

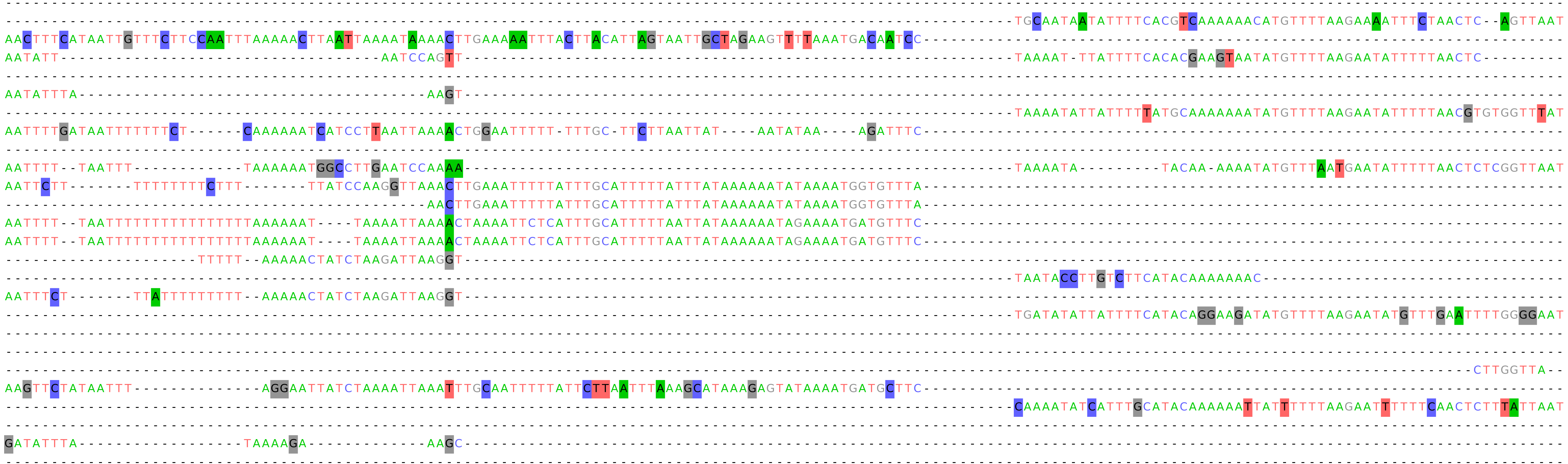
Start crop Point

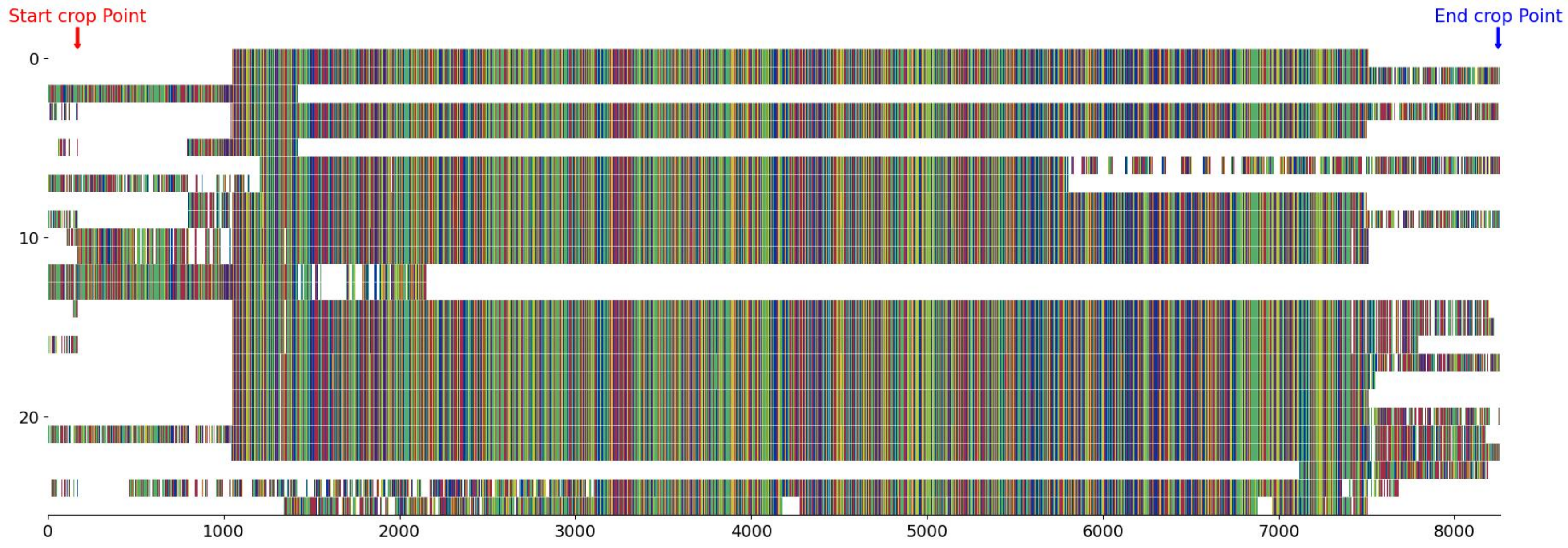


End crop Point



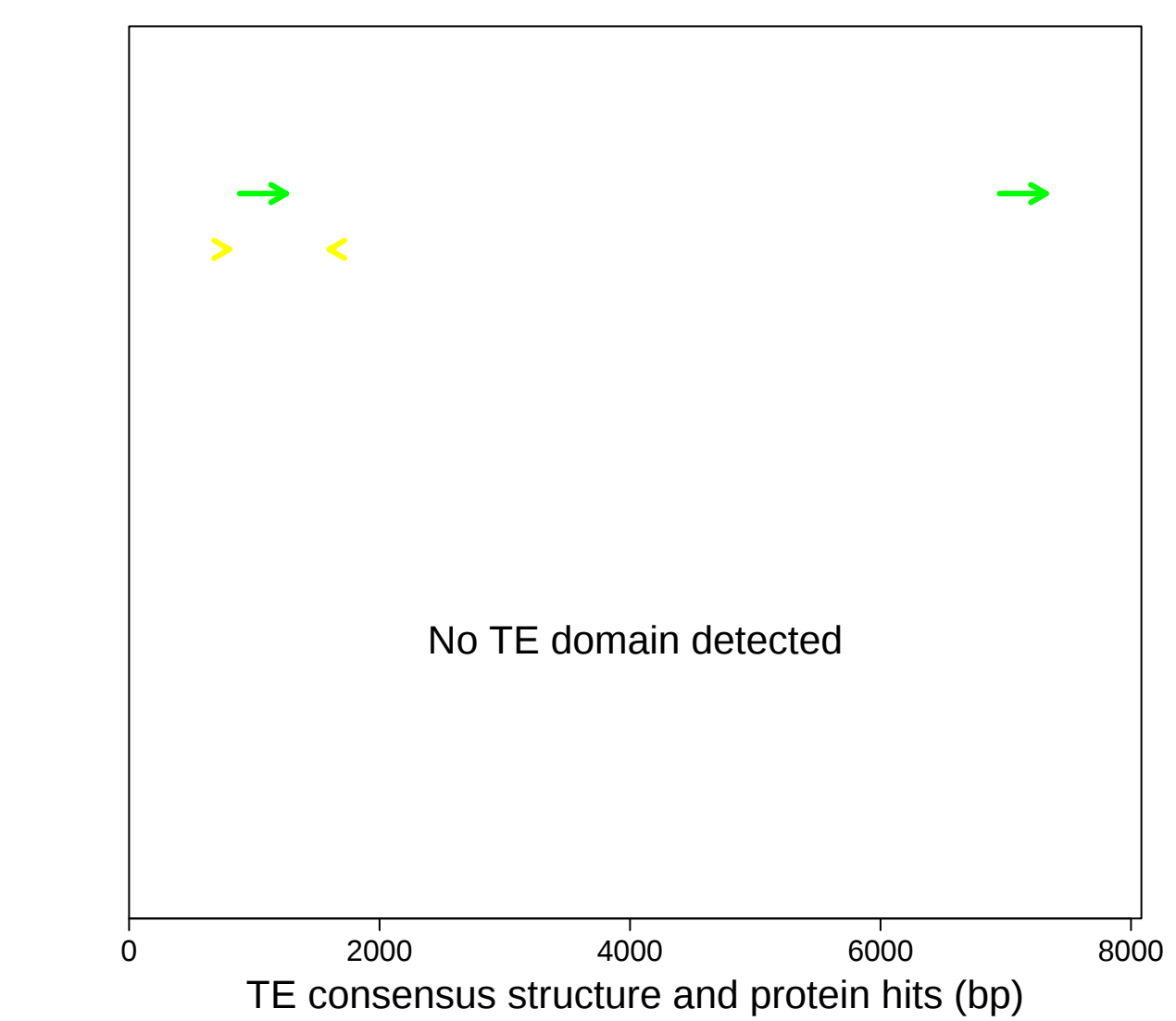
