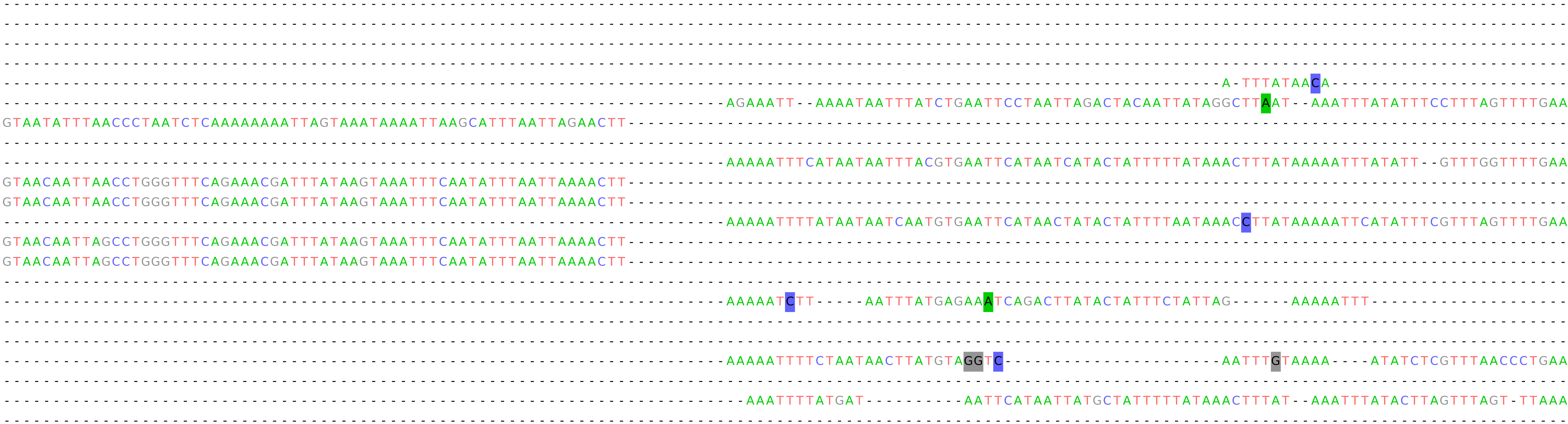


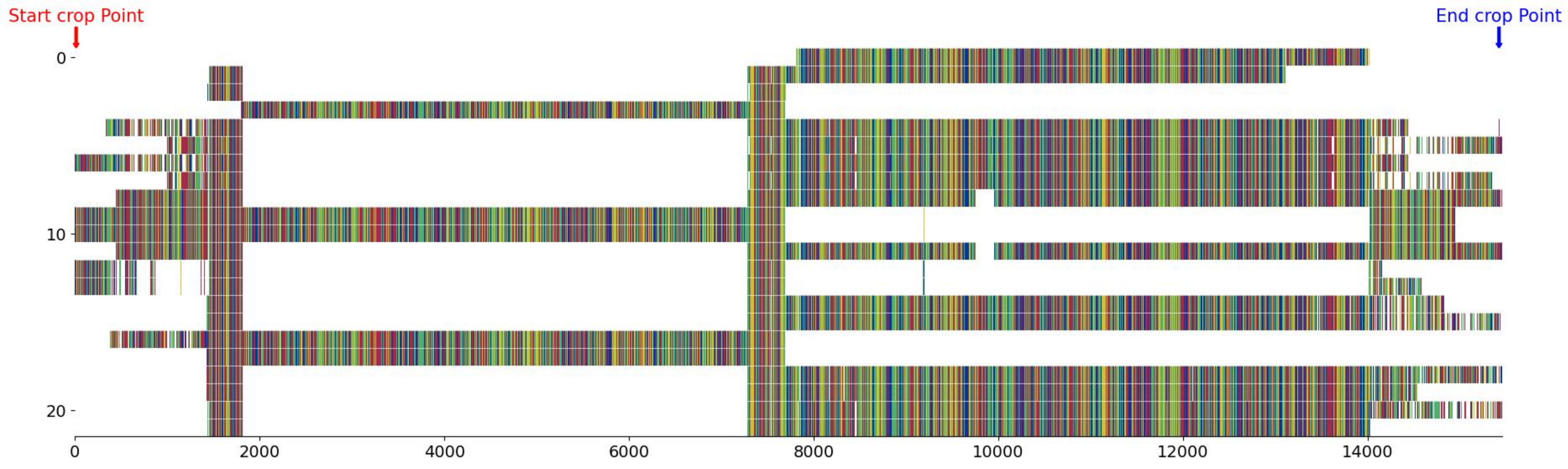
Start crop Point



