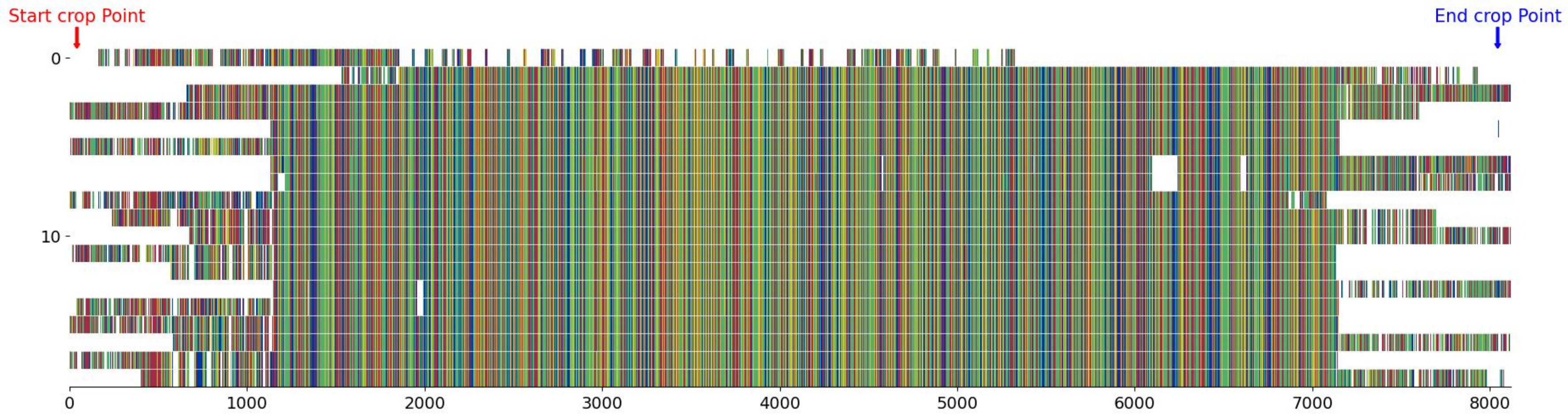


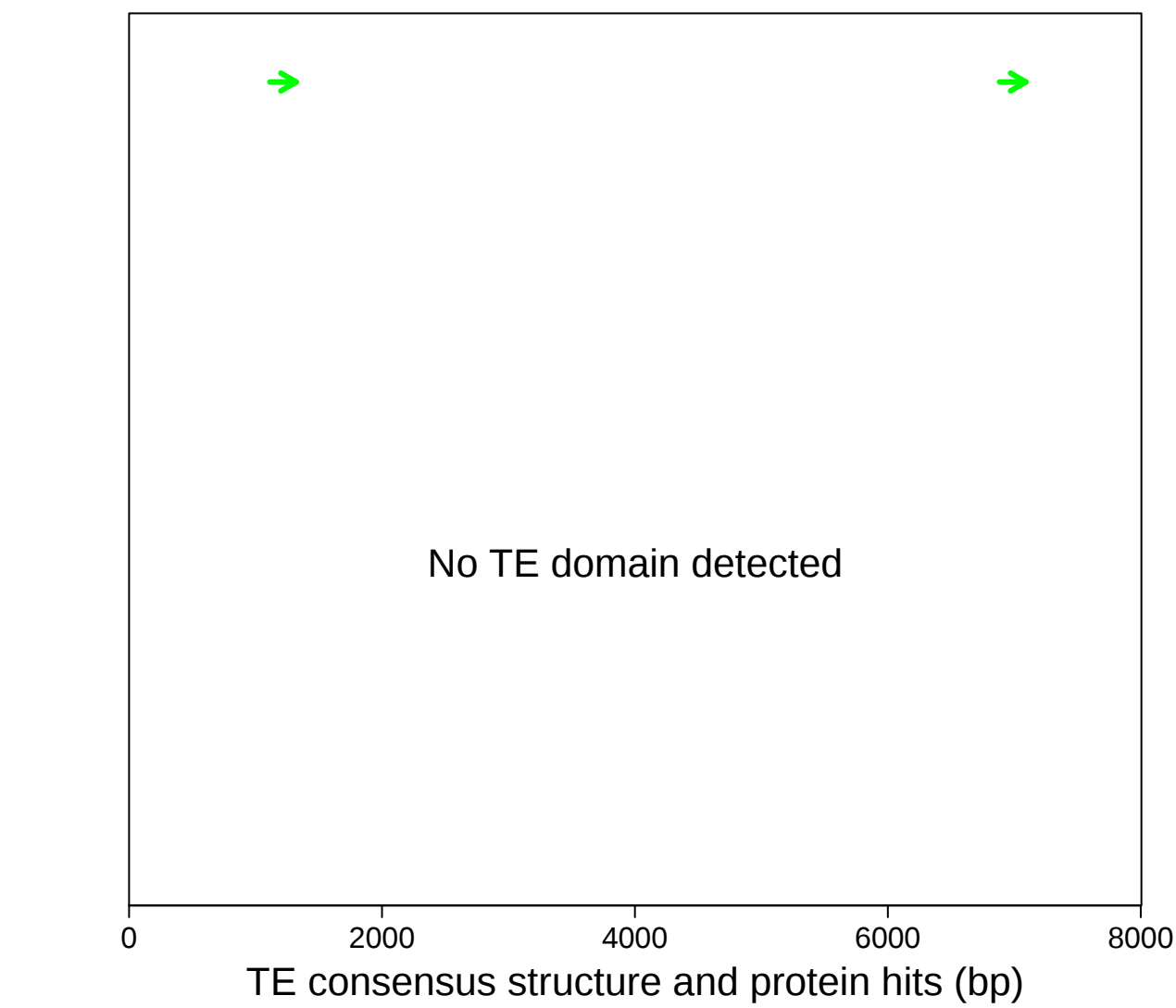
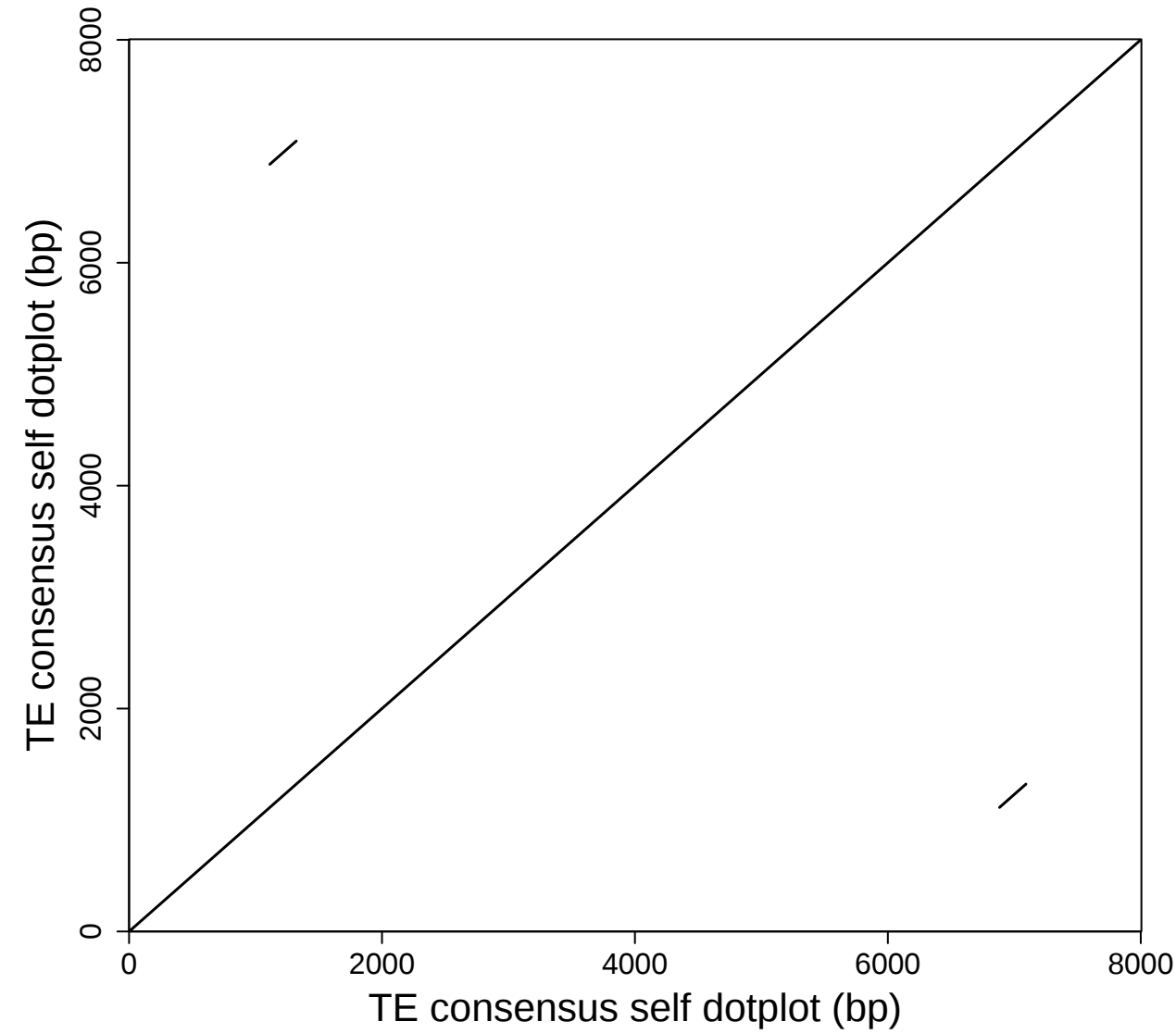
