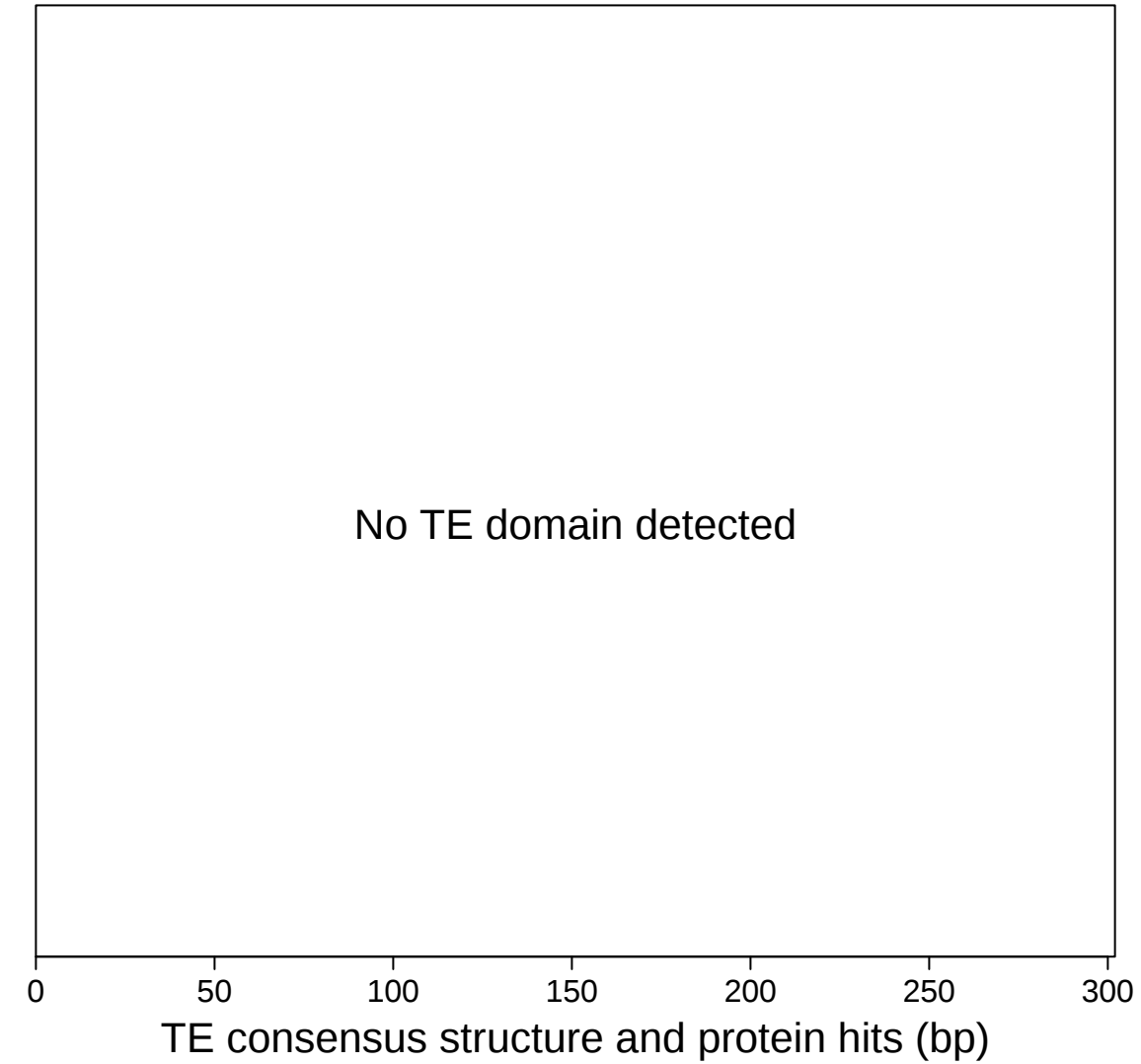
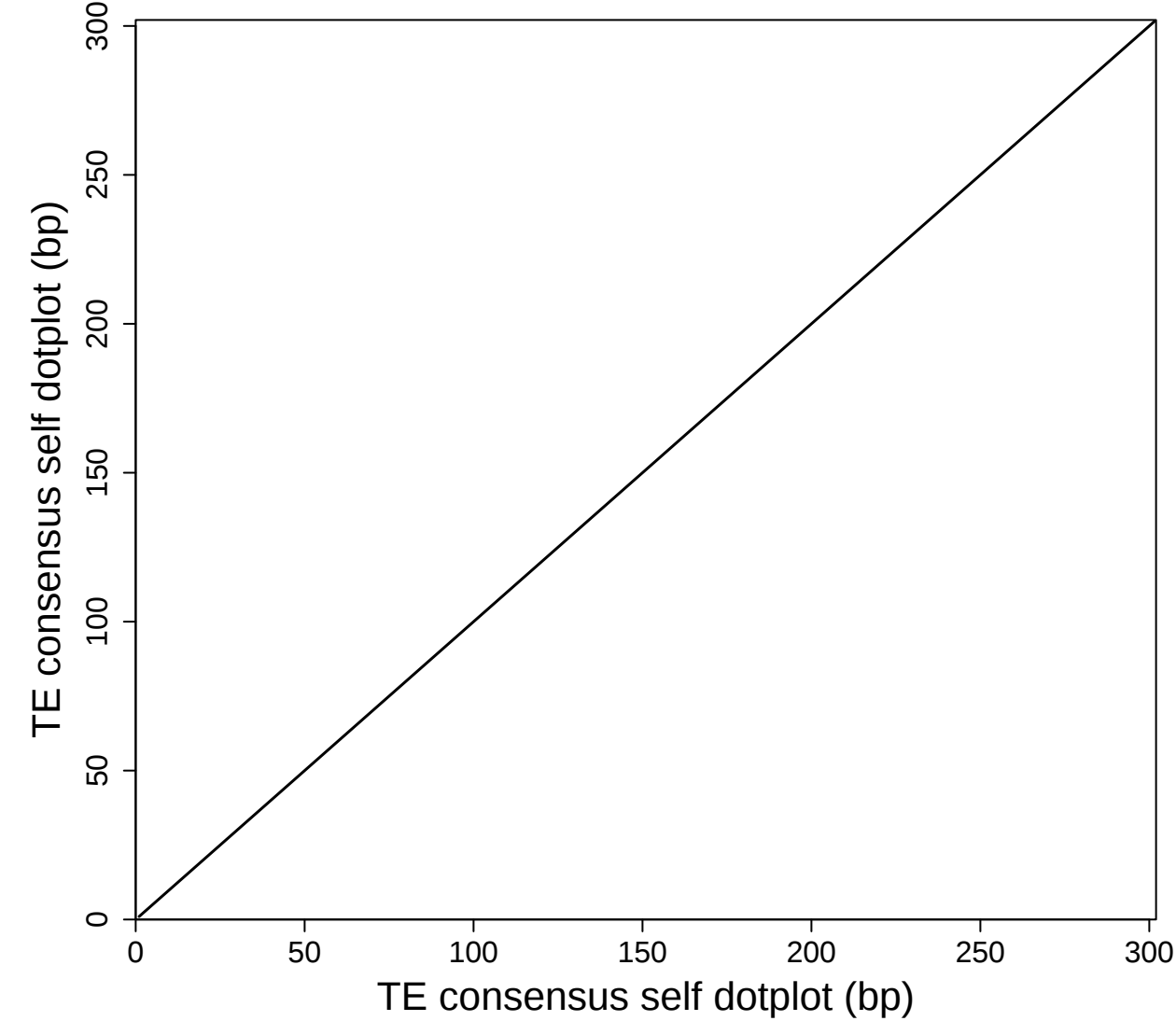
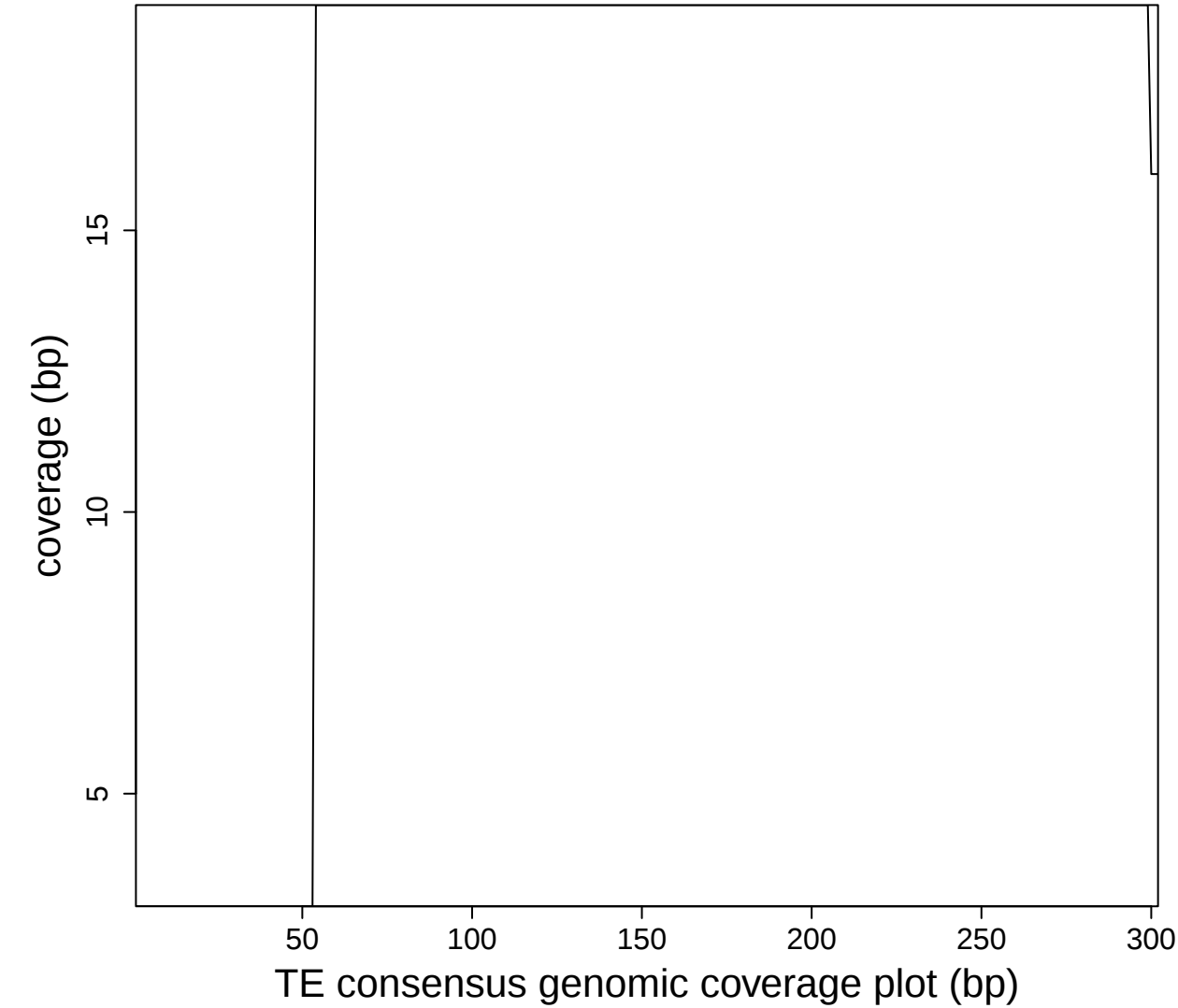