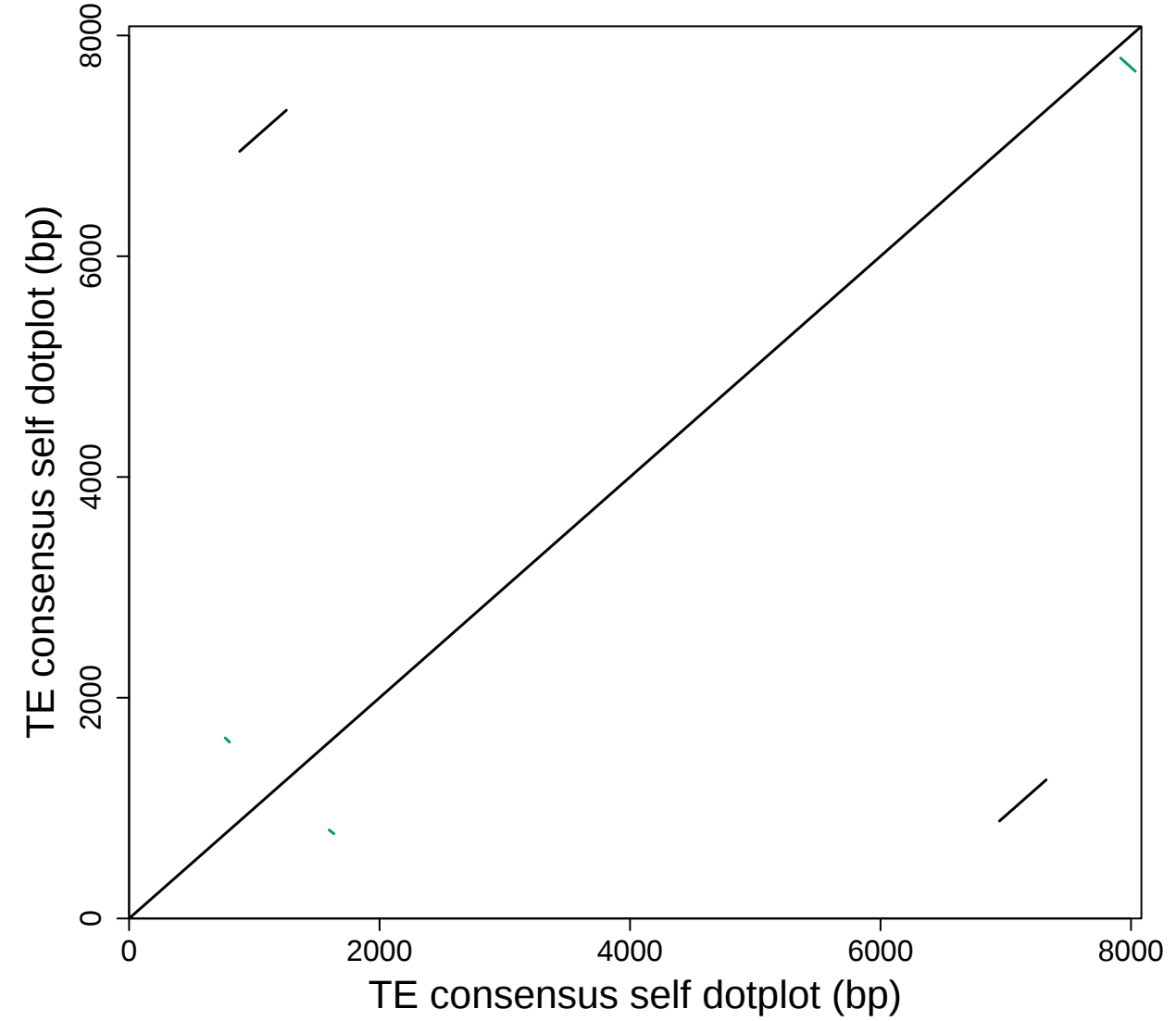
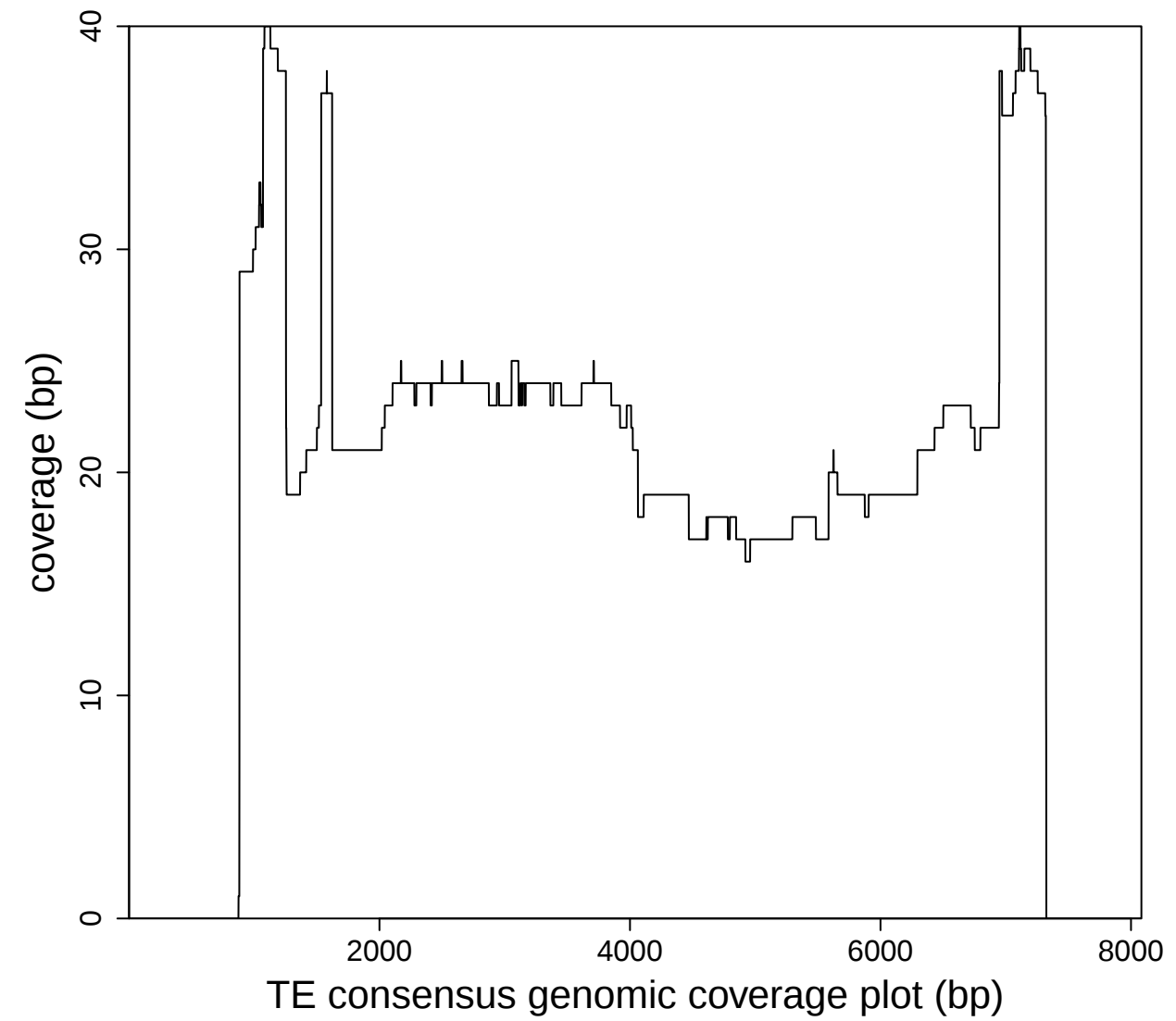
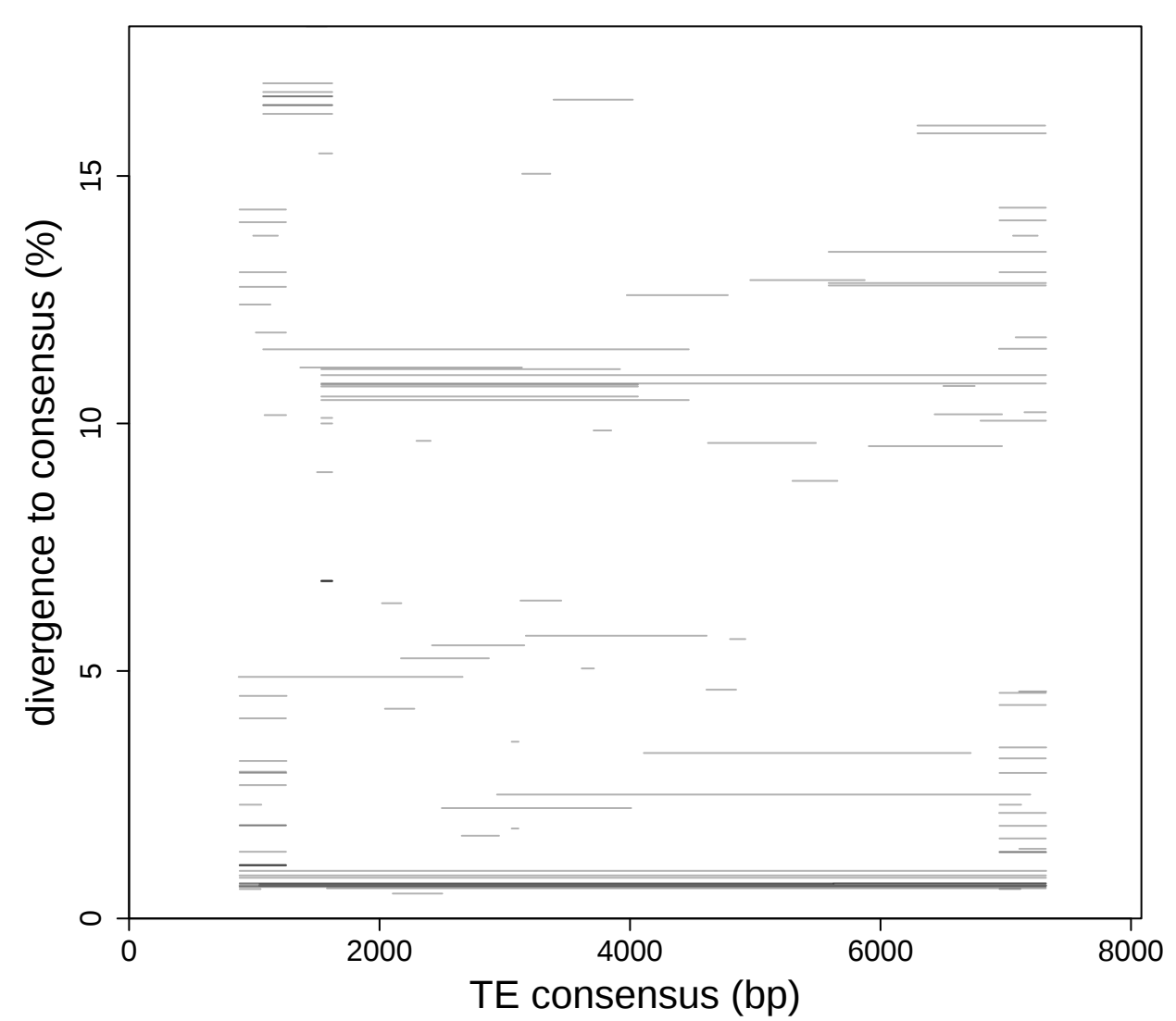
MSA length = 8083





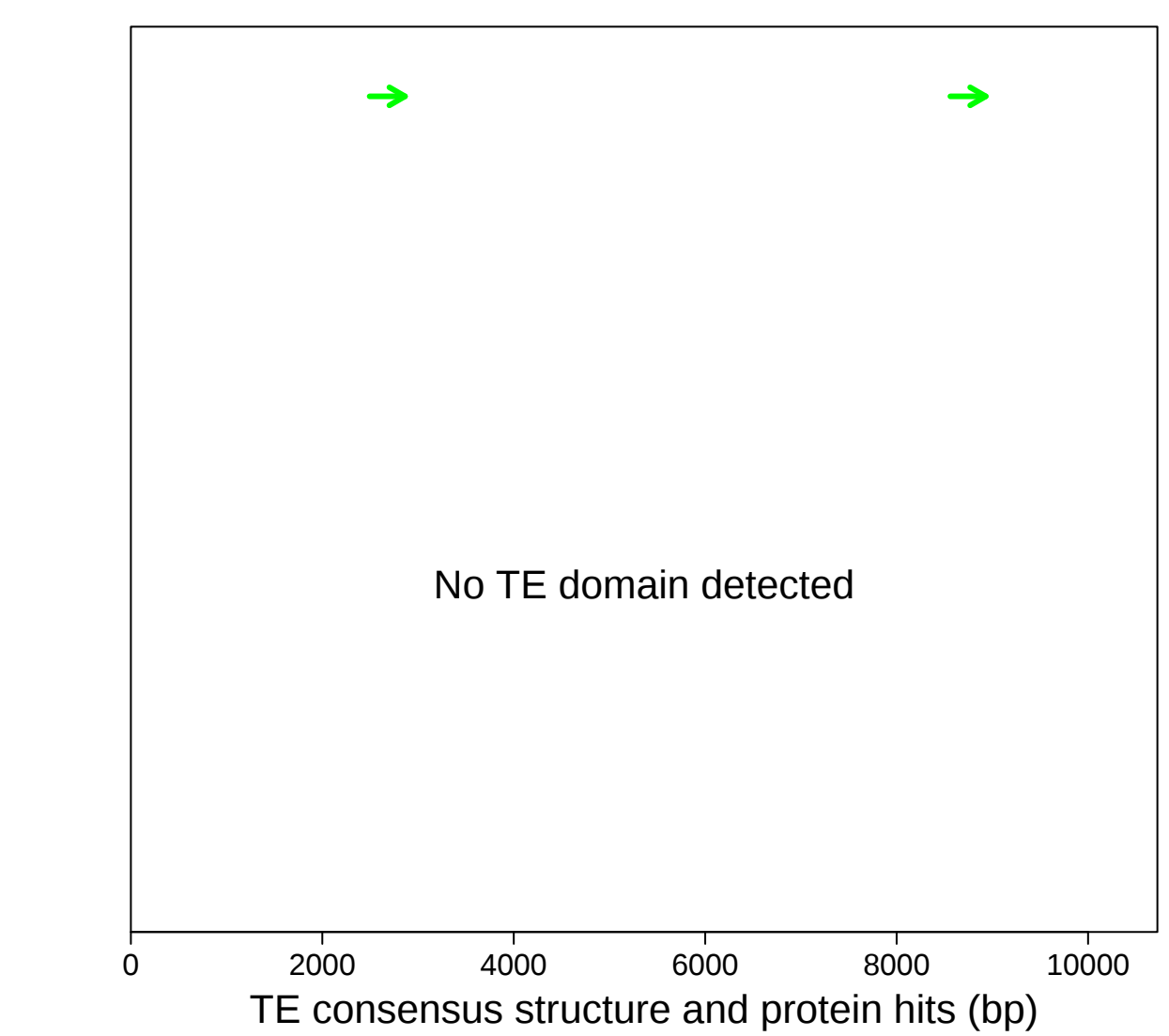