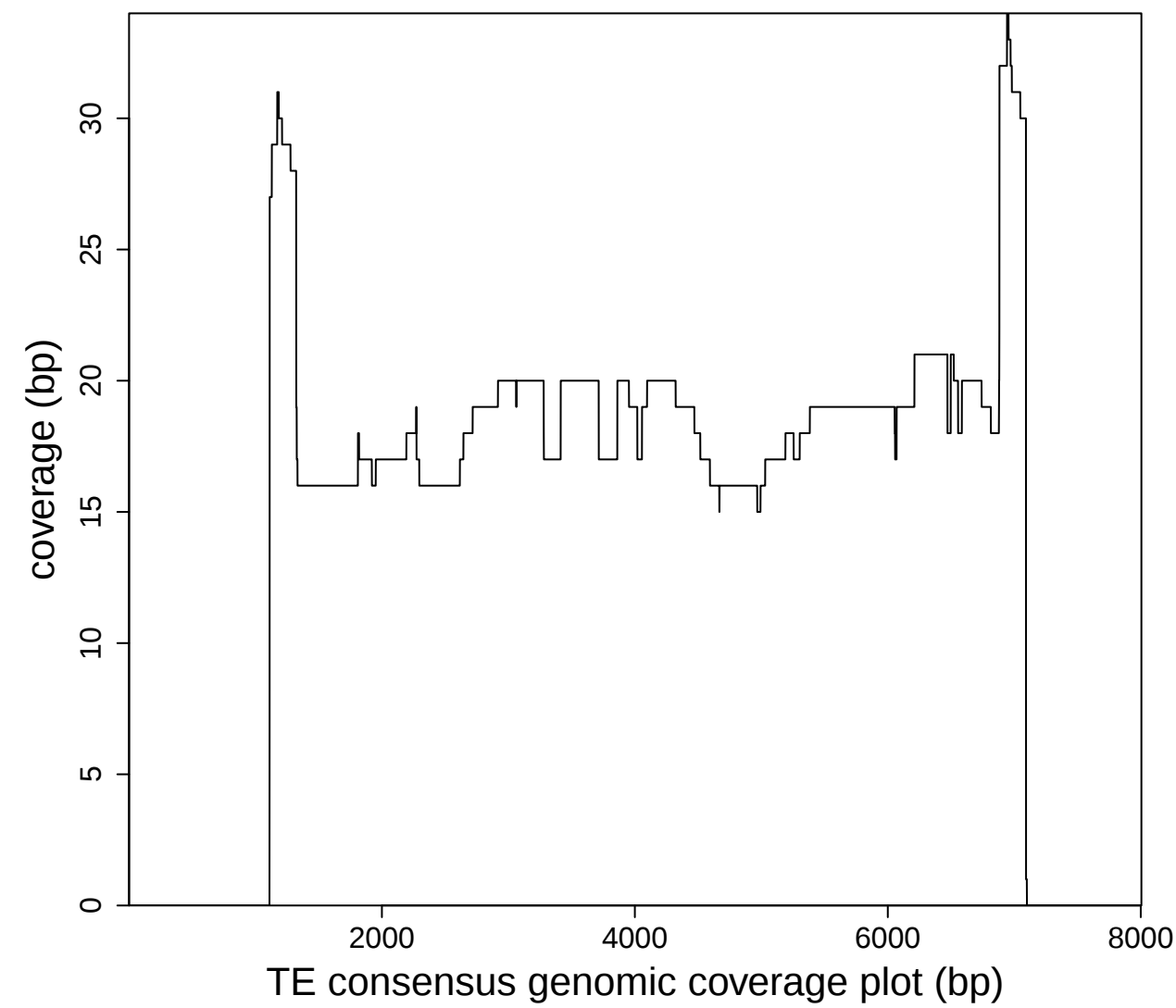
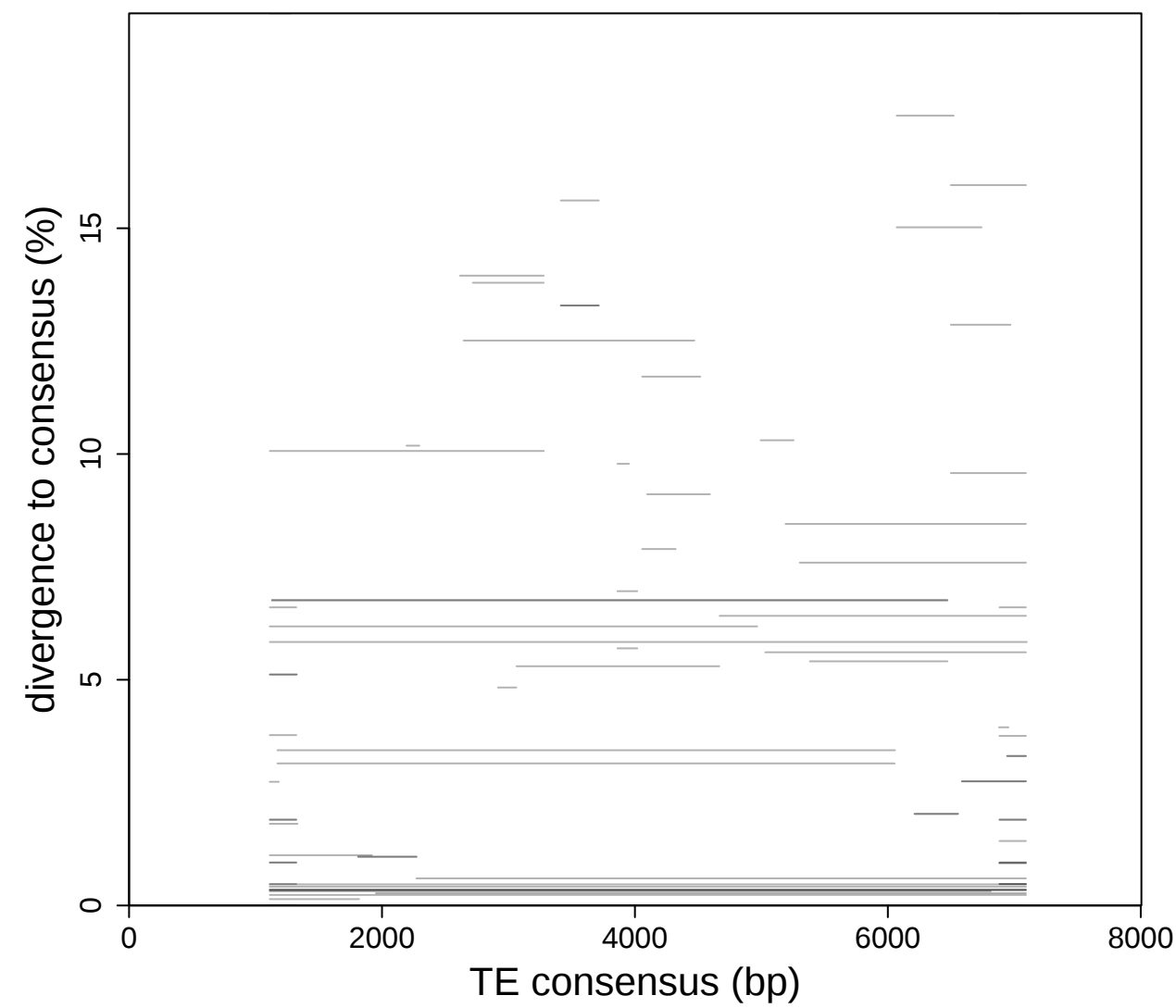
MSA length = 8005