MSA length = 15409

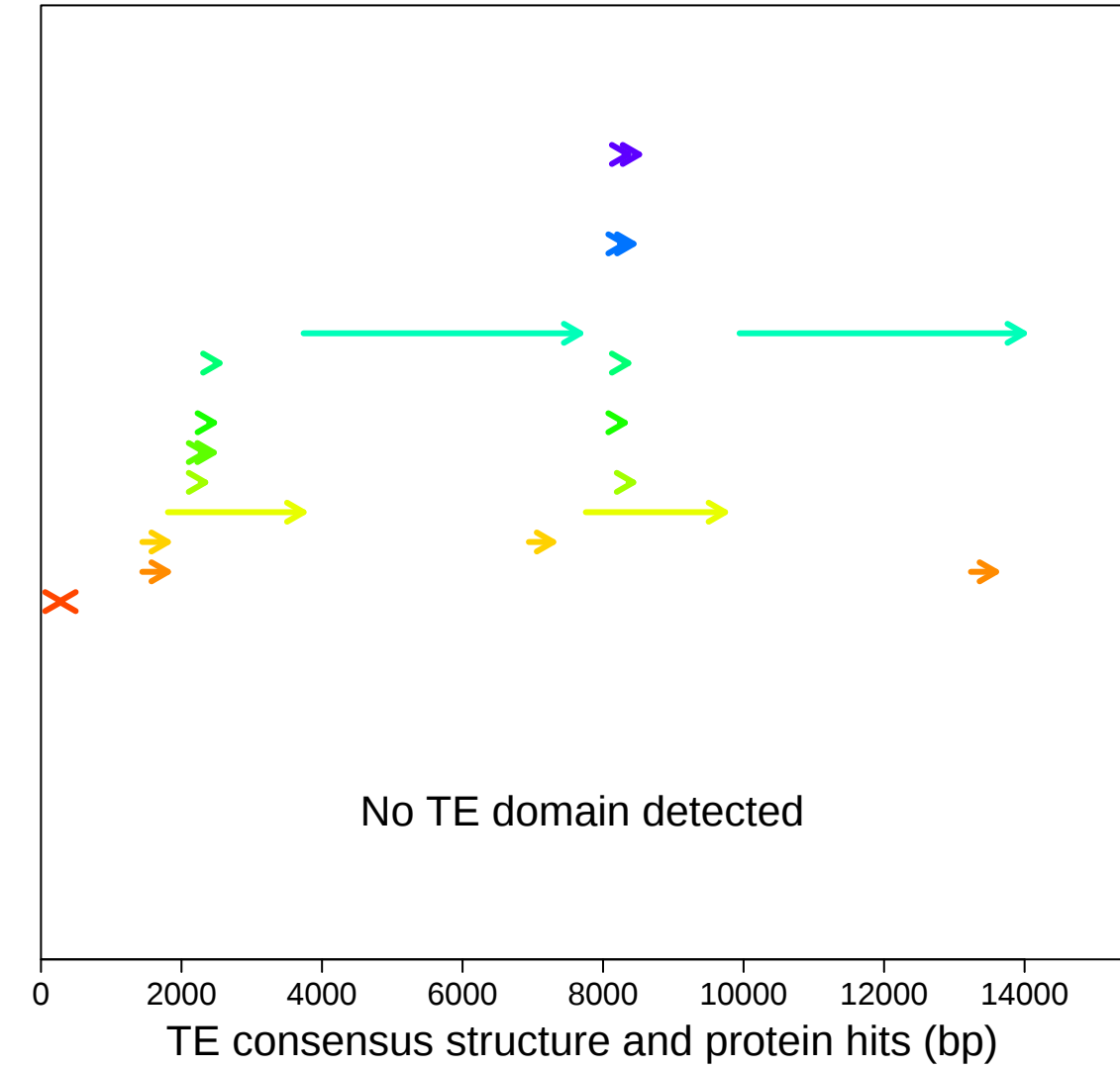
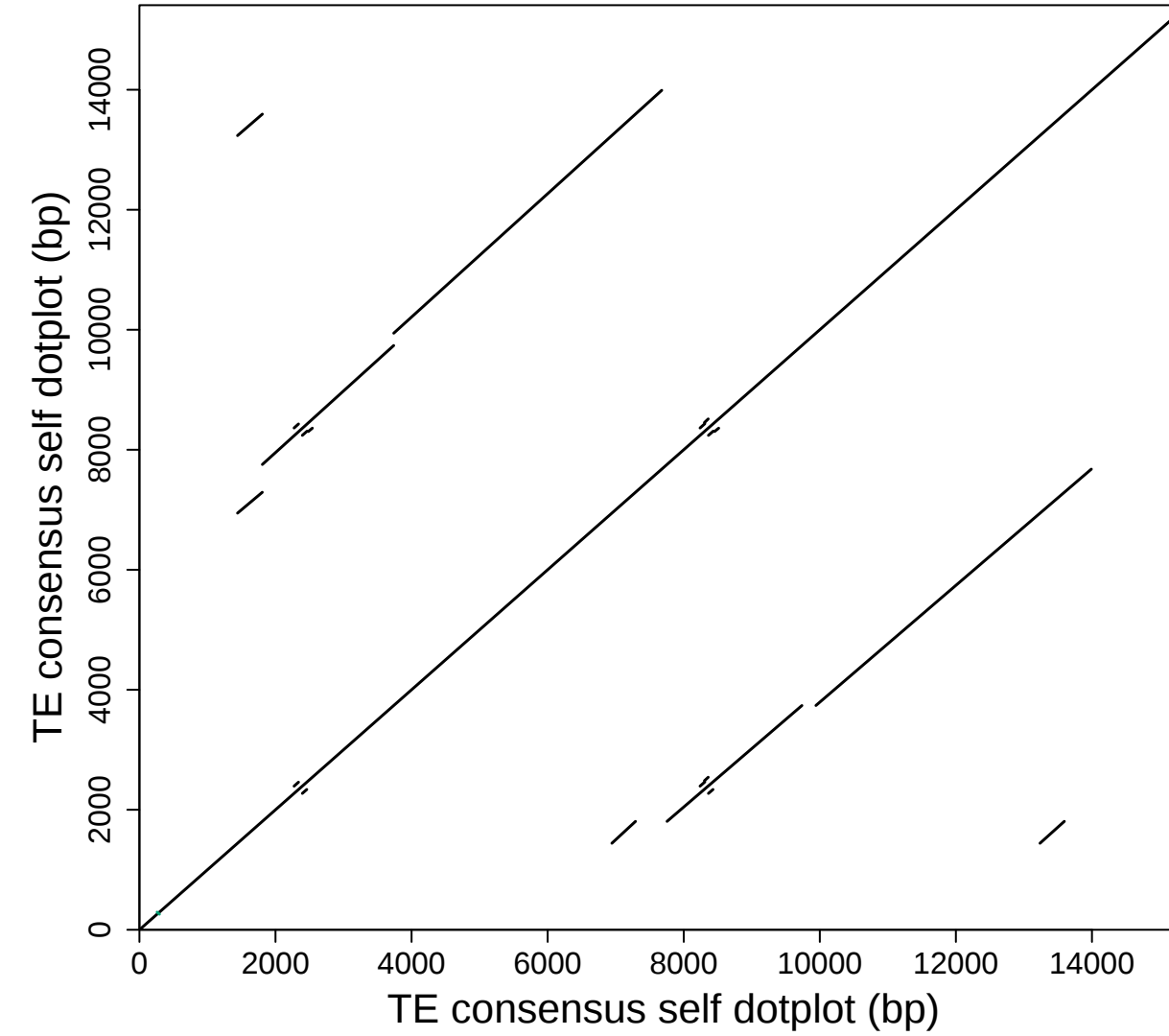