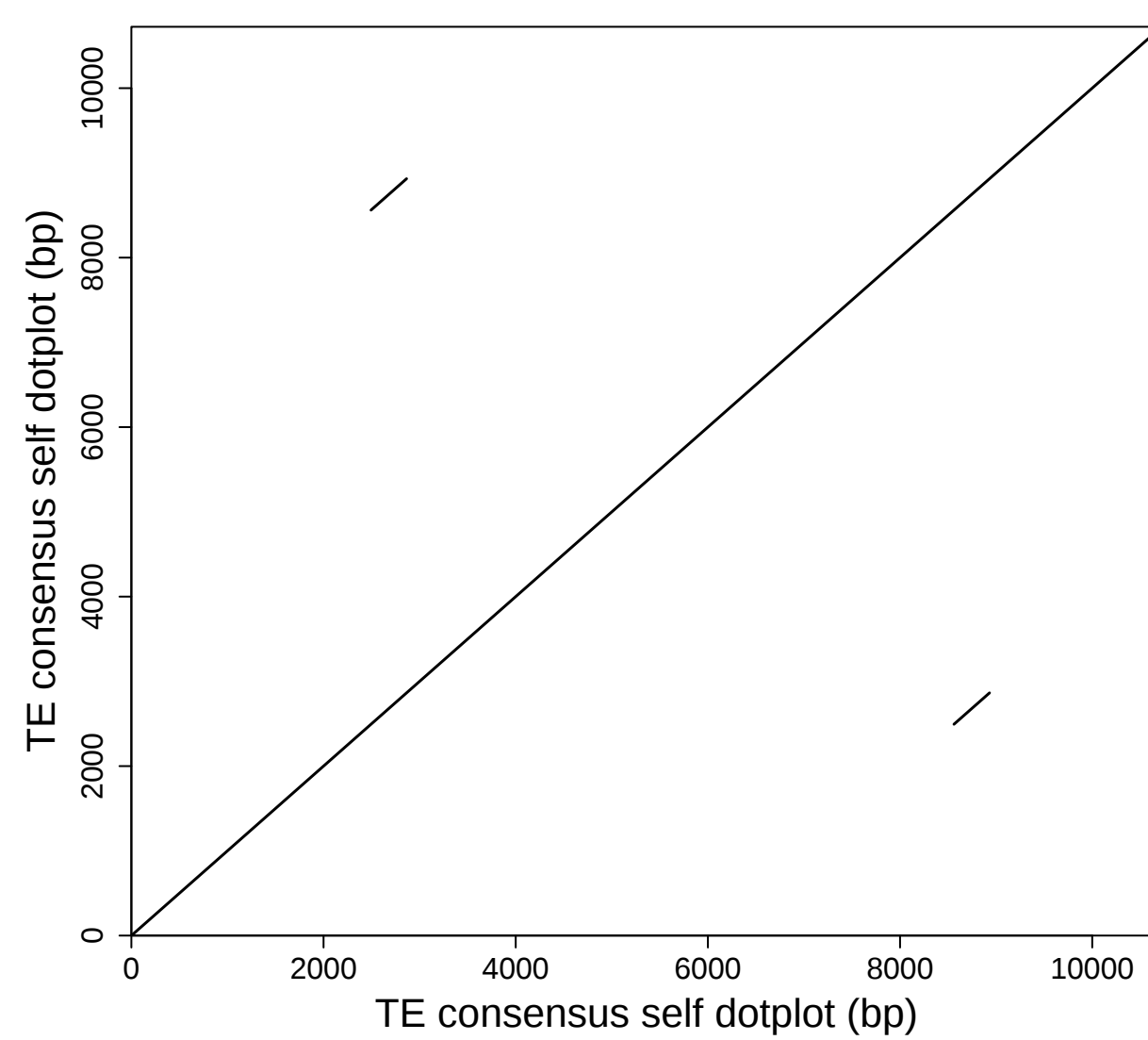
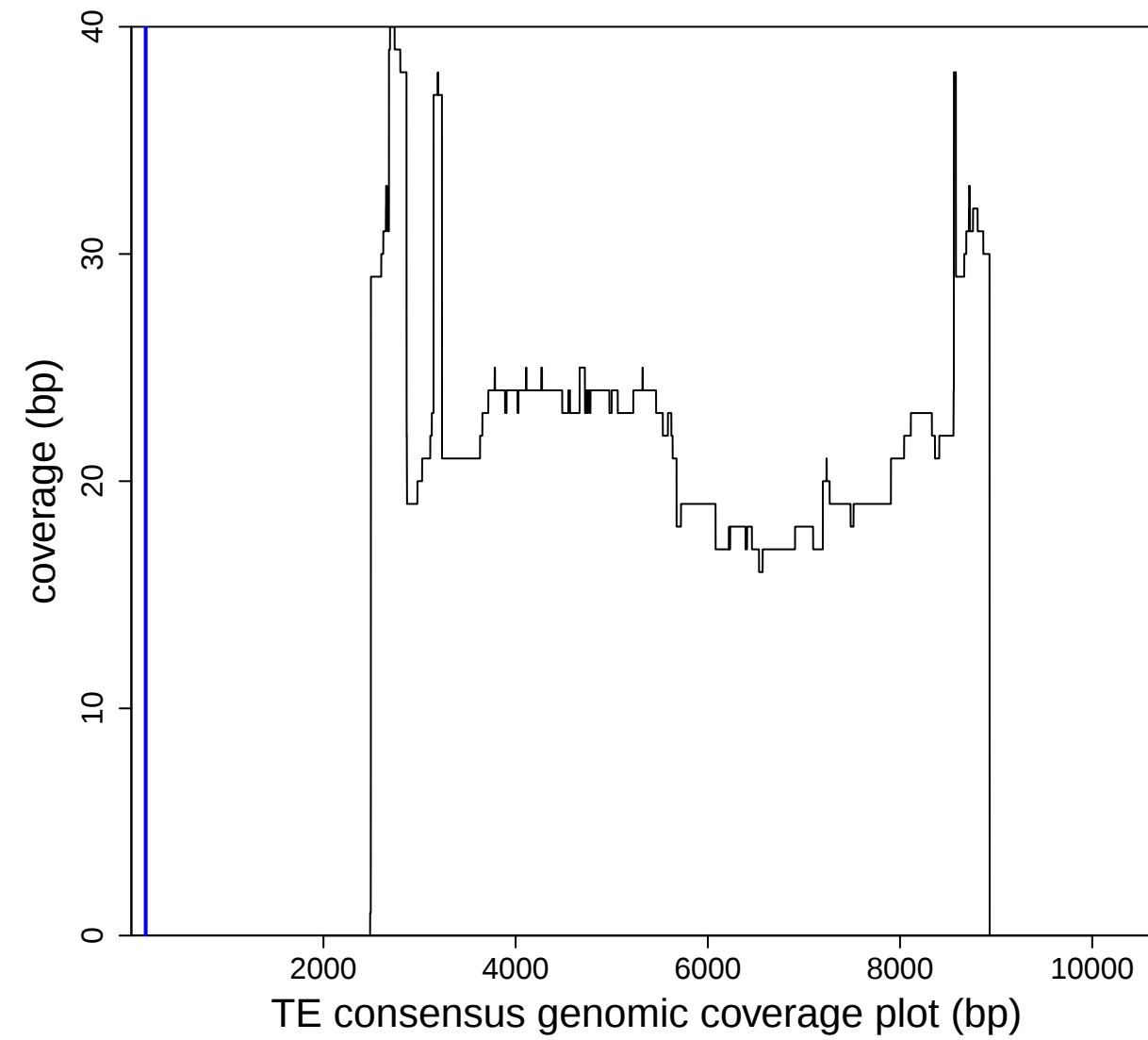
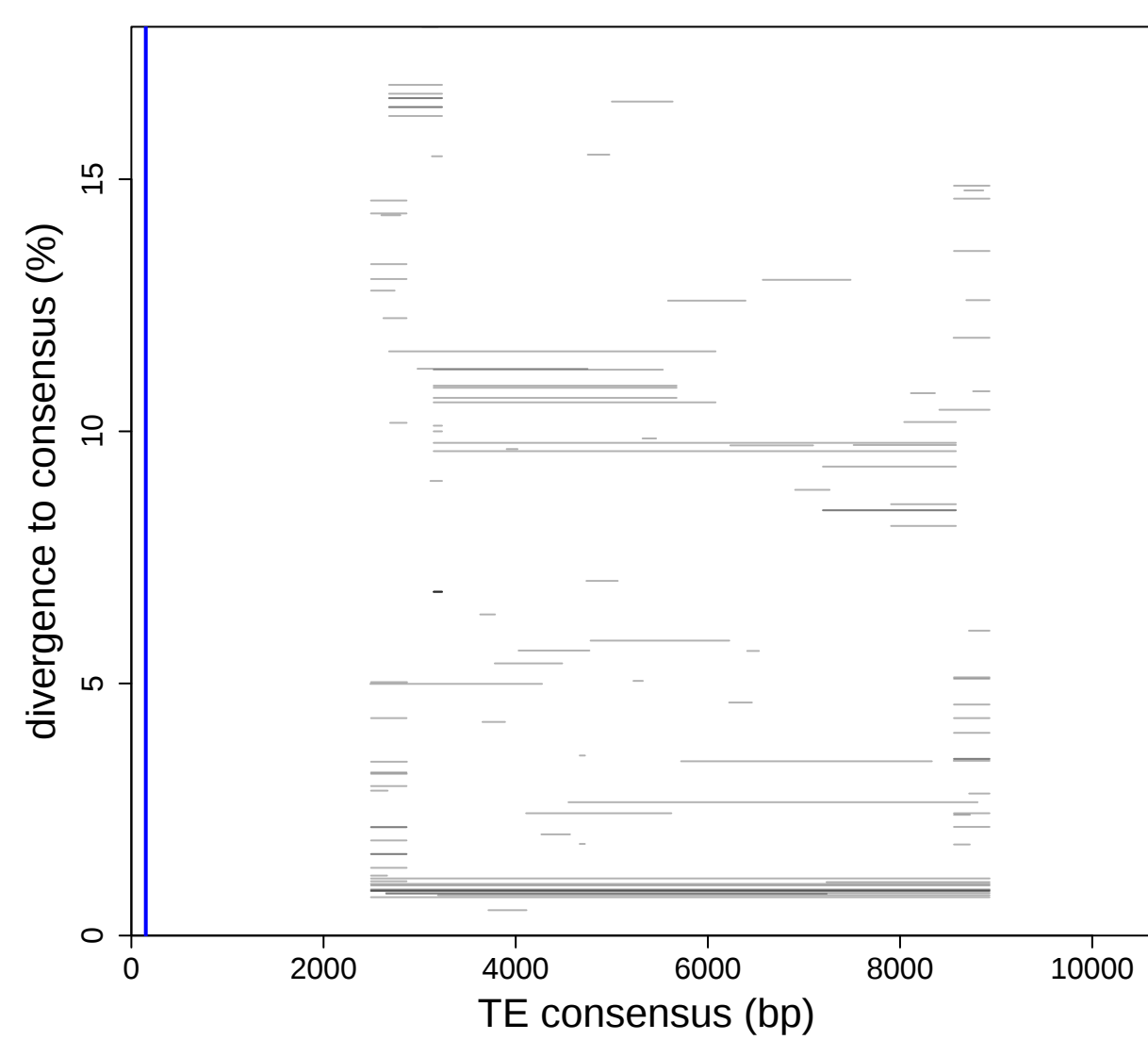
_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcIn.fa_al
size: 8083bp; fragments: 116; full length: 0 (>=7274.7bp)

After TEtrimmer 8083 bp

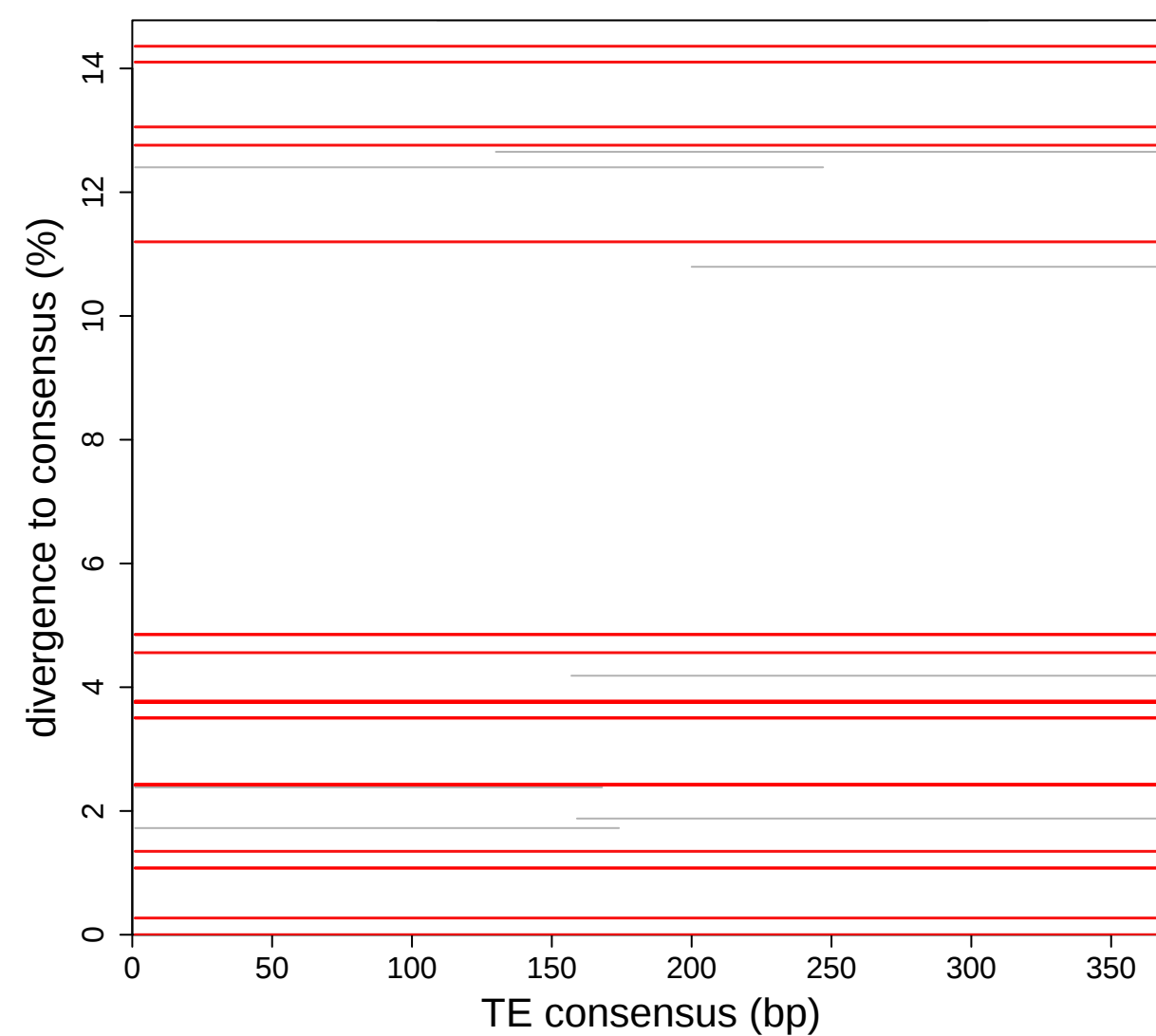


