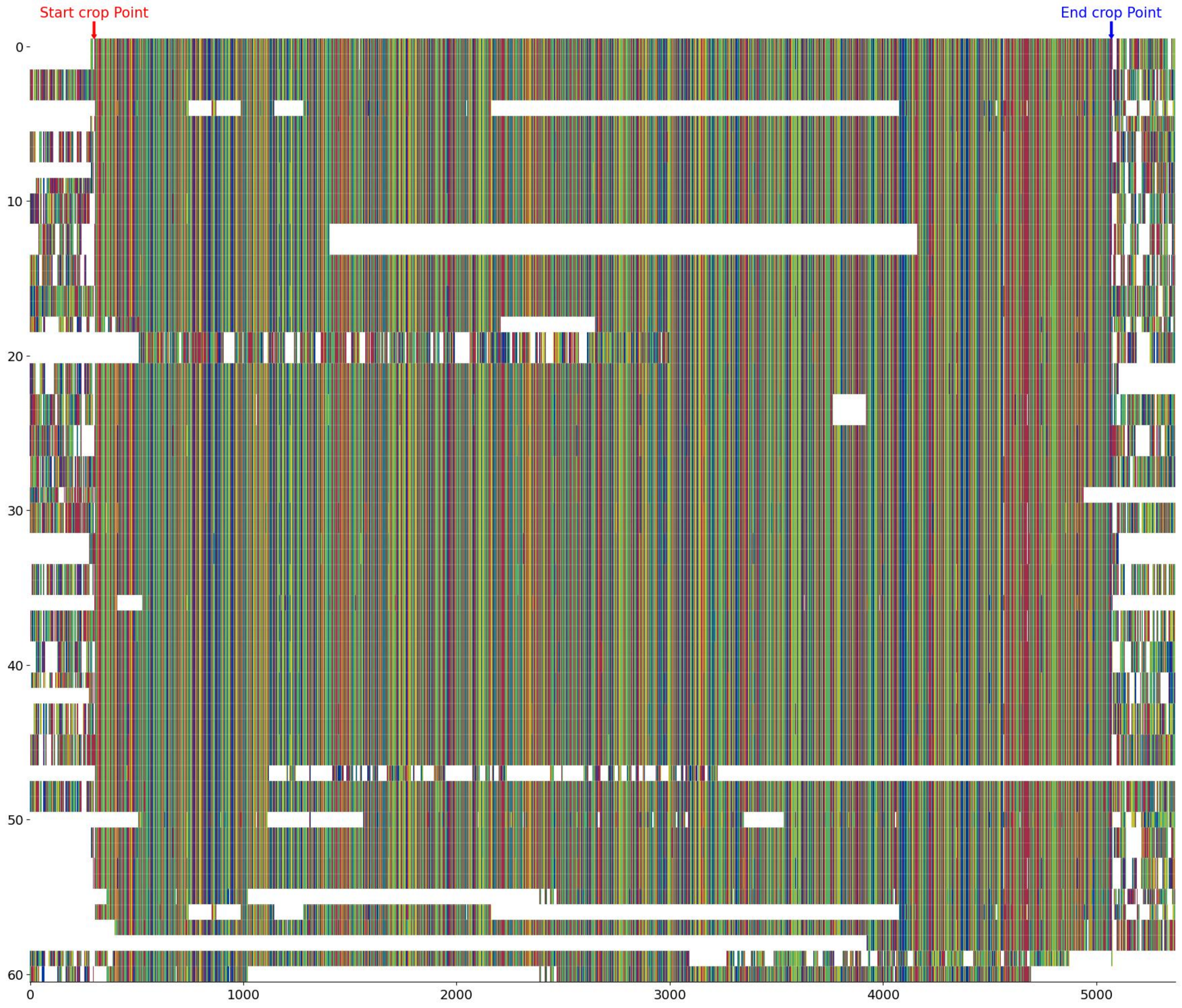
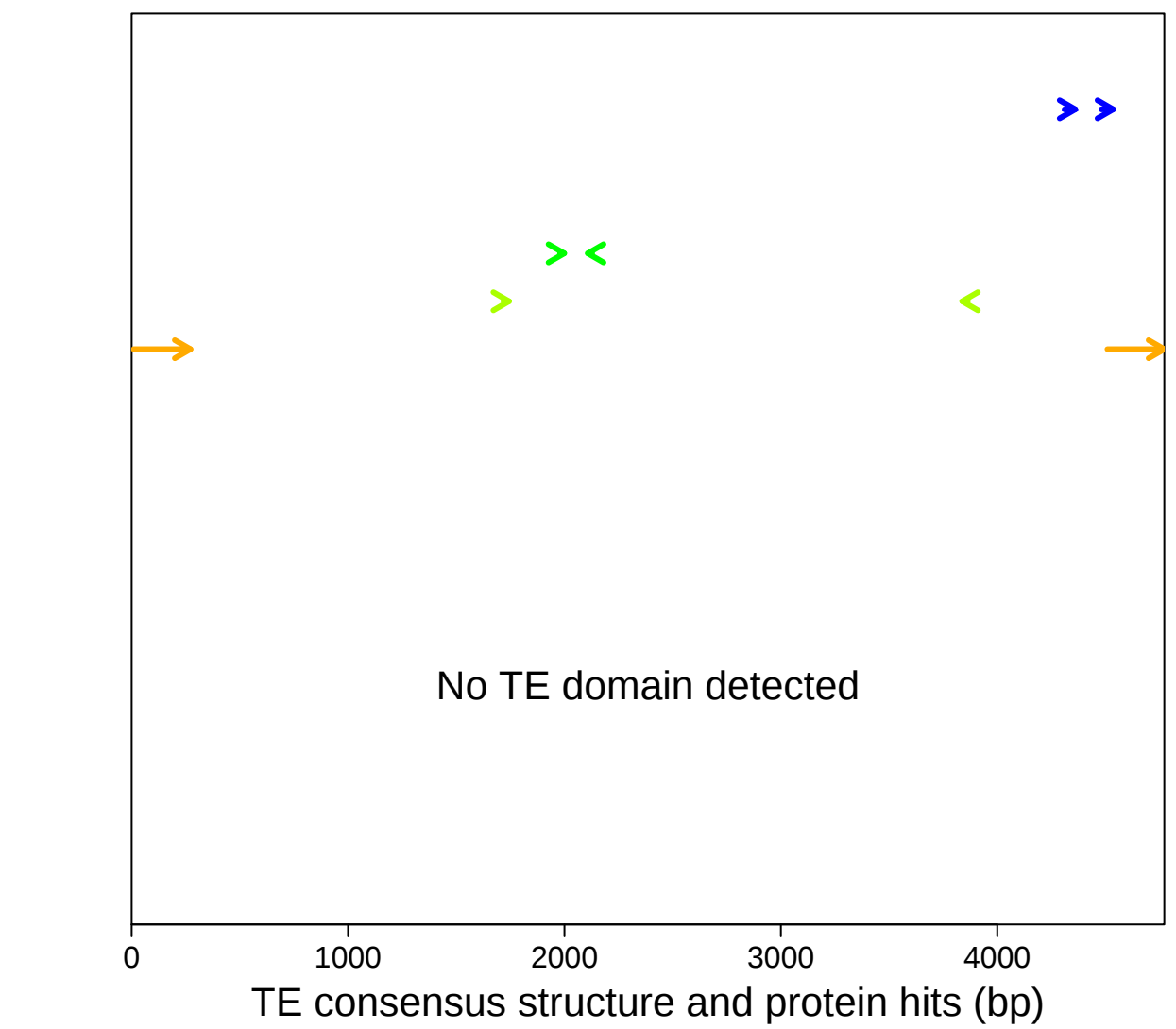
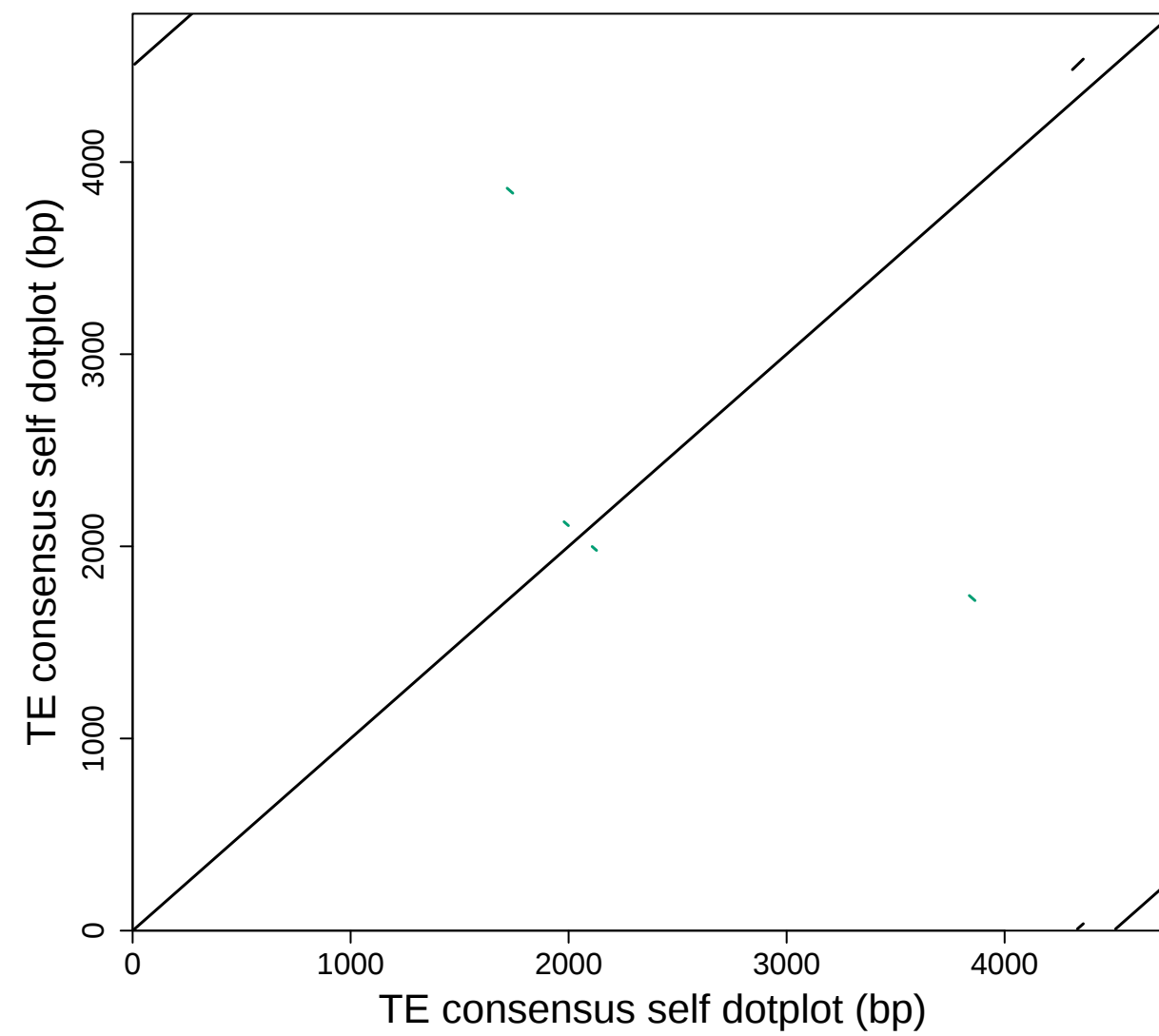