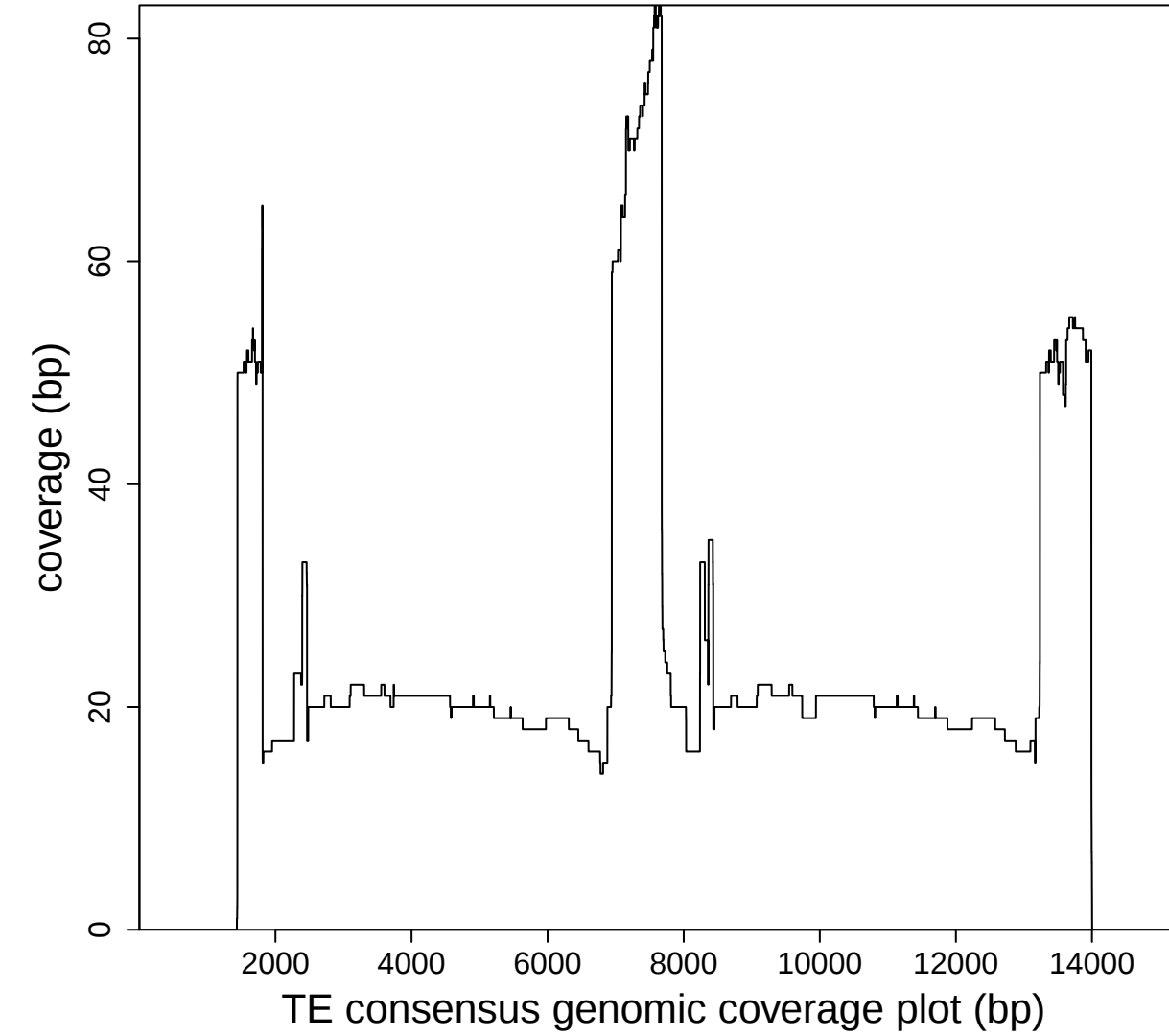
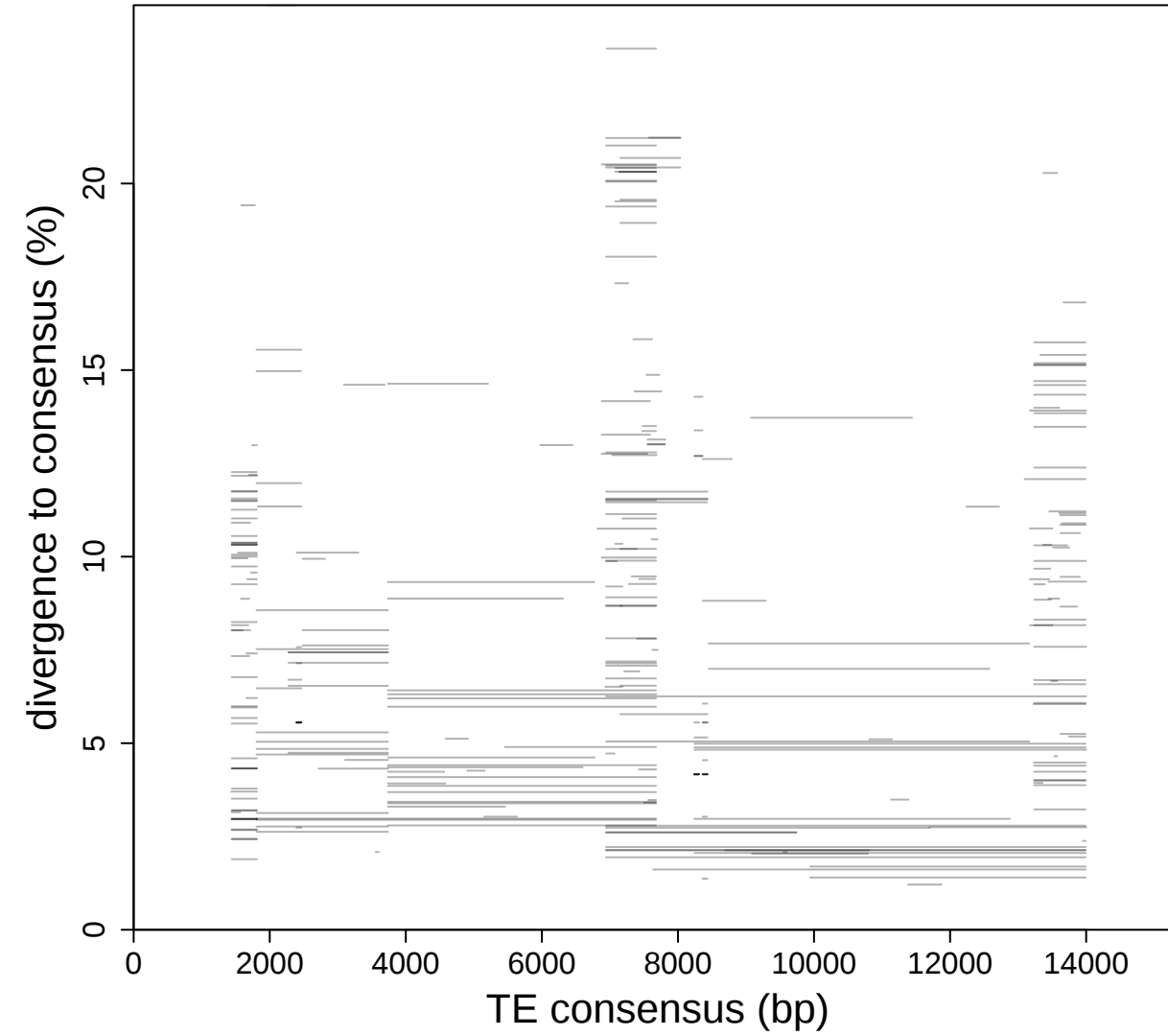
End crop Point