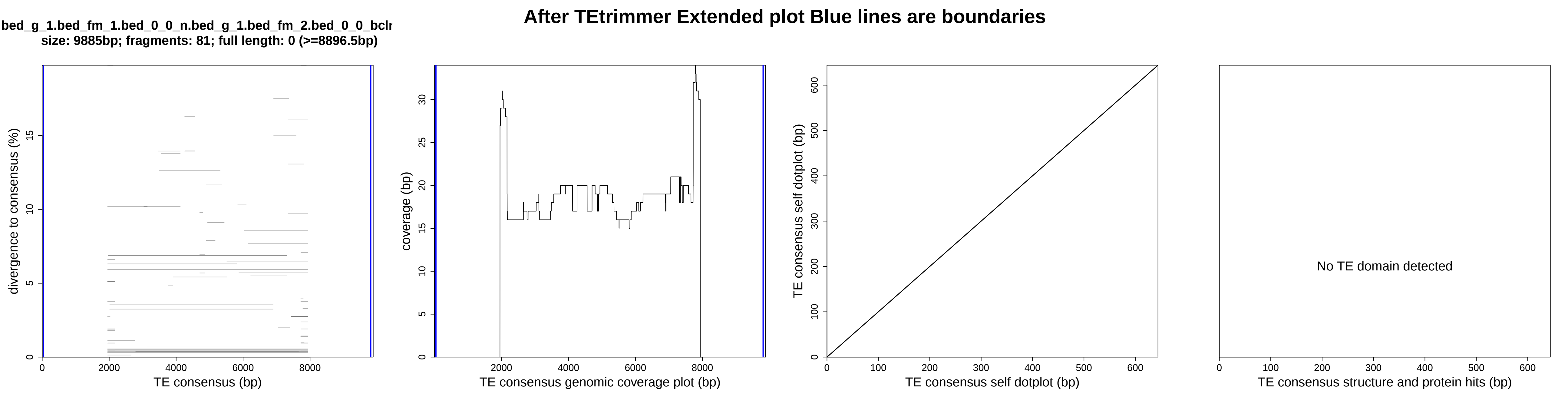
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