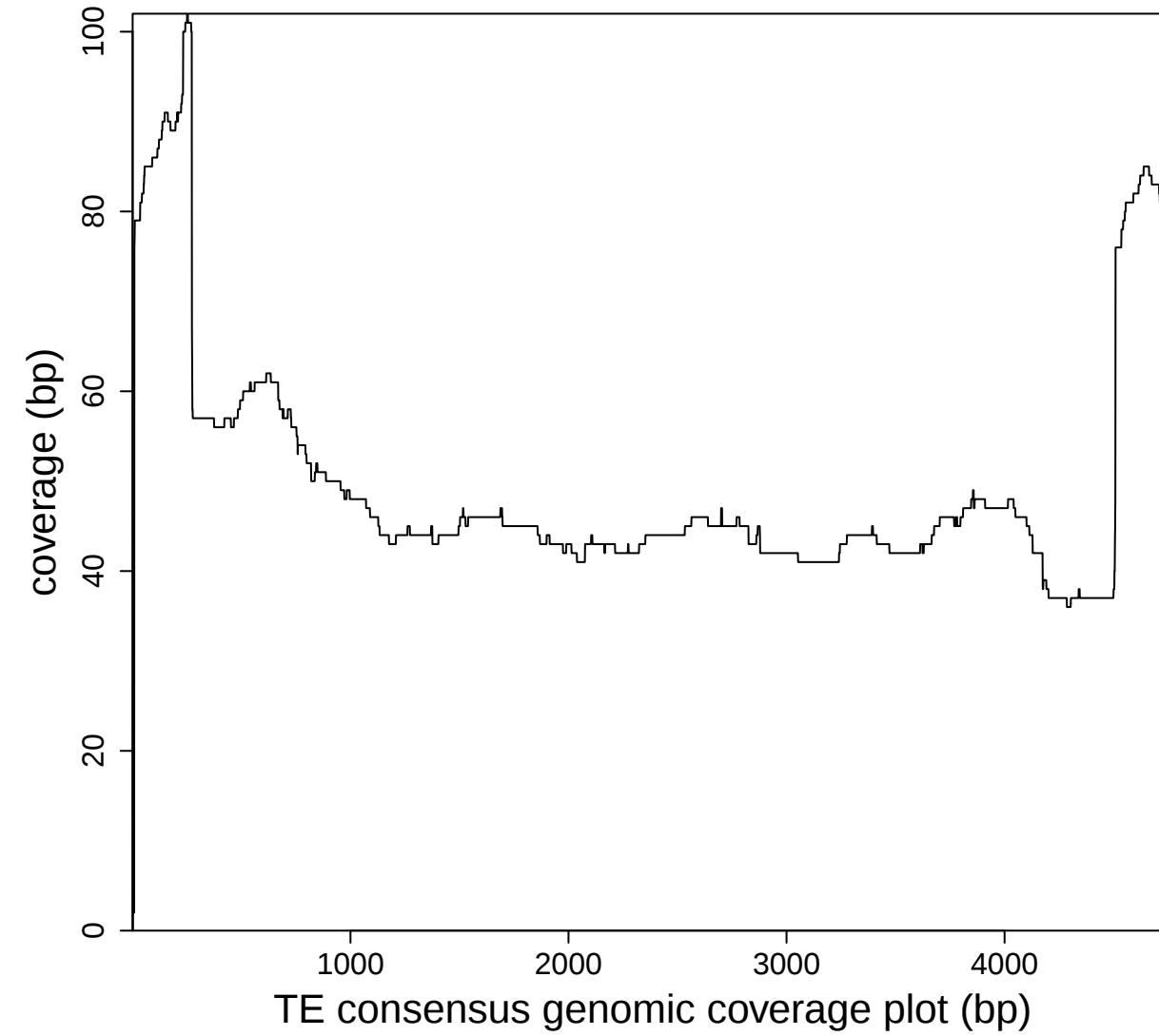
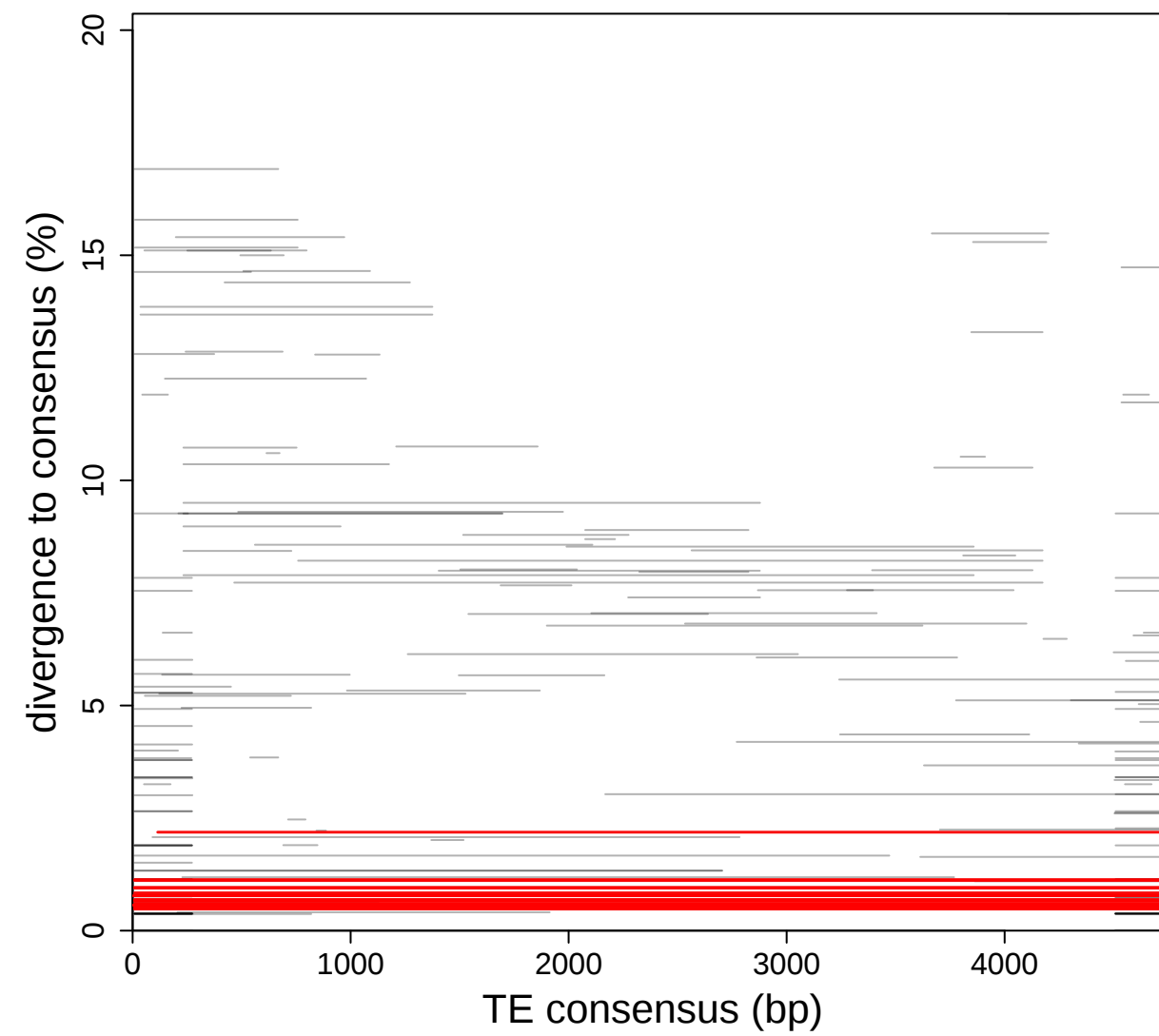






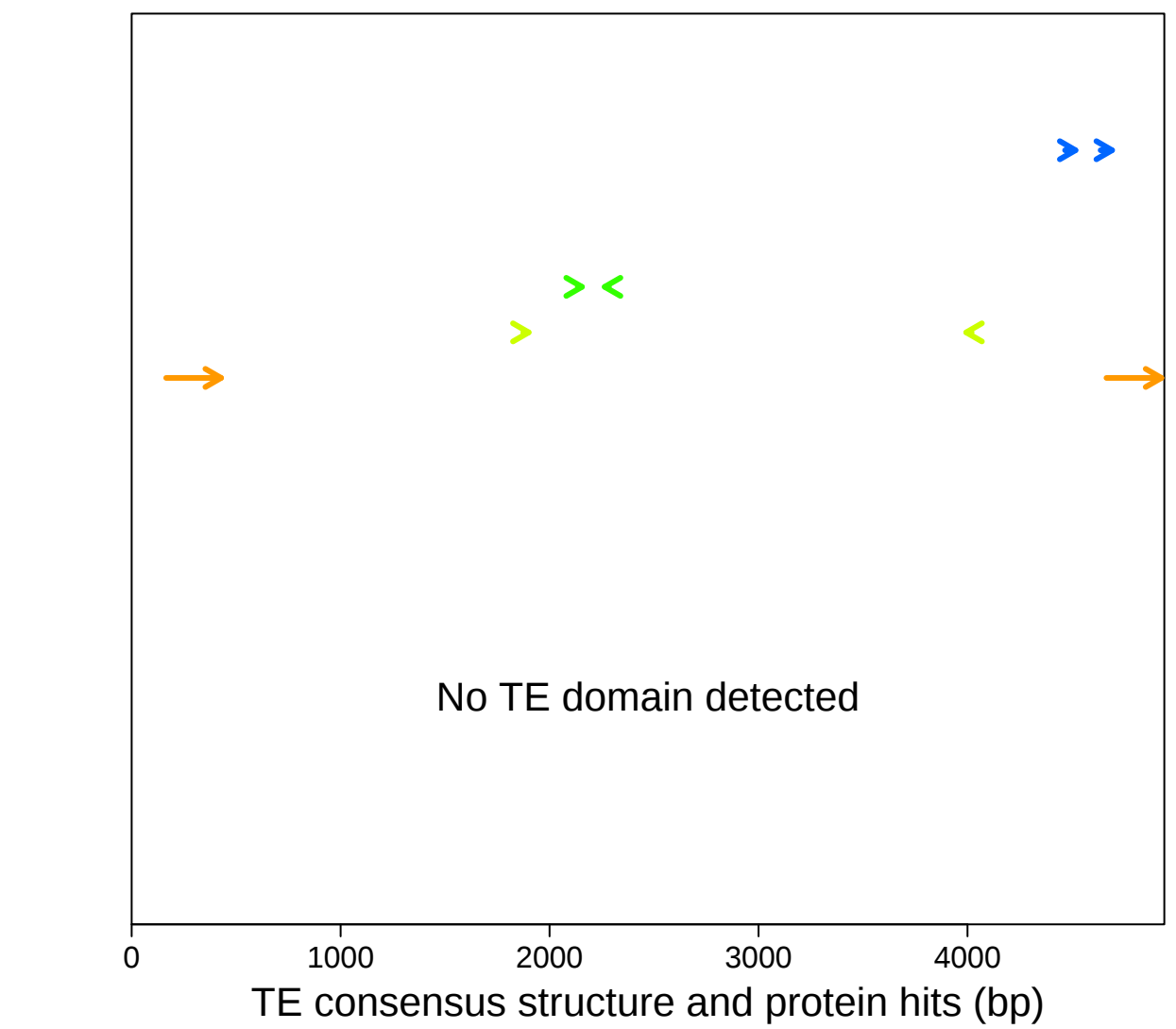
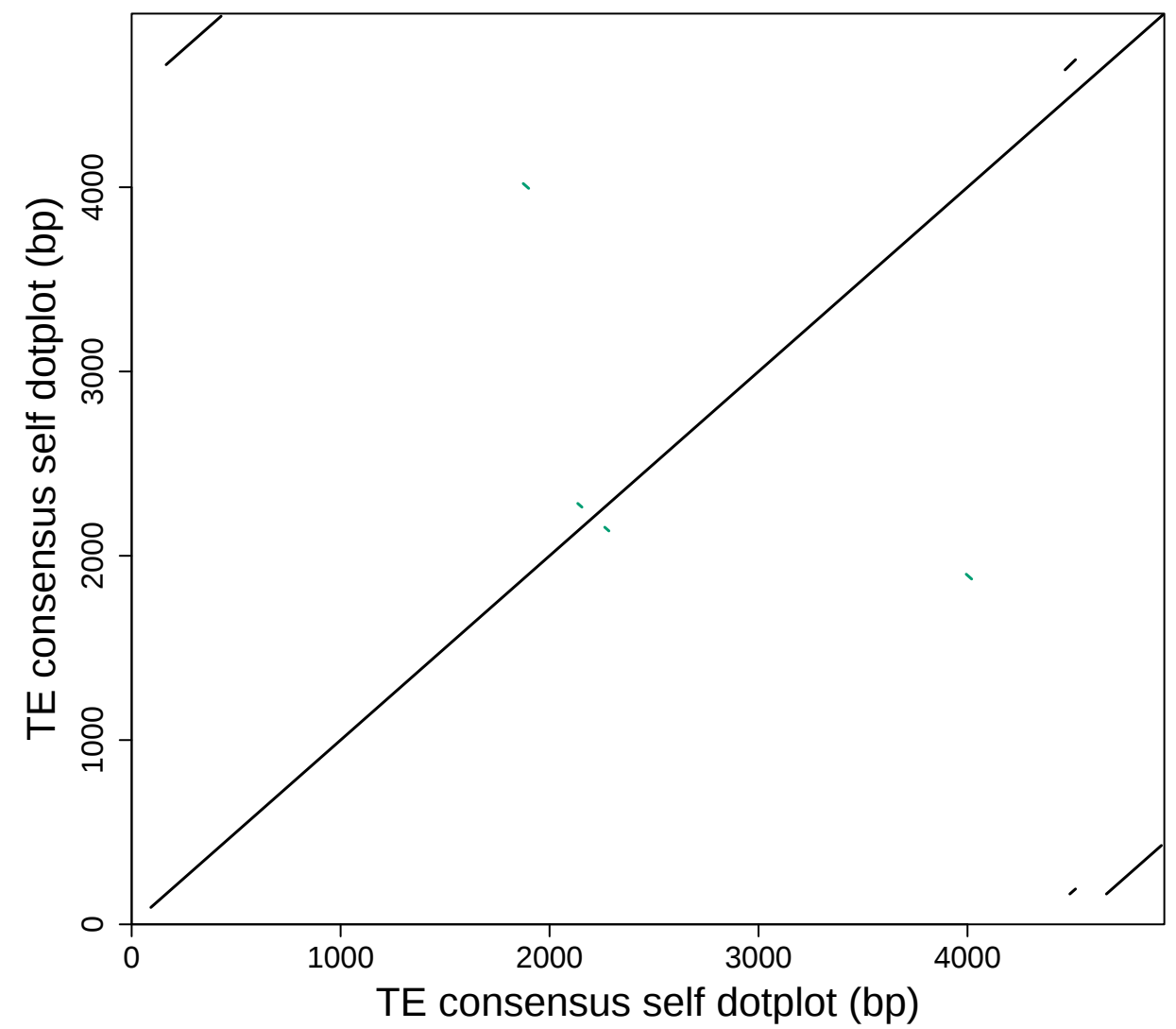