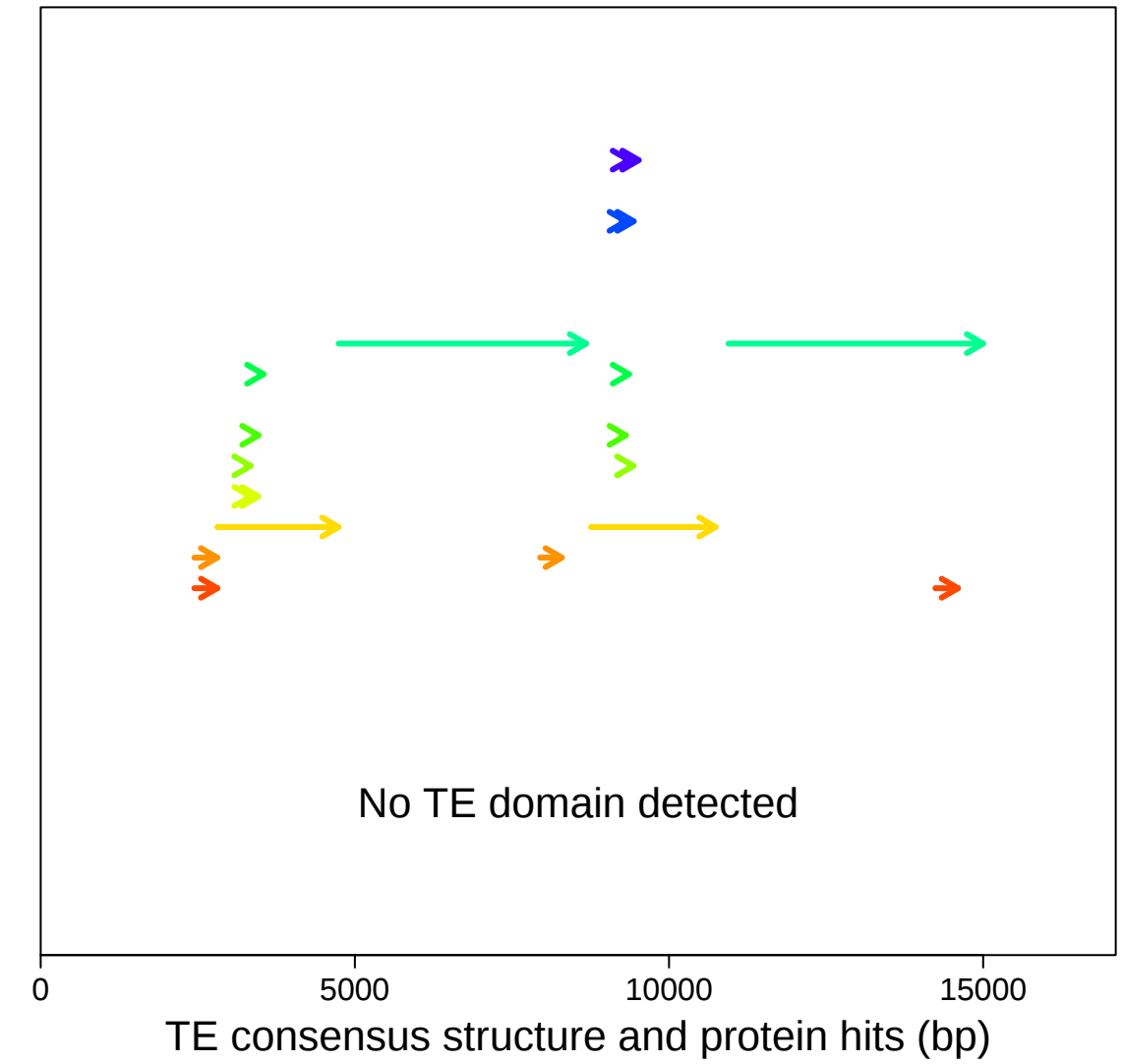
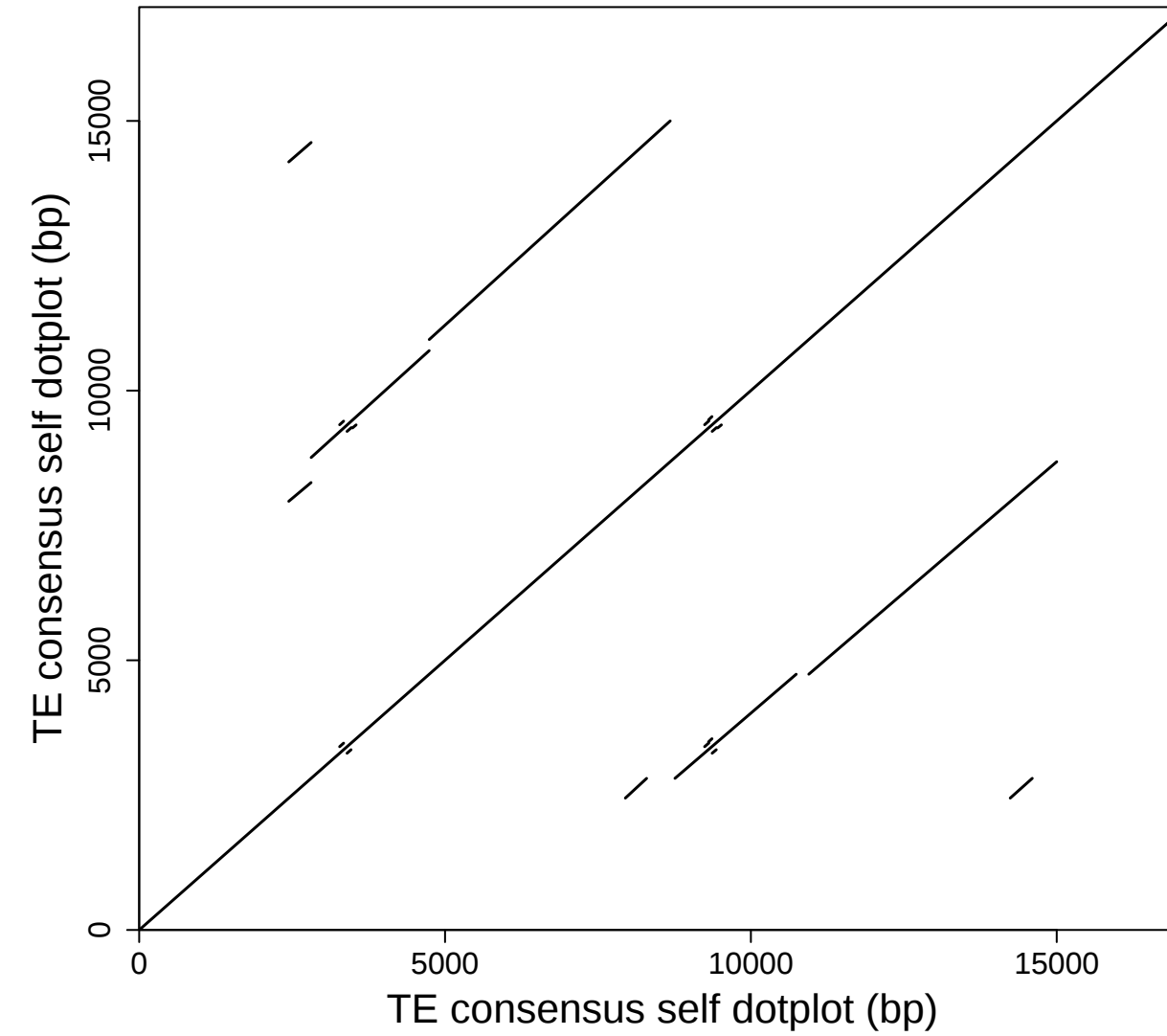
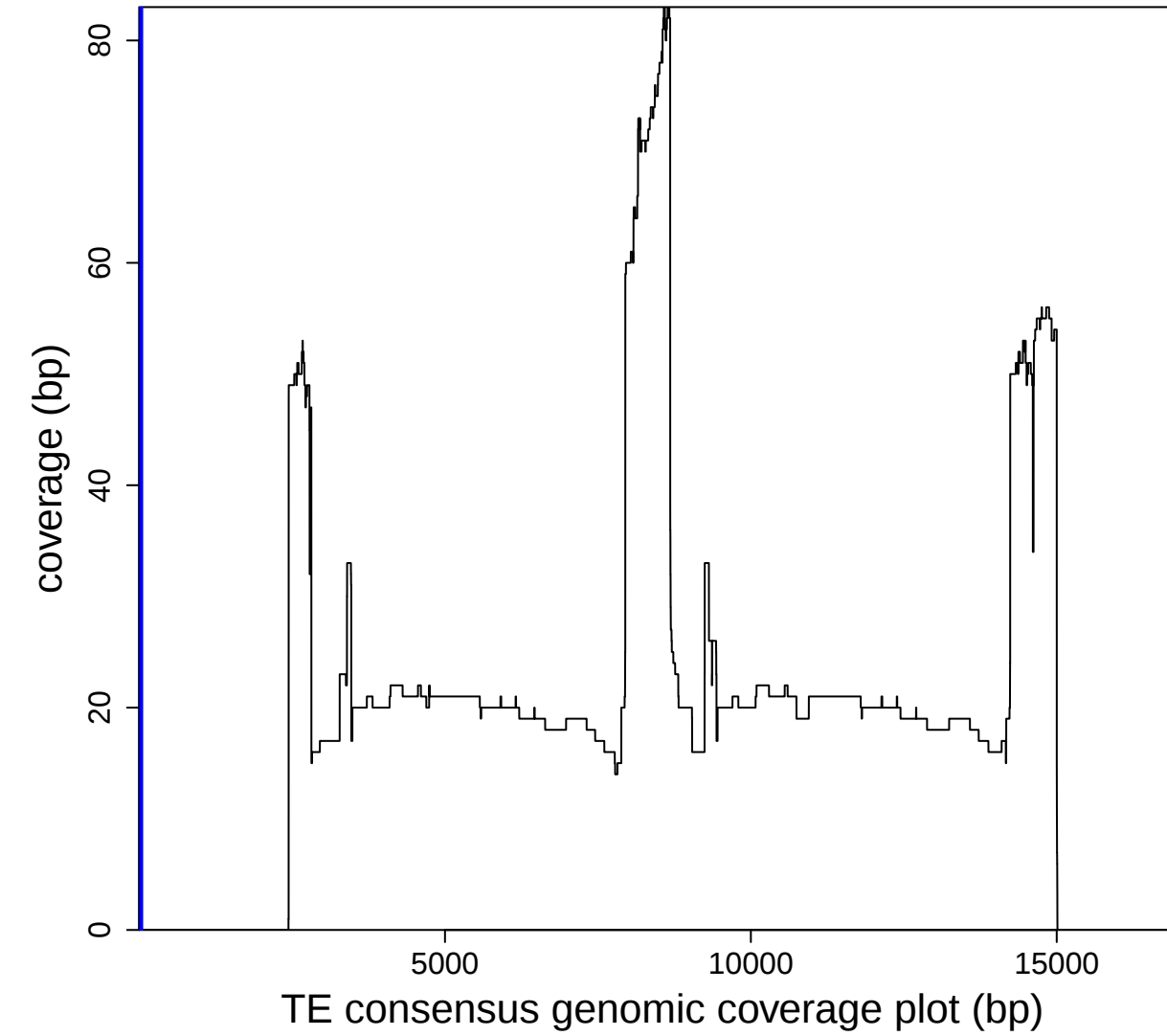