bed_g_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bclr
size: 10724bp; fragments: 116; full length: 0 (>=9651.6bp)

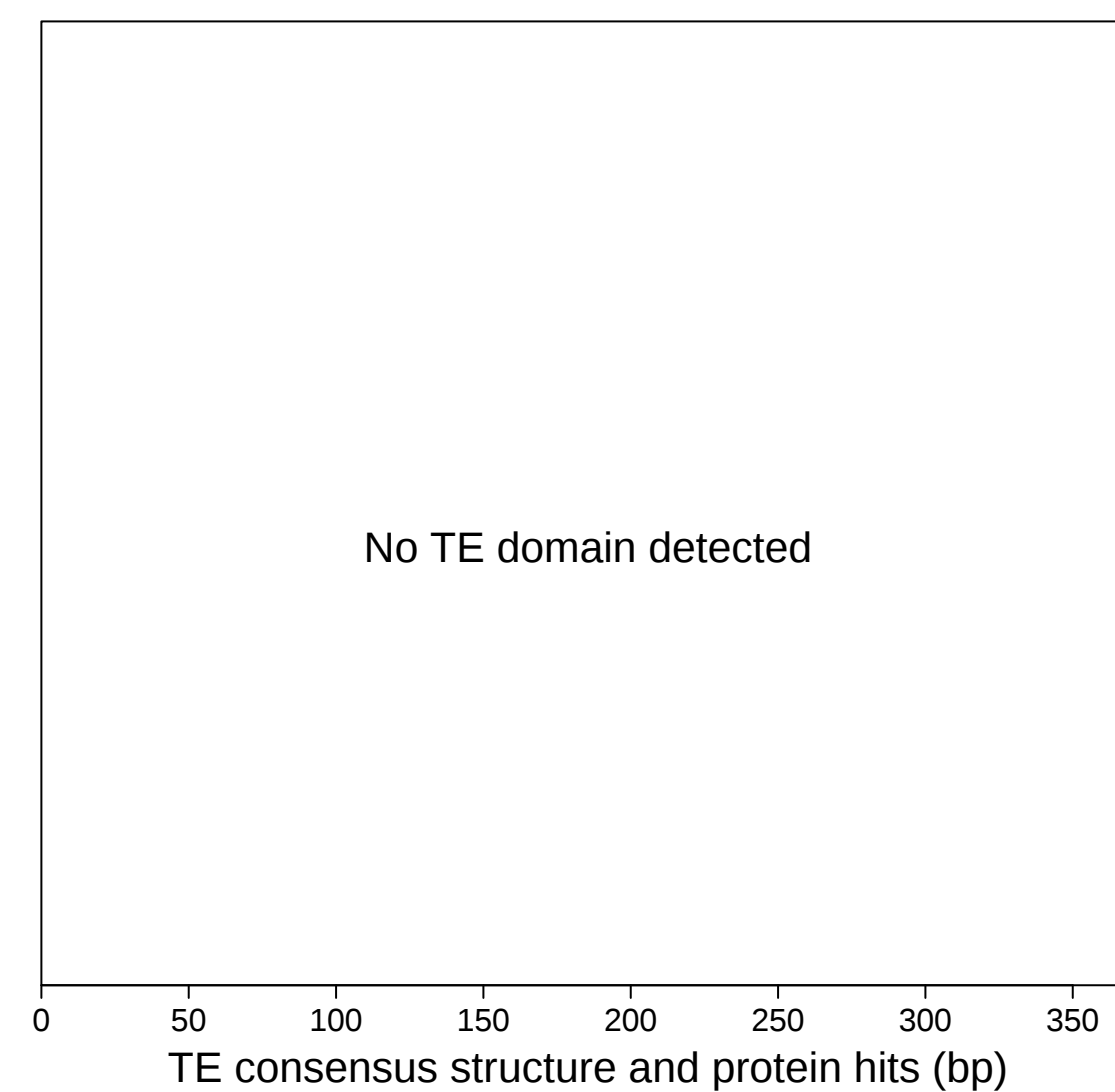