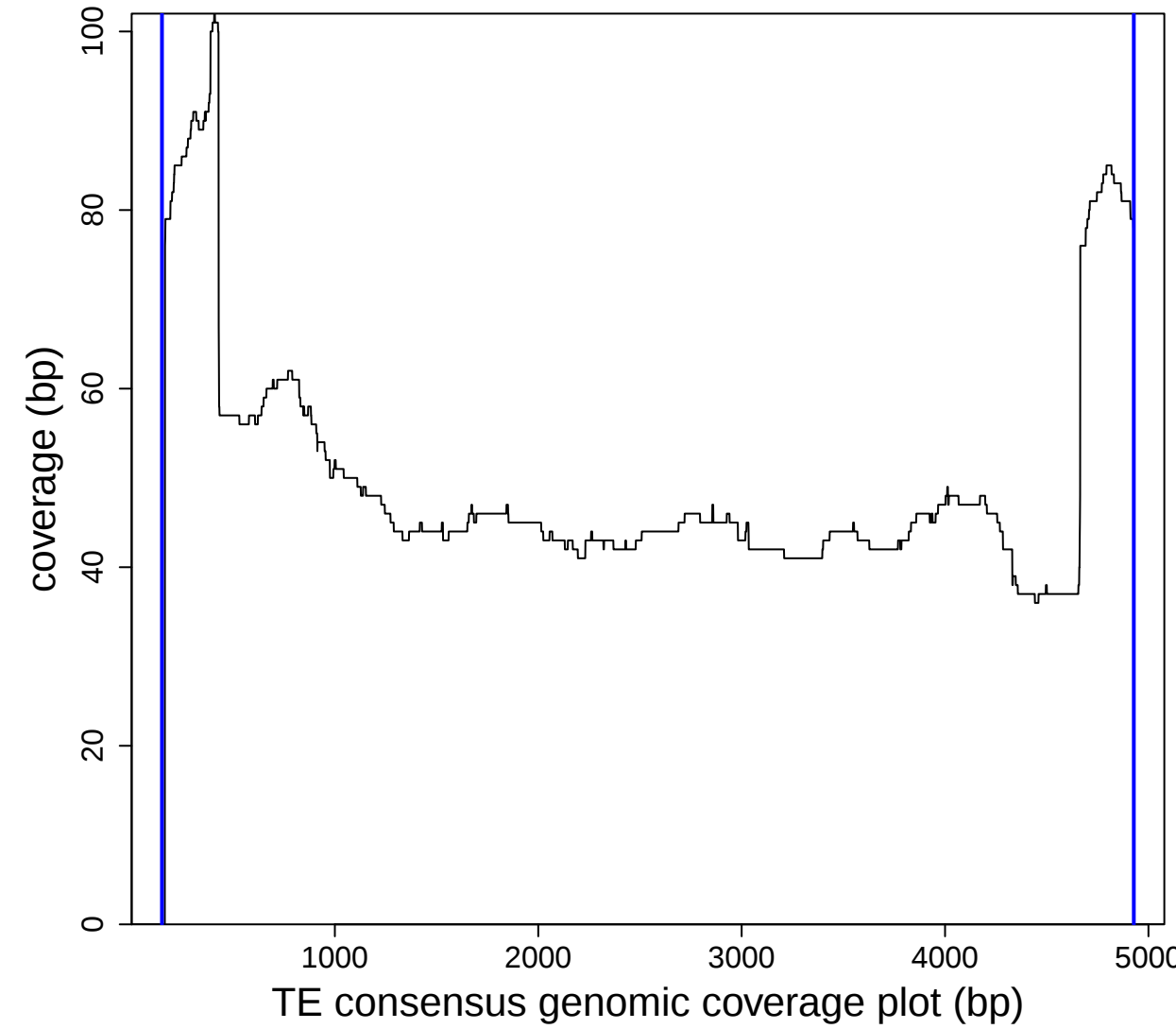
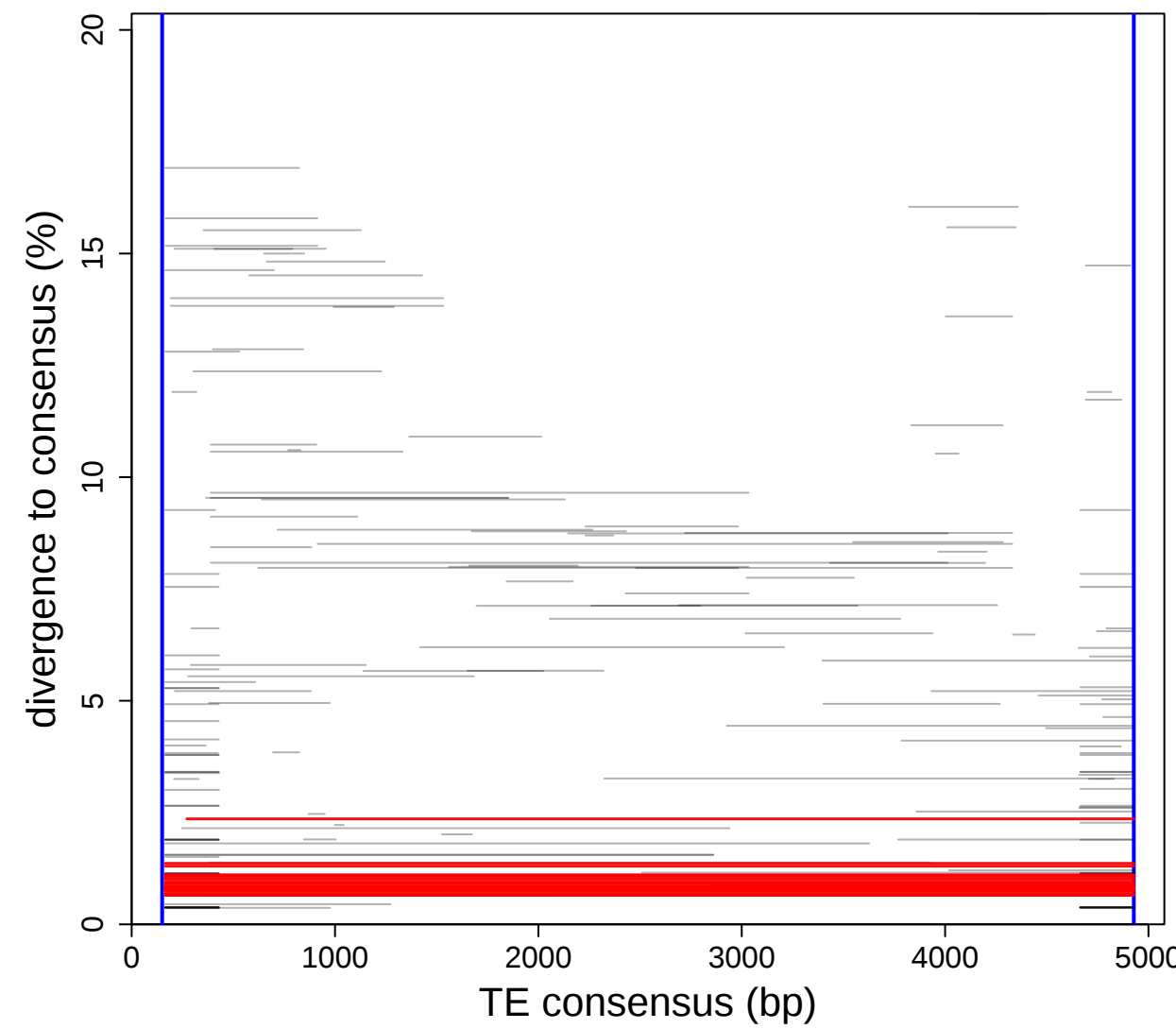
_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcIn.fa_al