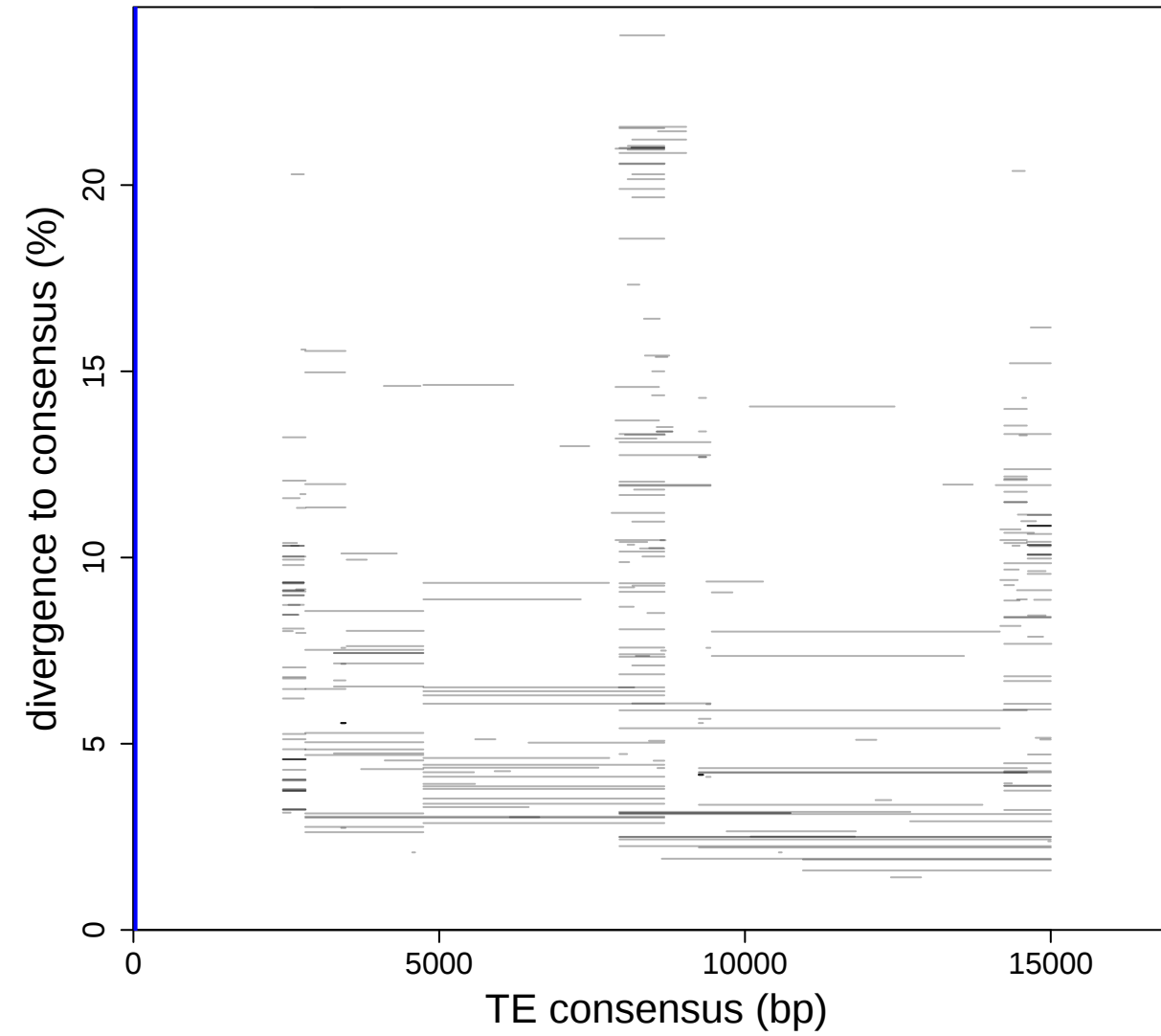
_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcIn.fa_al