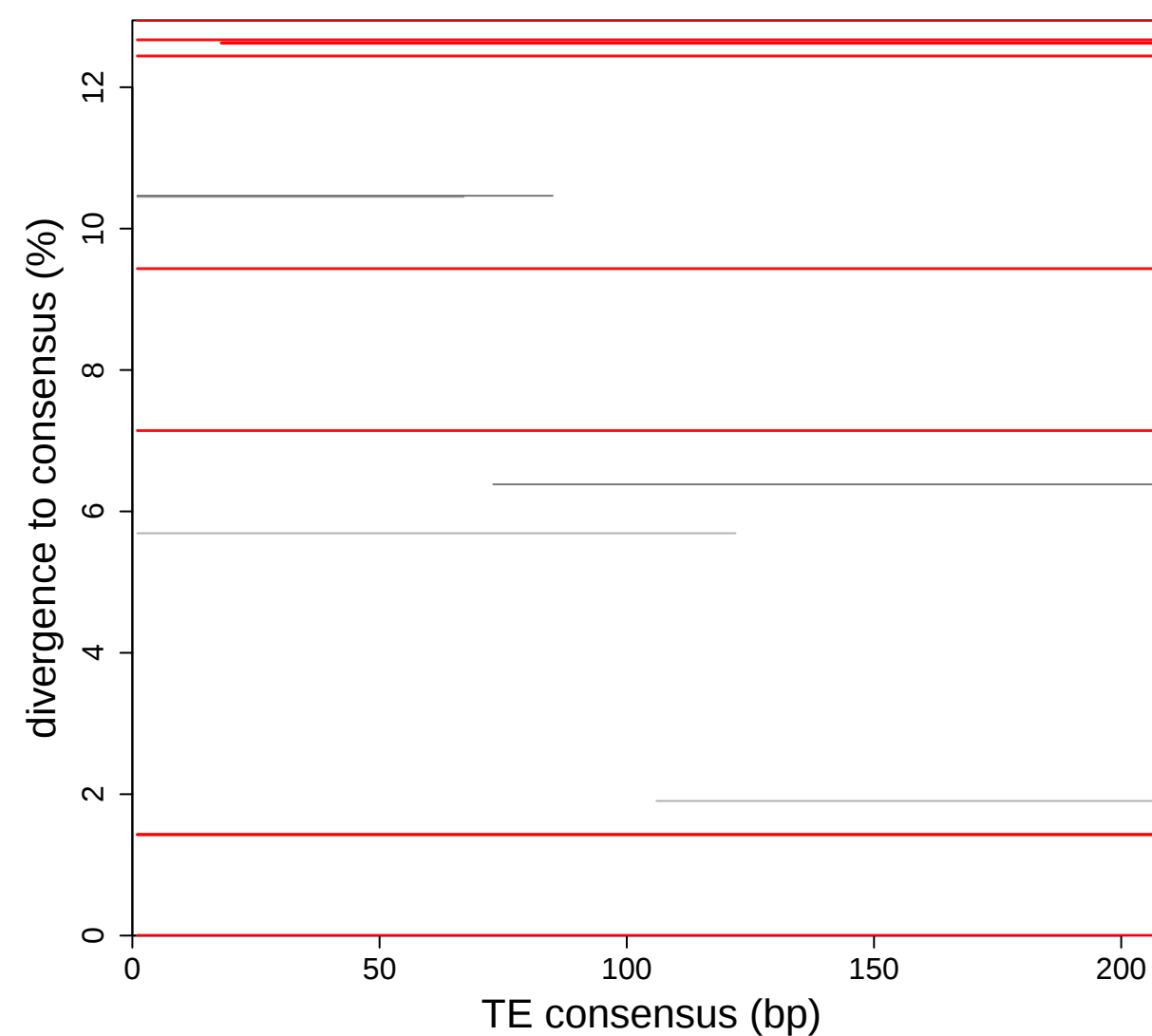
_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcIn.fa_al
size: 8005bp; fragments: 