size: 4772bp; fragments: 211; full length: 24 (>=4294.8bp)

After TEtrimmer 4772 bp

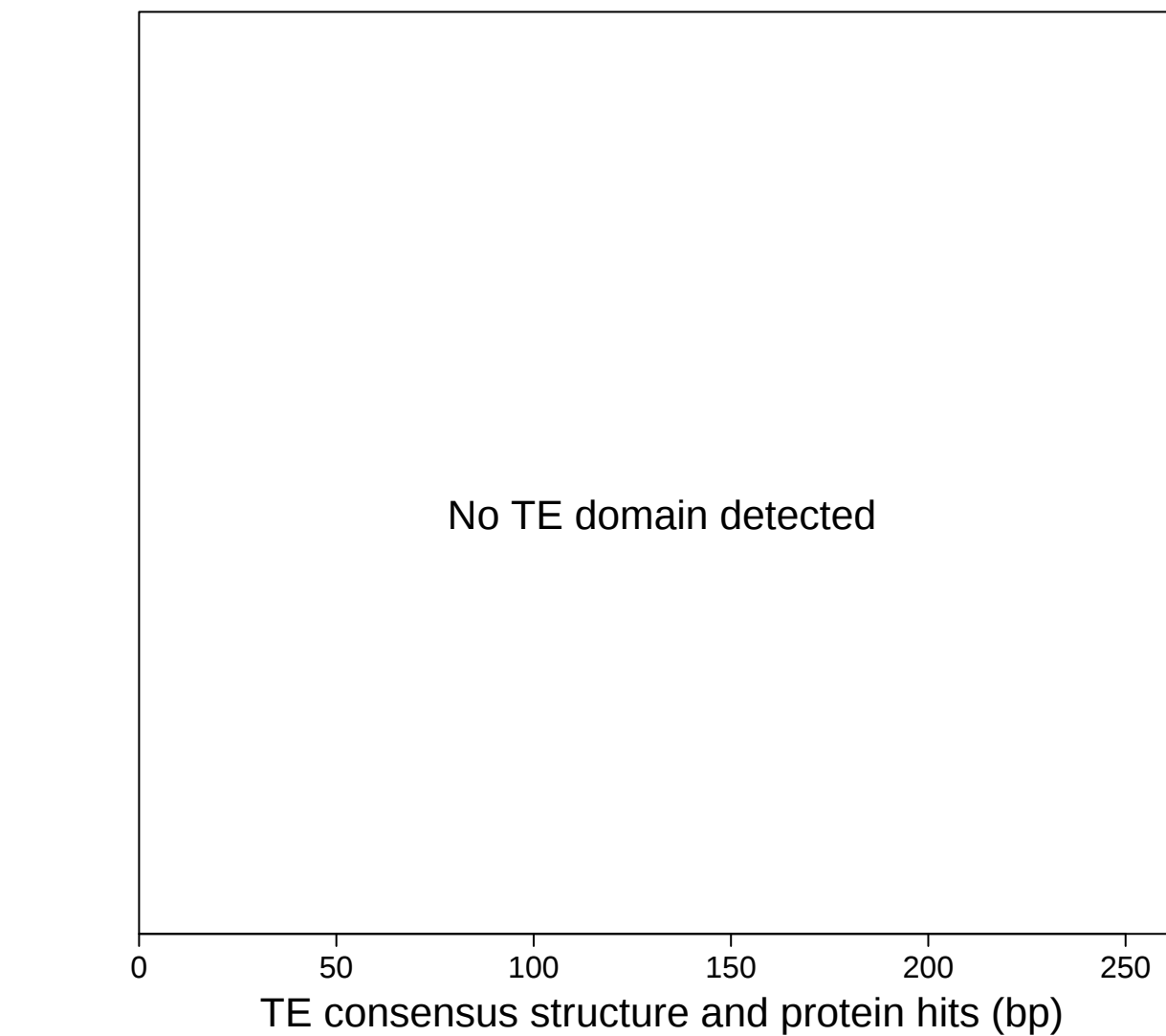