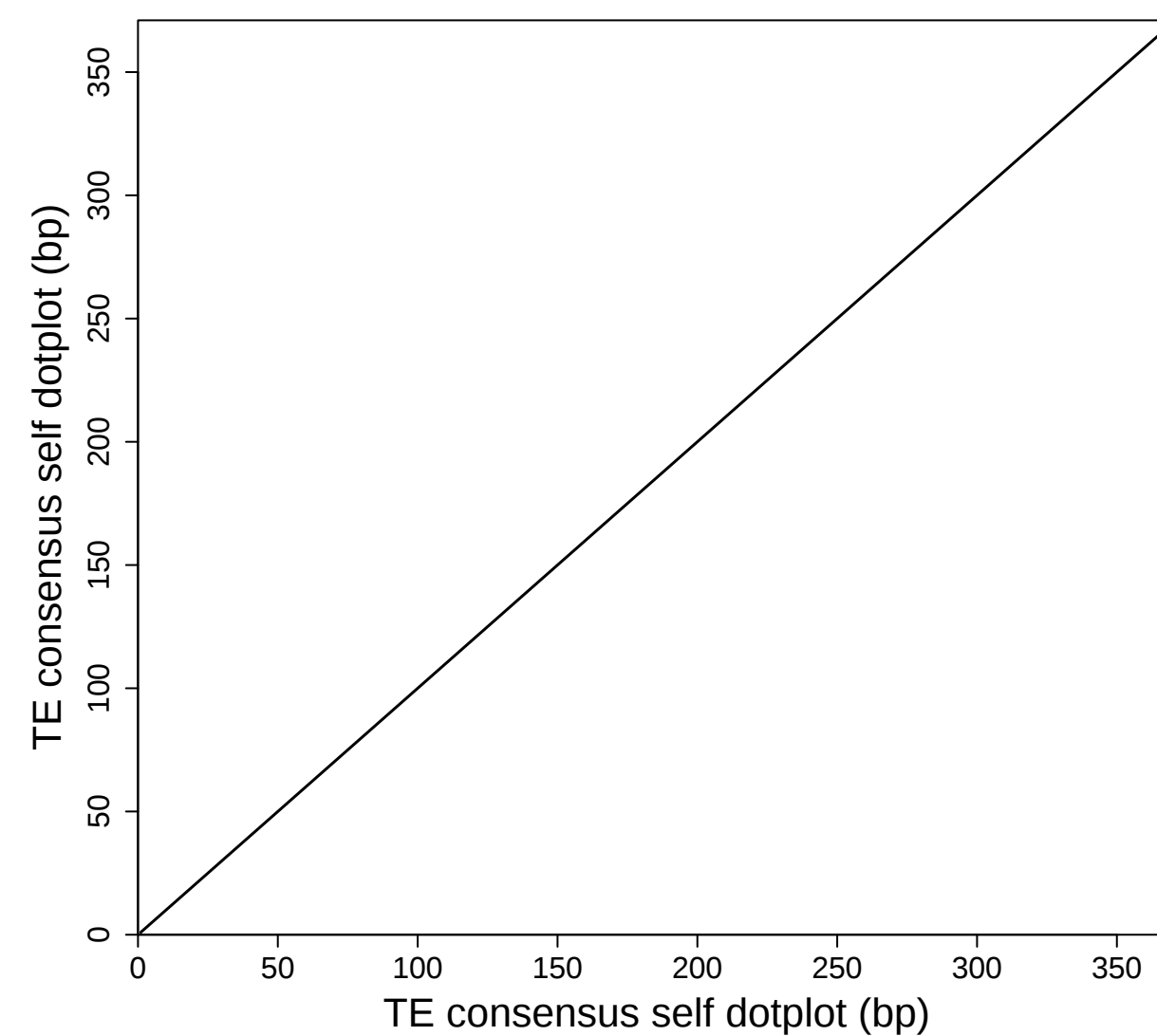
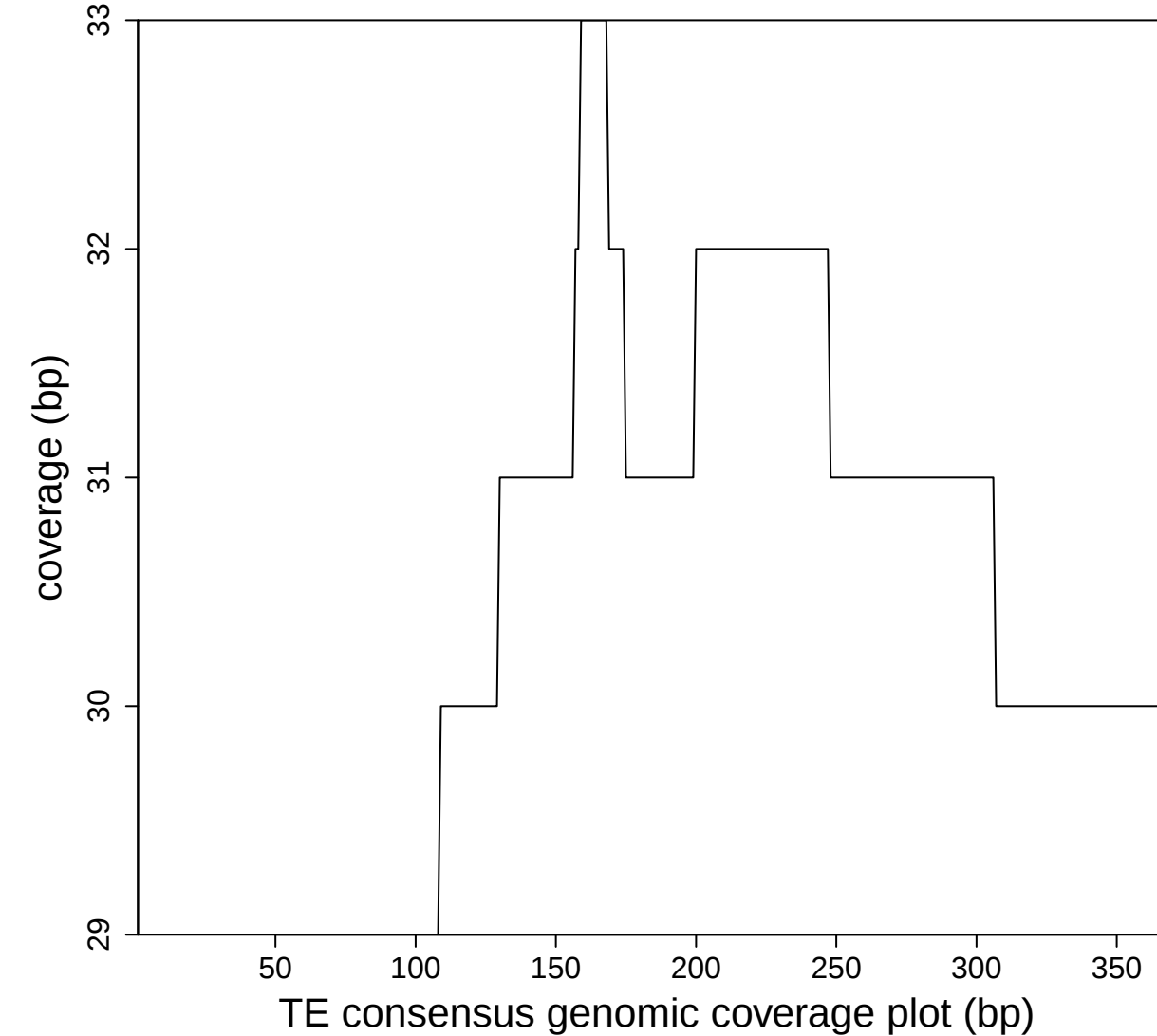
After TEtrimmer Extended plot Blue lines are boundaries



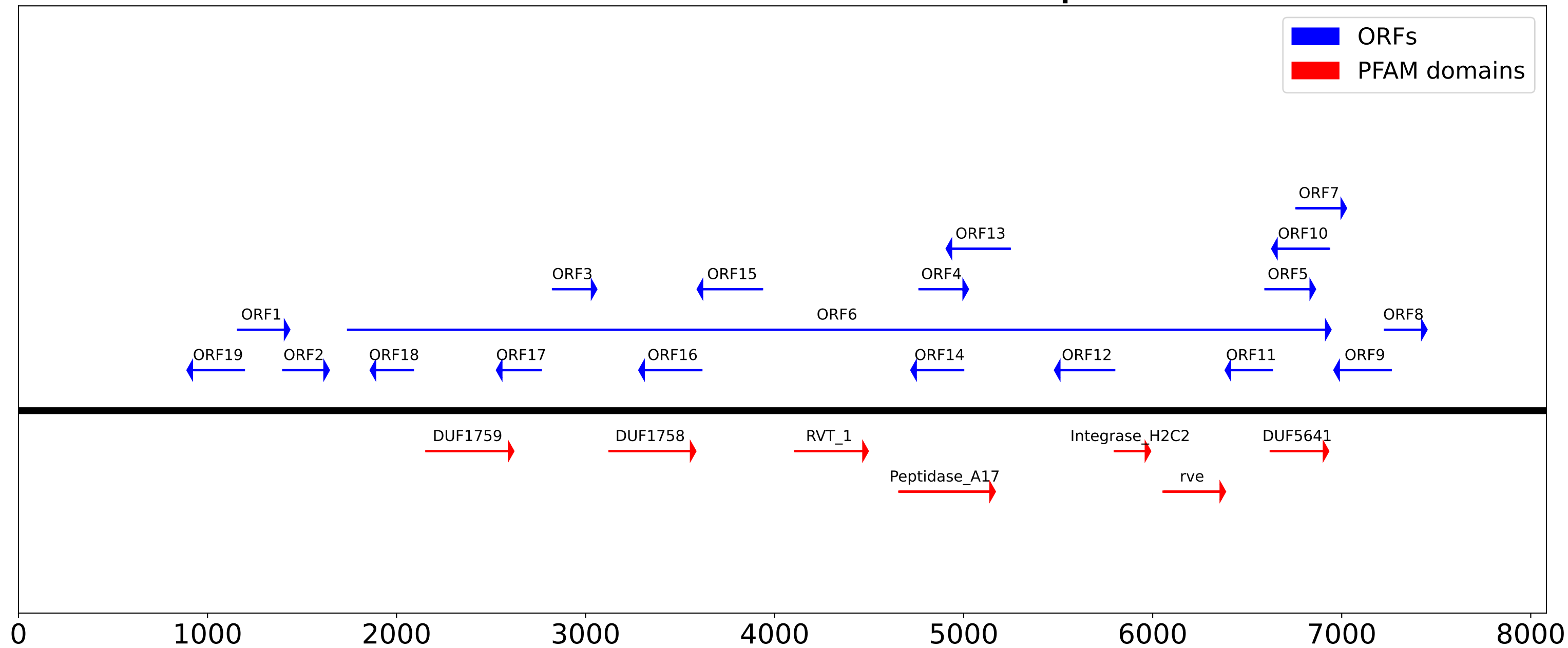