81; full length: 0 (>=7204.5bp)

After TEtrimmer 8005 bp

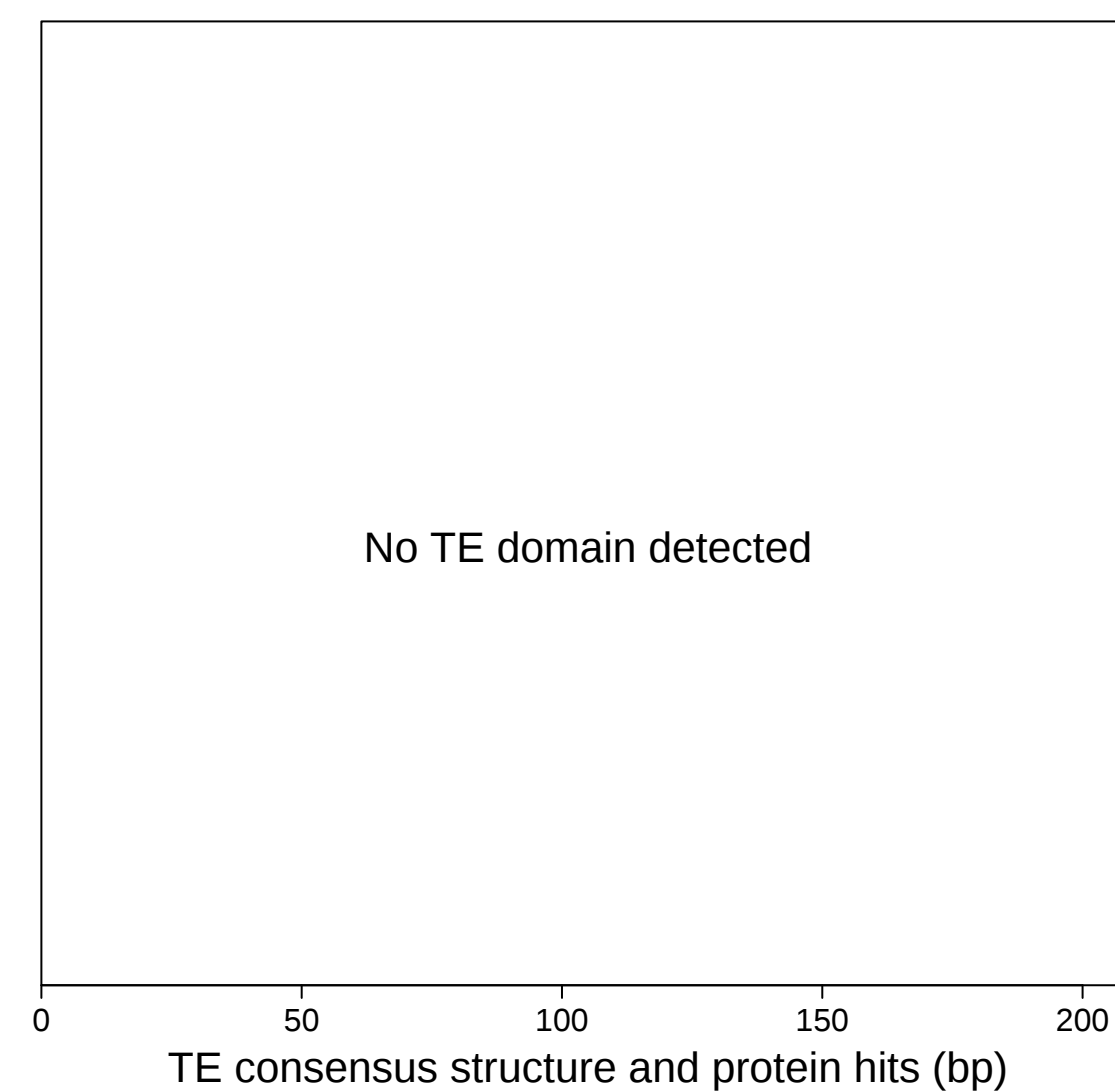