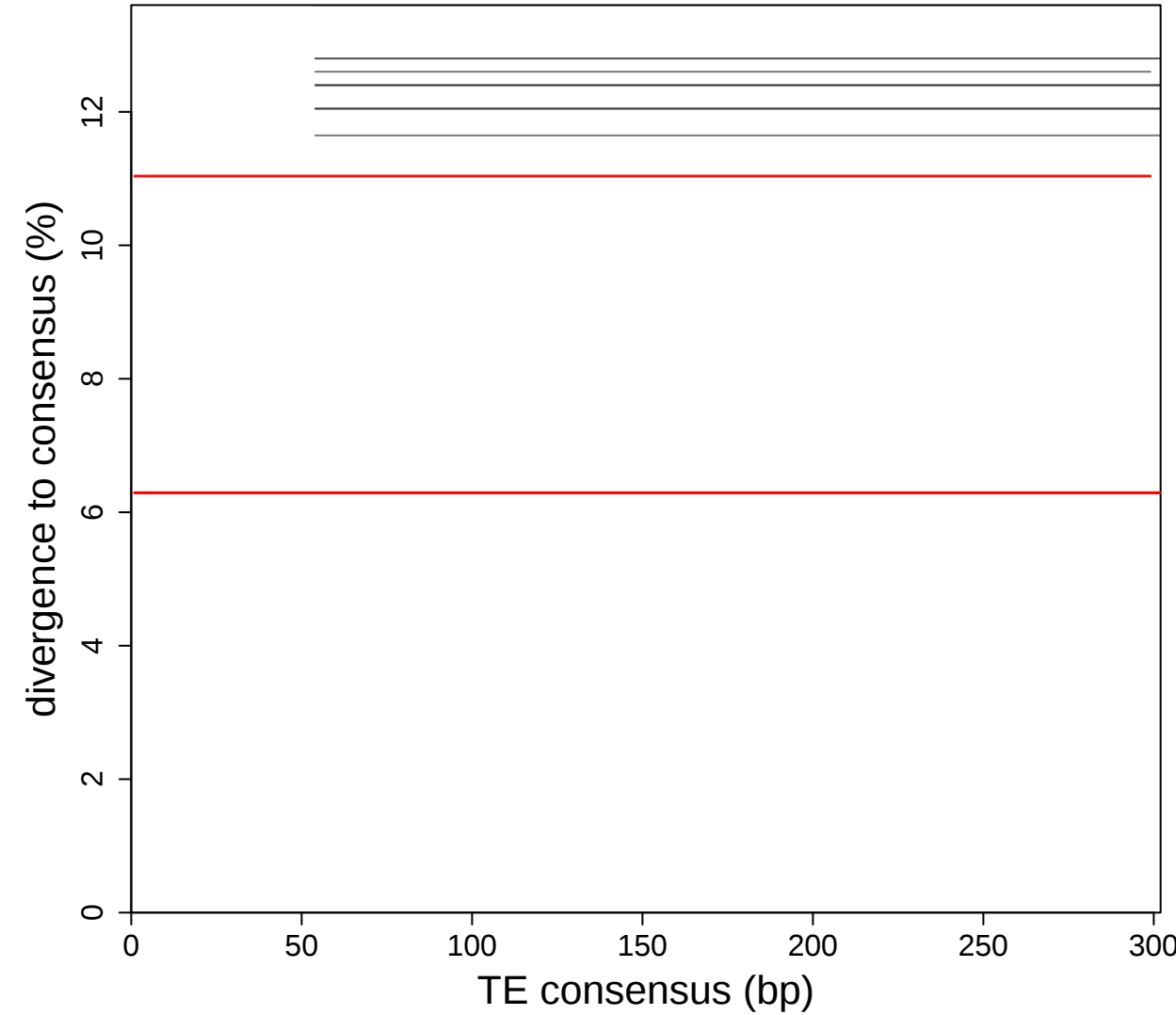
After TEtrimmer 7385 bp



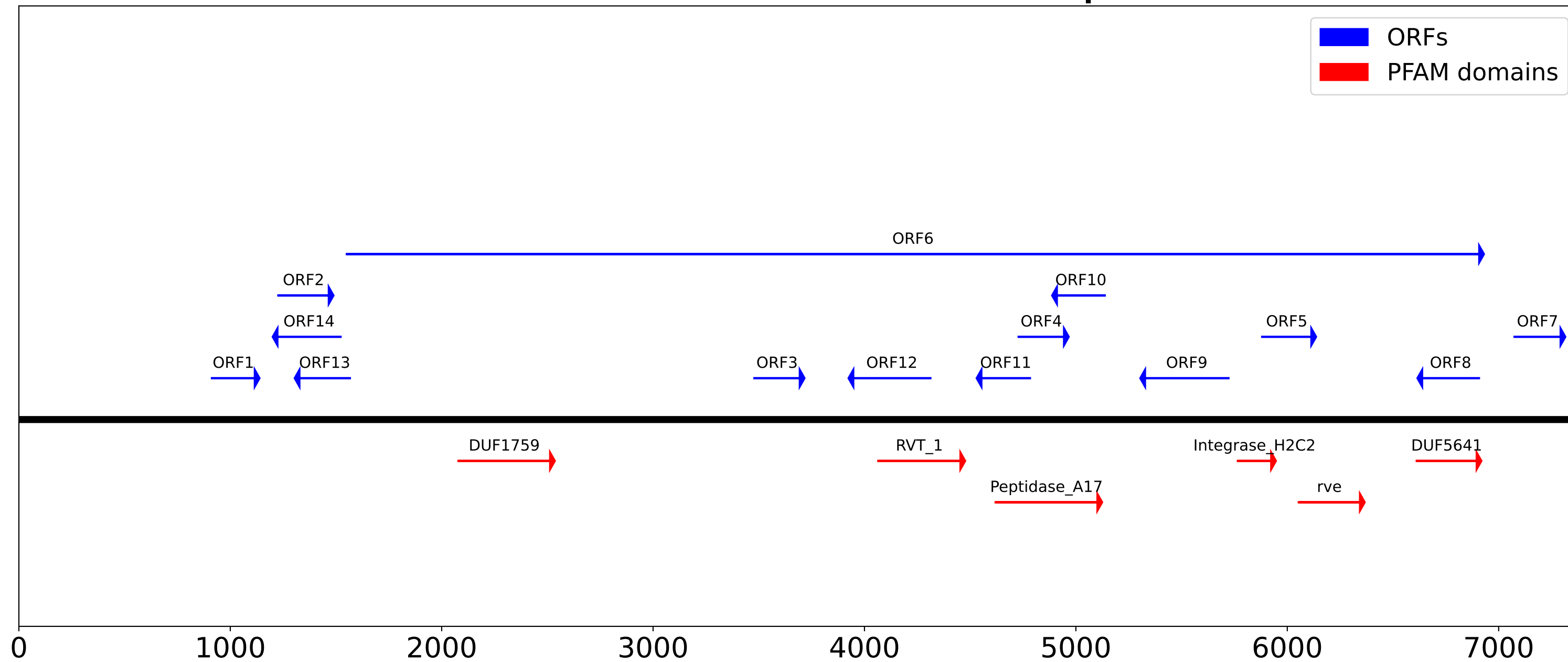


Before TEtrimmer 302 bp

TE: ltr_1_family_182
size: 302bp; fragments: 19; full length: 3 (≥ 271.8 bp)



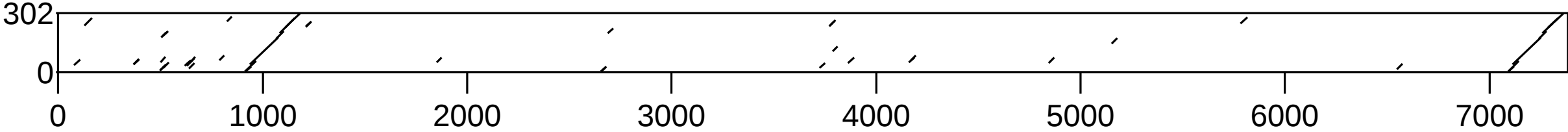
After TEtrimmer ORF and PFAM domain plot



Dotplot

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

nsus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)