TE: ltr_1_family_415
size: 371bp; fragments: 34; full length: 26 (≥ 333.9 bp)



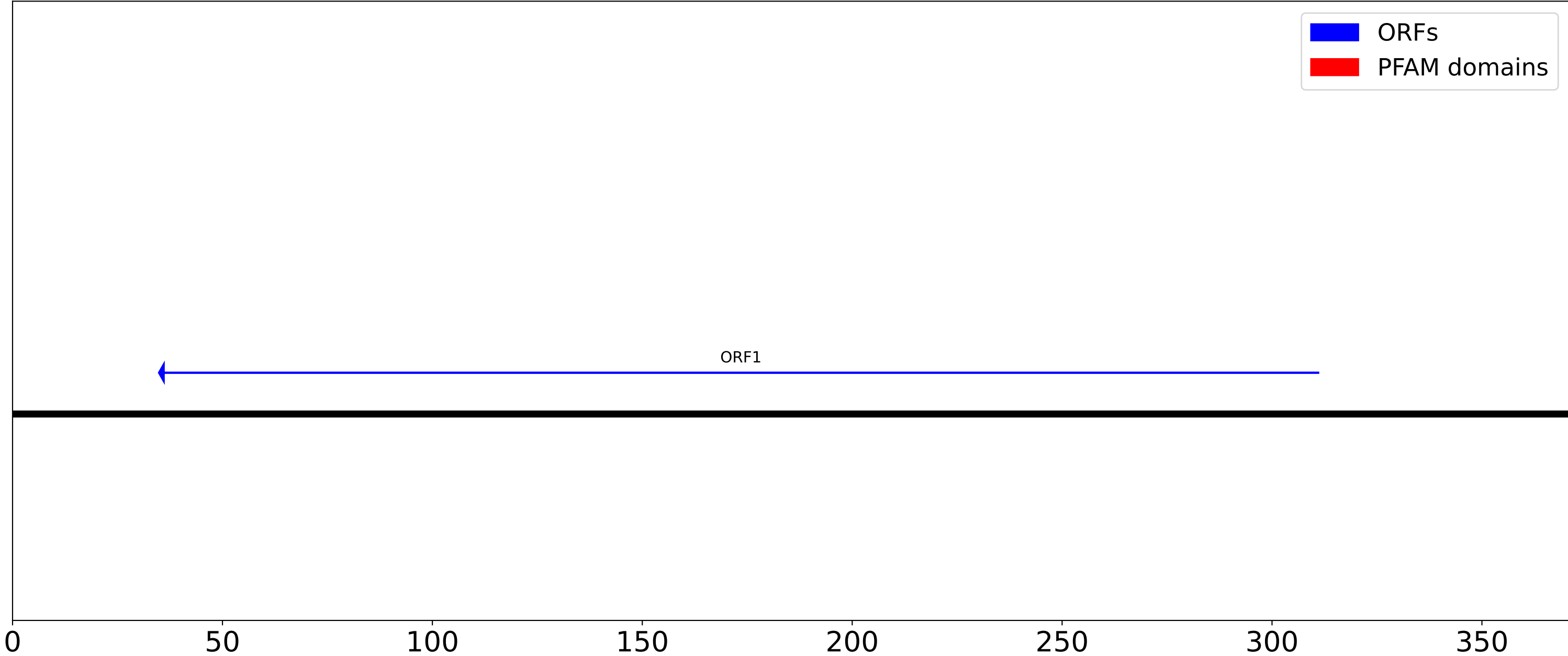
Before TEtrimmer 371 bp



After TEtrimmer ORF and PFAM domain plot



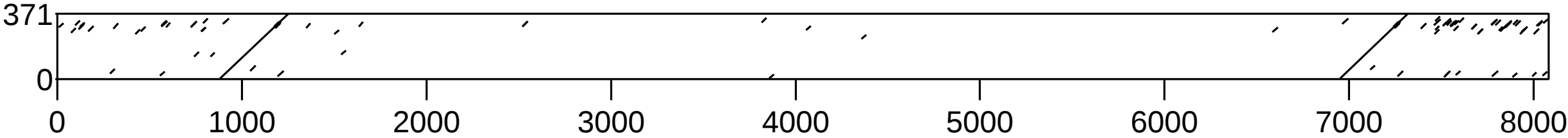
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

nsus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)