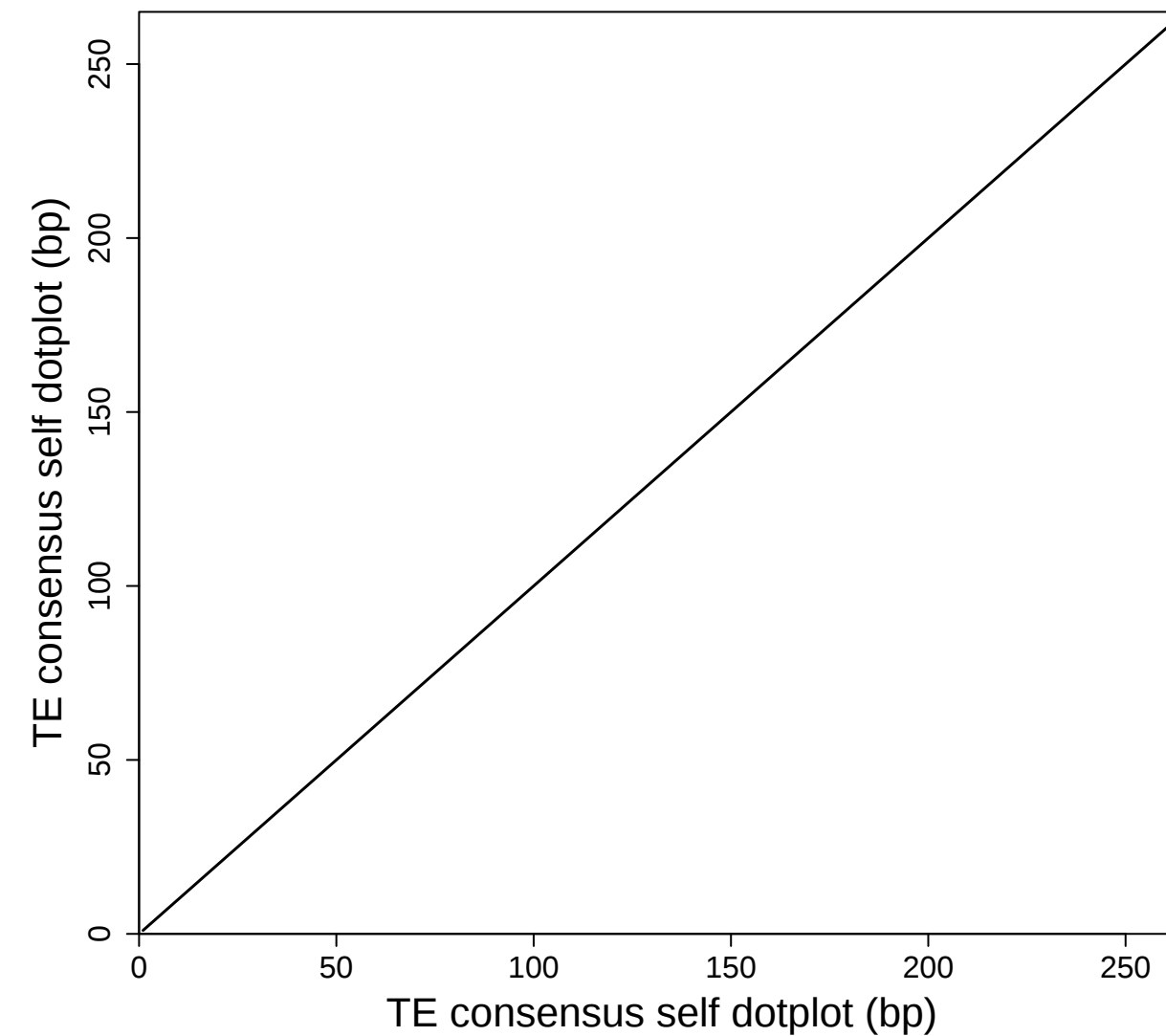
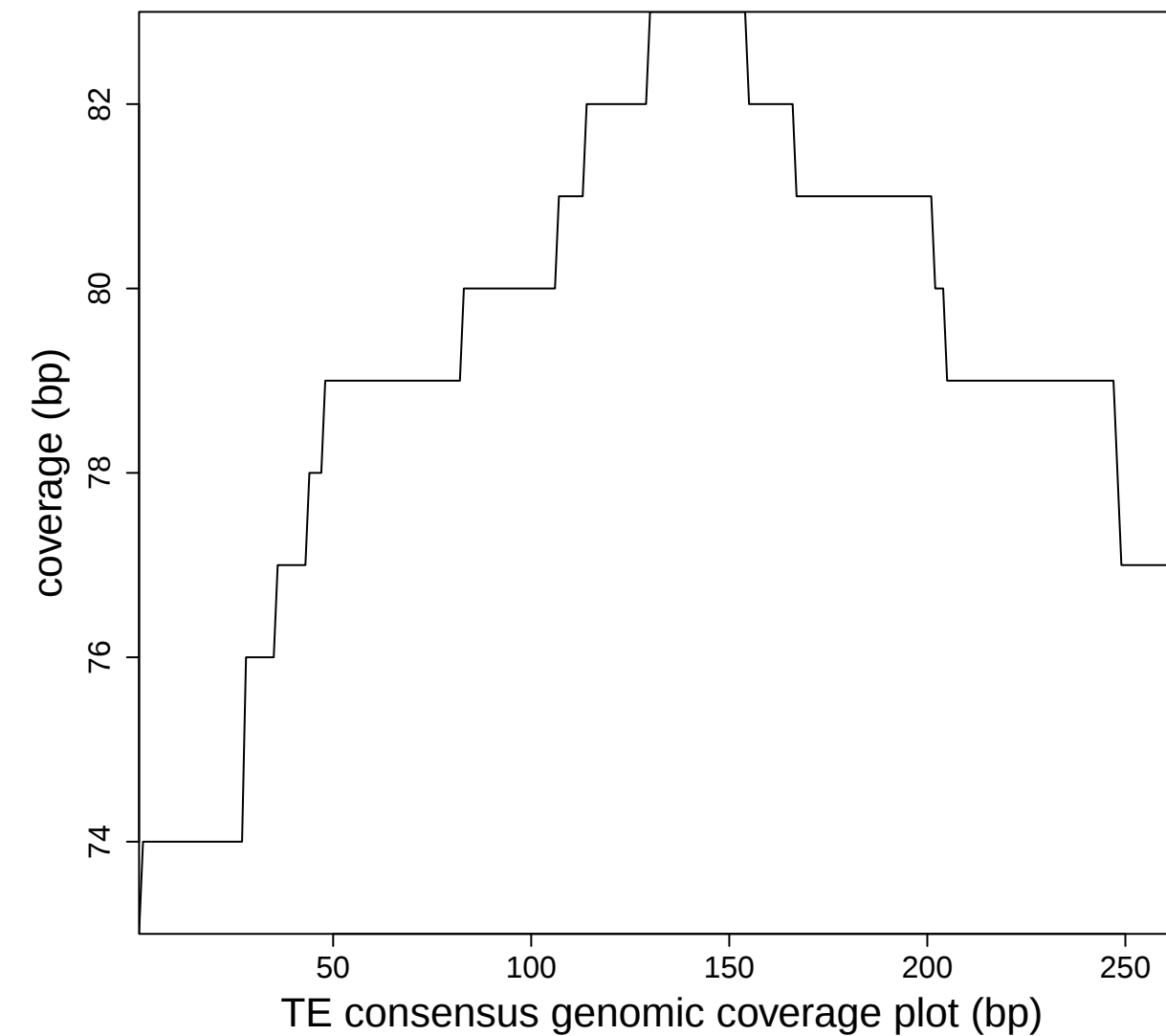
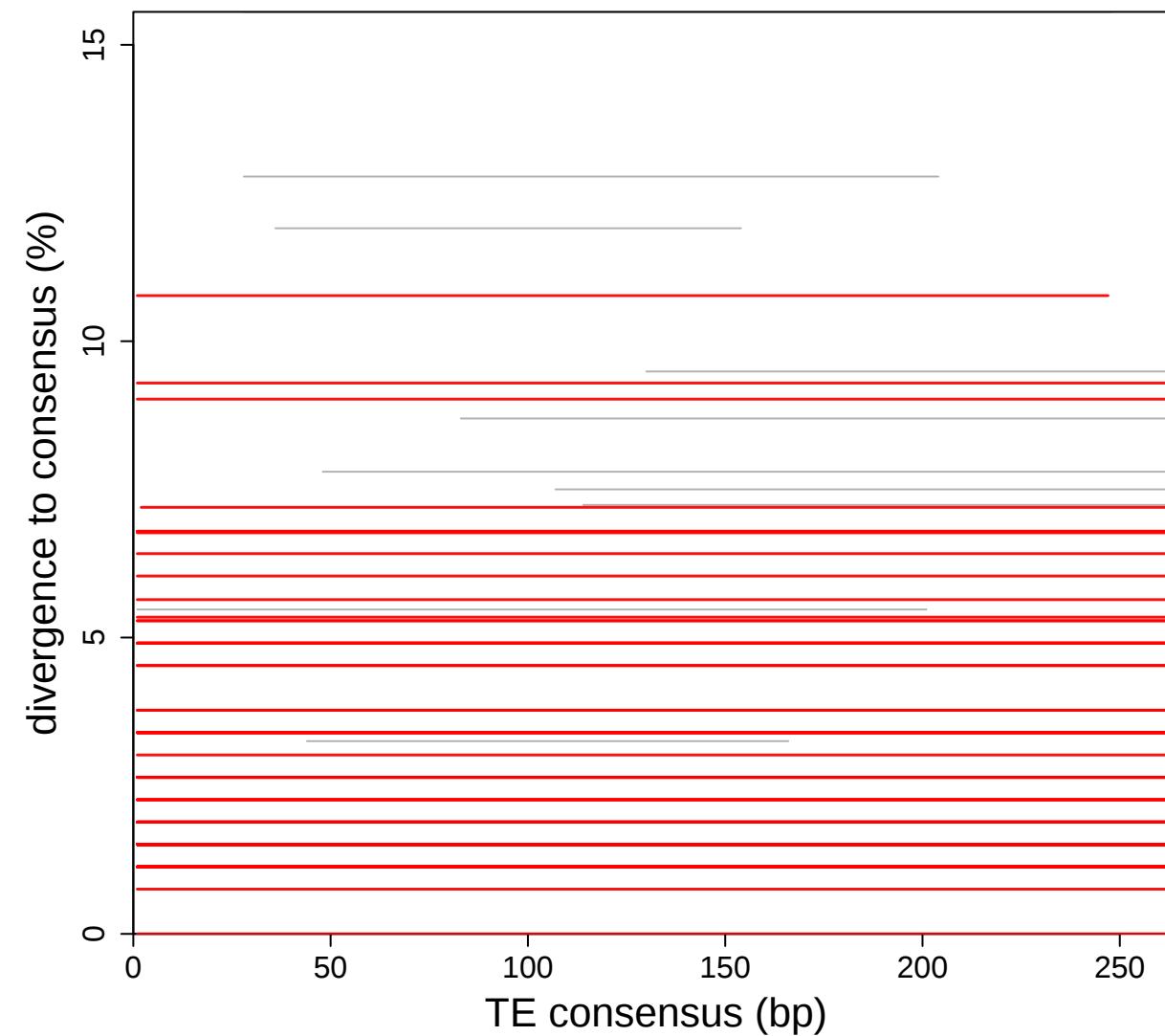