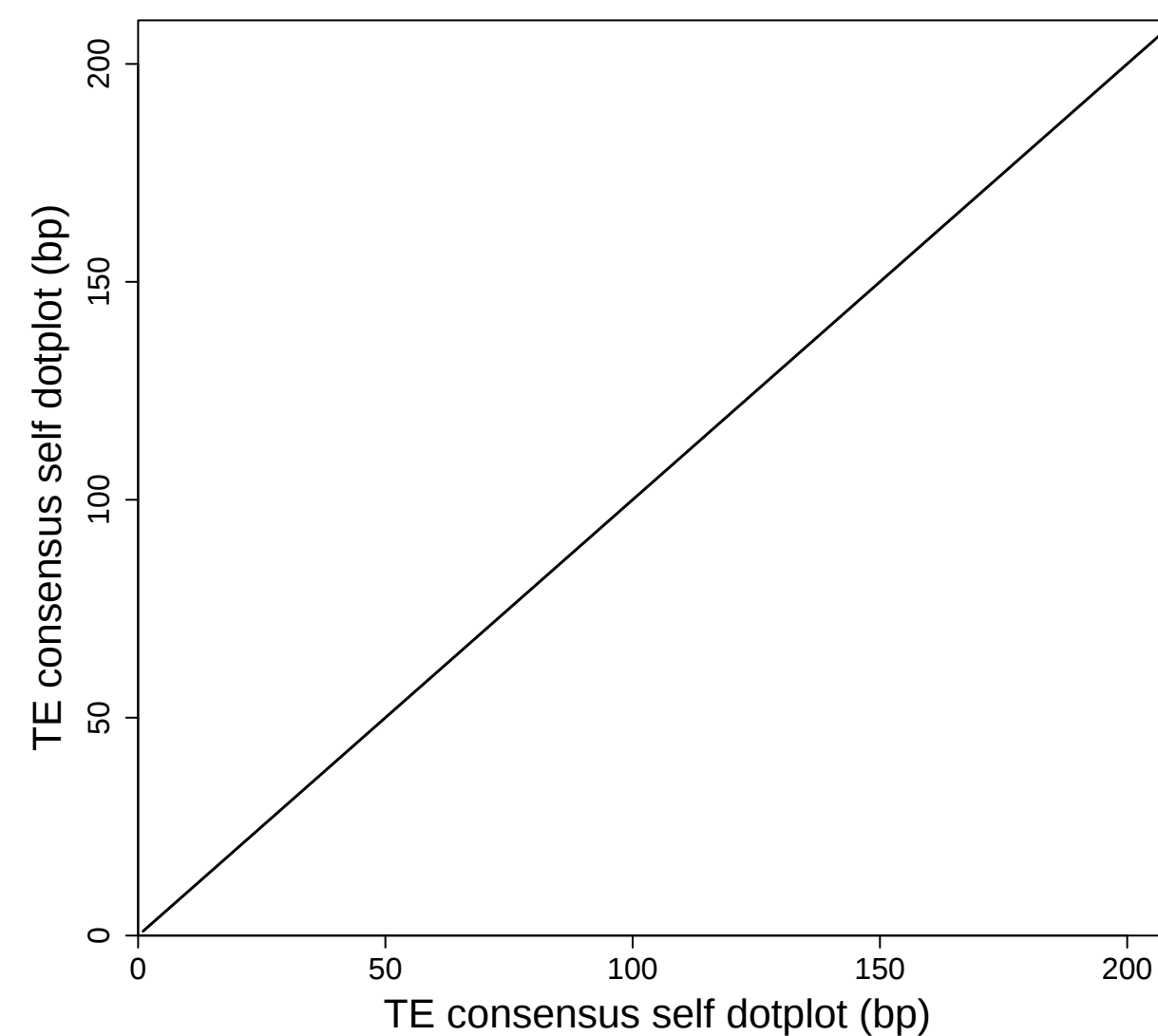
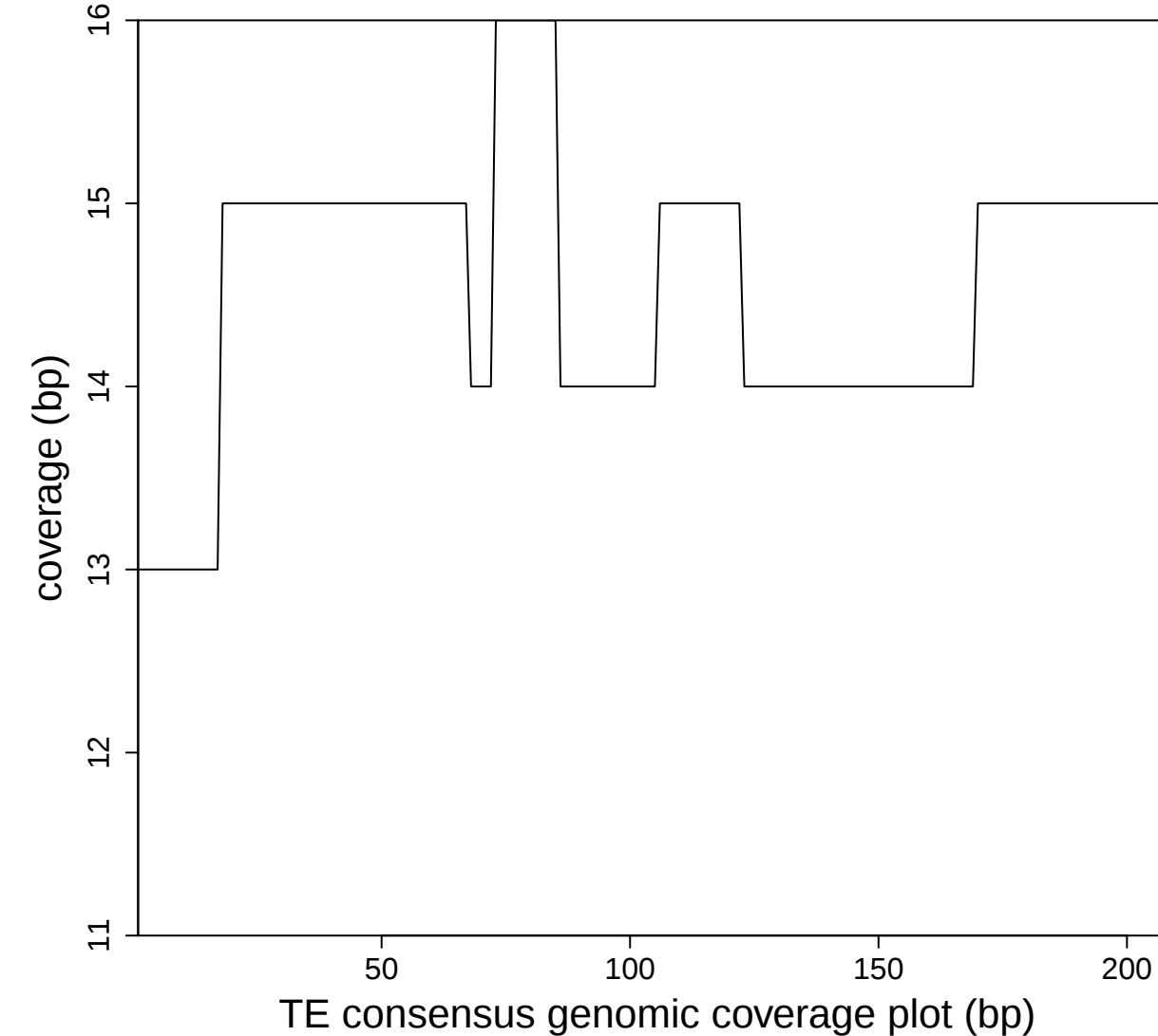




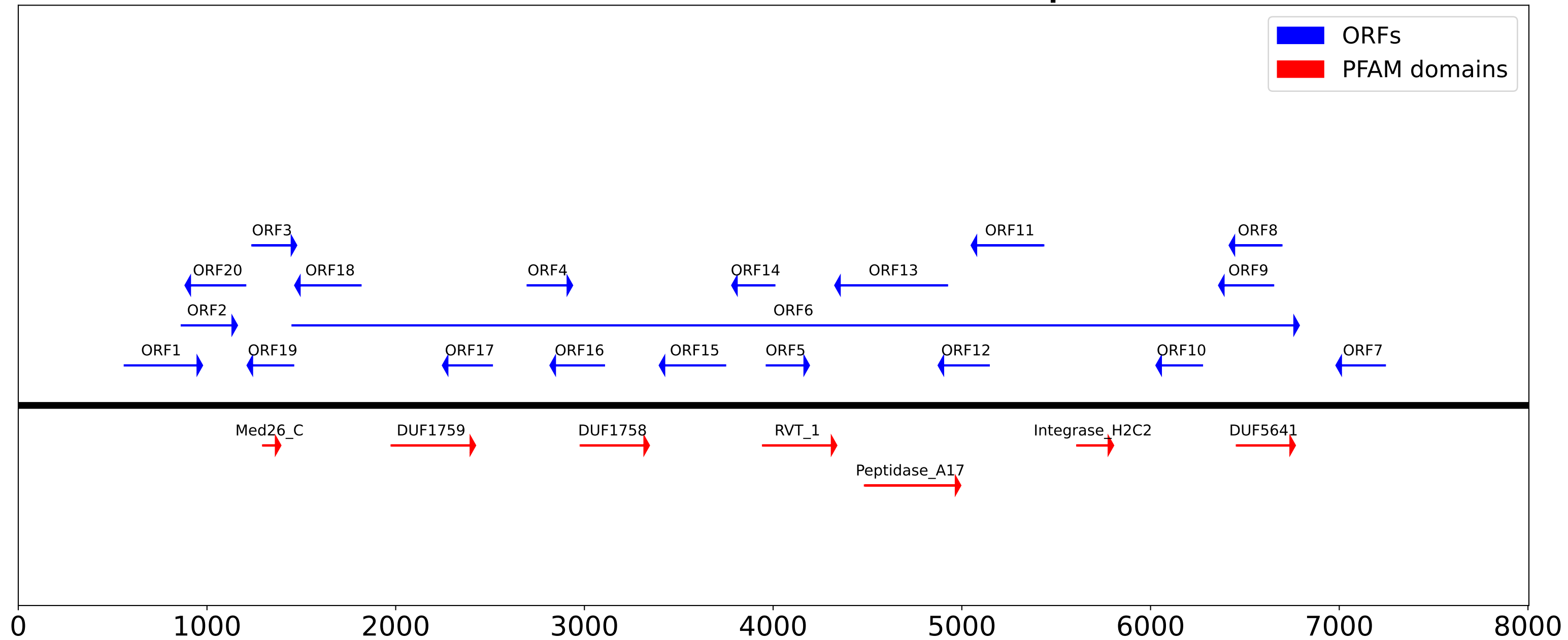
TE: ltr_1_family_284
size: 210bp; fragments: 19; full length: 11 (≥ 189 bp)



Before TEtrimmer 210 bp



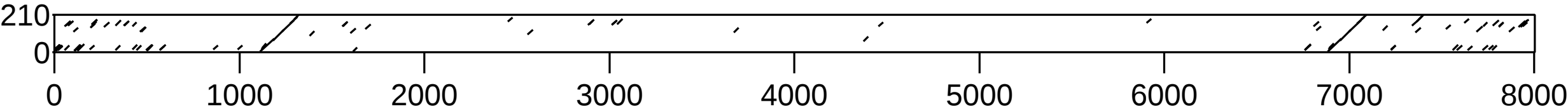
After TETrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)