size: 15409bp; fragments: 305; full length: 0 (>=13868.1bp)

After TEtrimmer 15409 bp

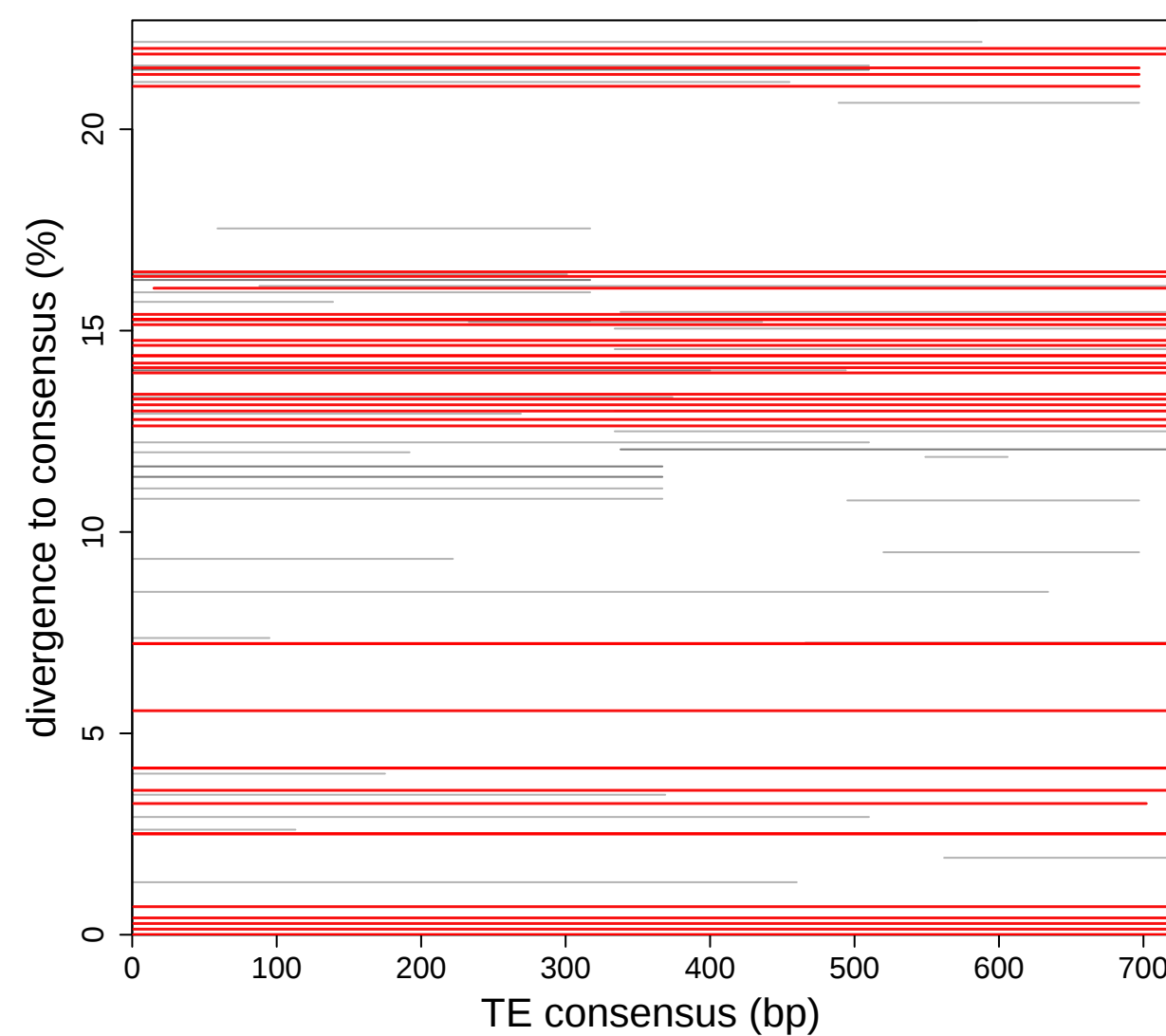