bed_g_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bclr
size: 5078bp; fragments: 211; full length: 24 (≥ 4570.2 bp)

After TEtrimmer Extended plot Blue lines are boundaries

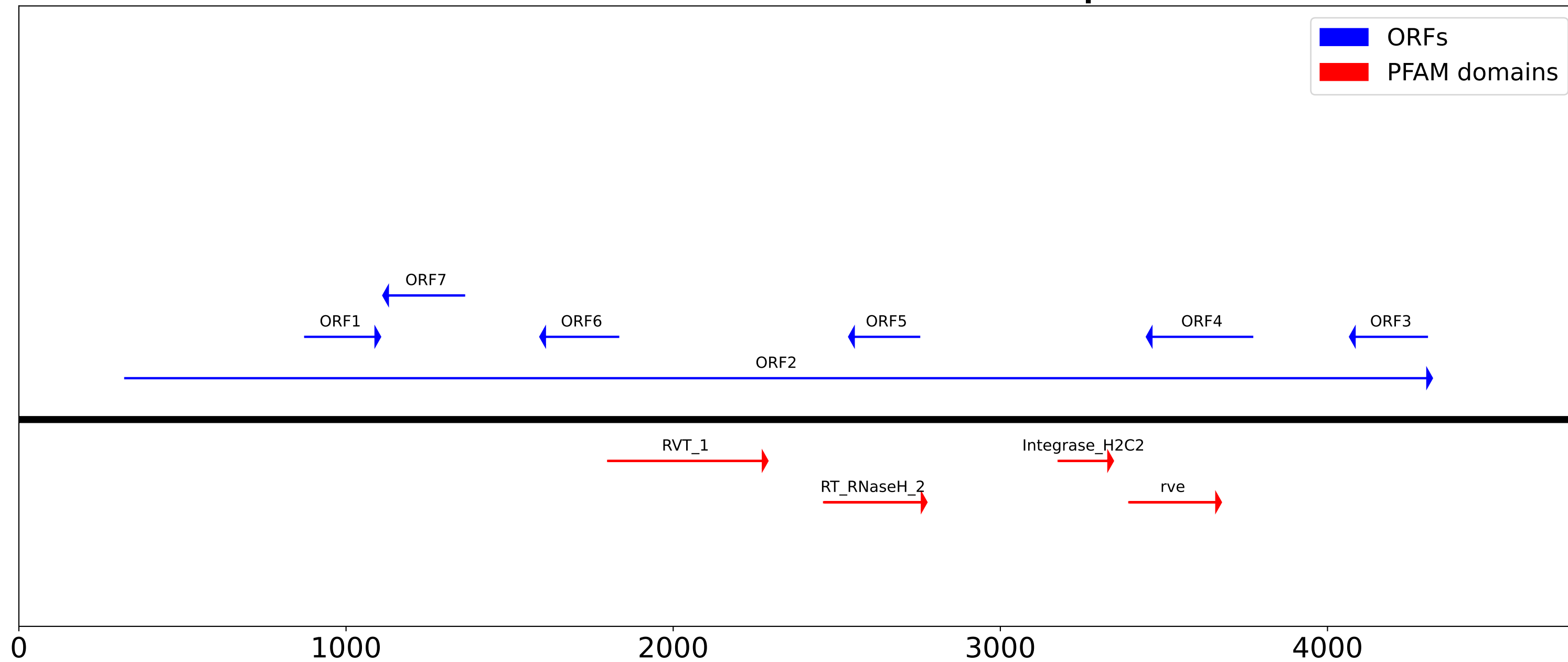


TE: ltr_1_family_142
size: 265bp; fragments: 83; full length: 73 (≥ 238.5 bp)



Before TEtrimmer 265 bp

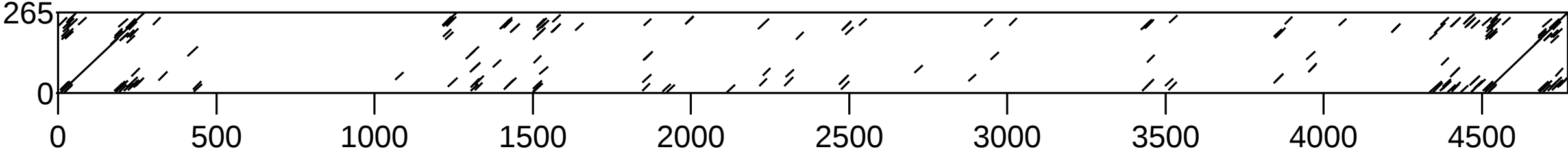
After TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)