d_g_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcln.fa
size: 17110bp; fragments: 310; full length: 0 (>=15399bp)

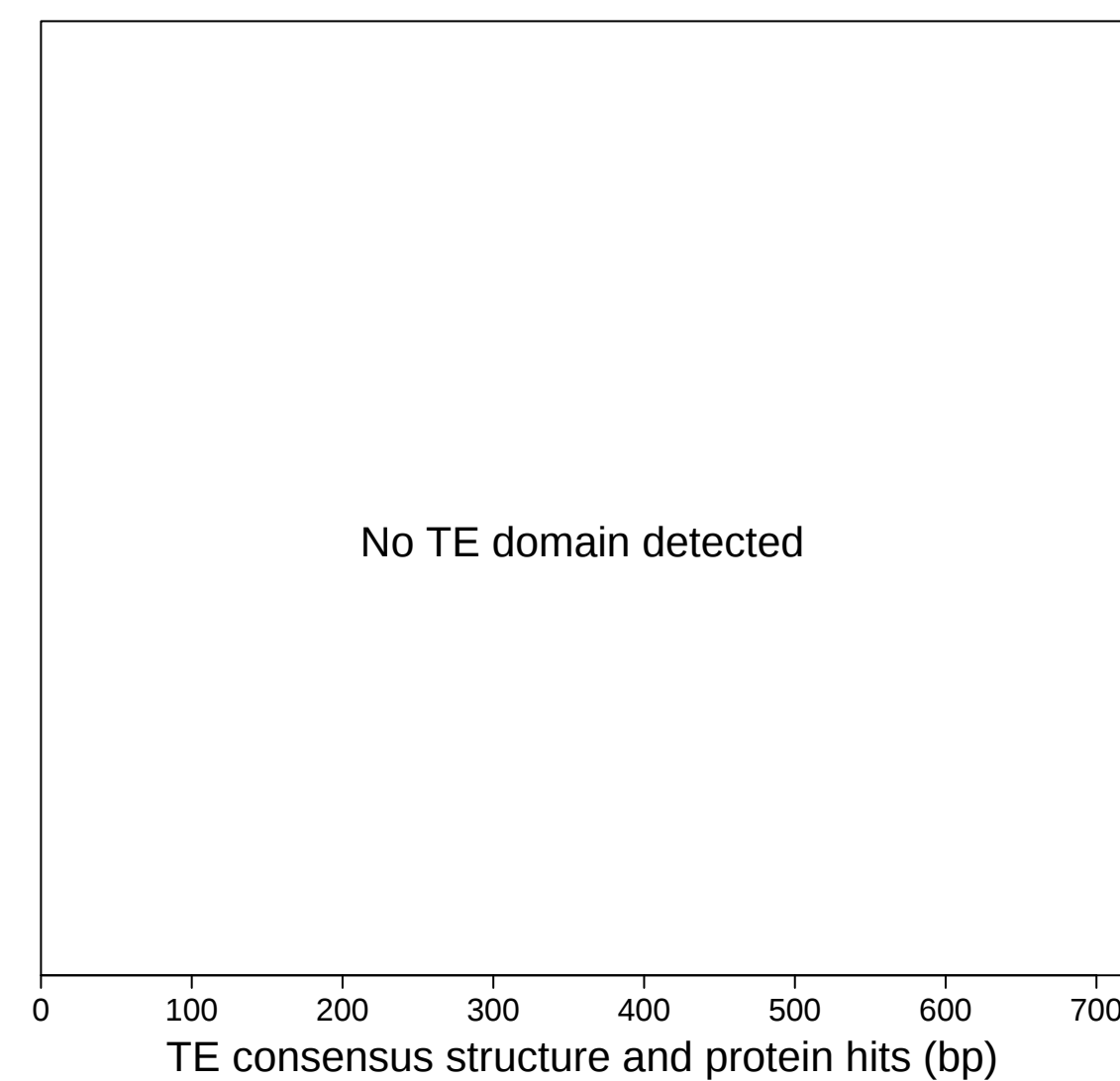
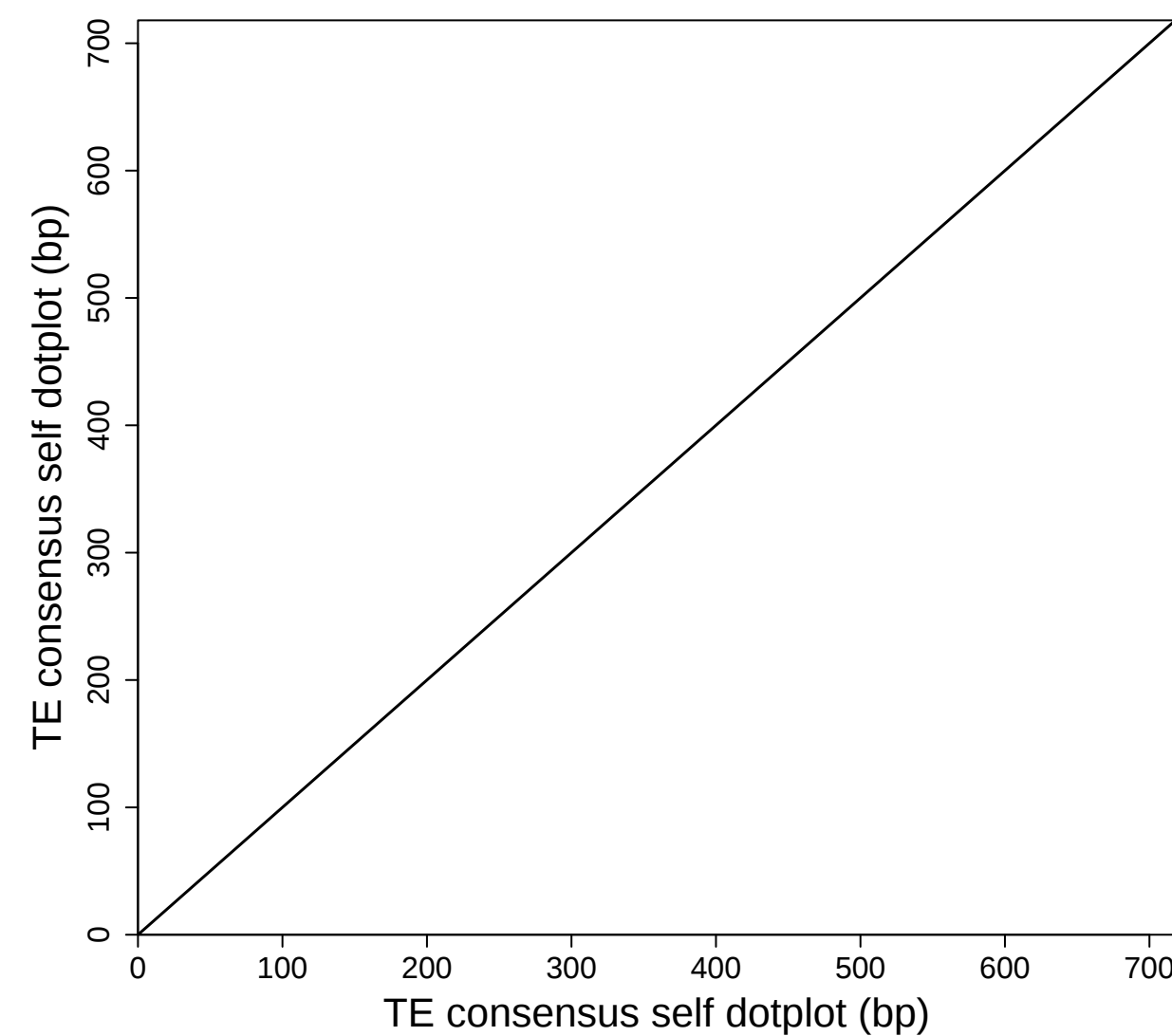
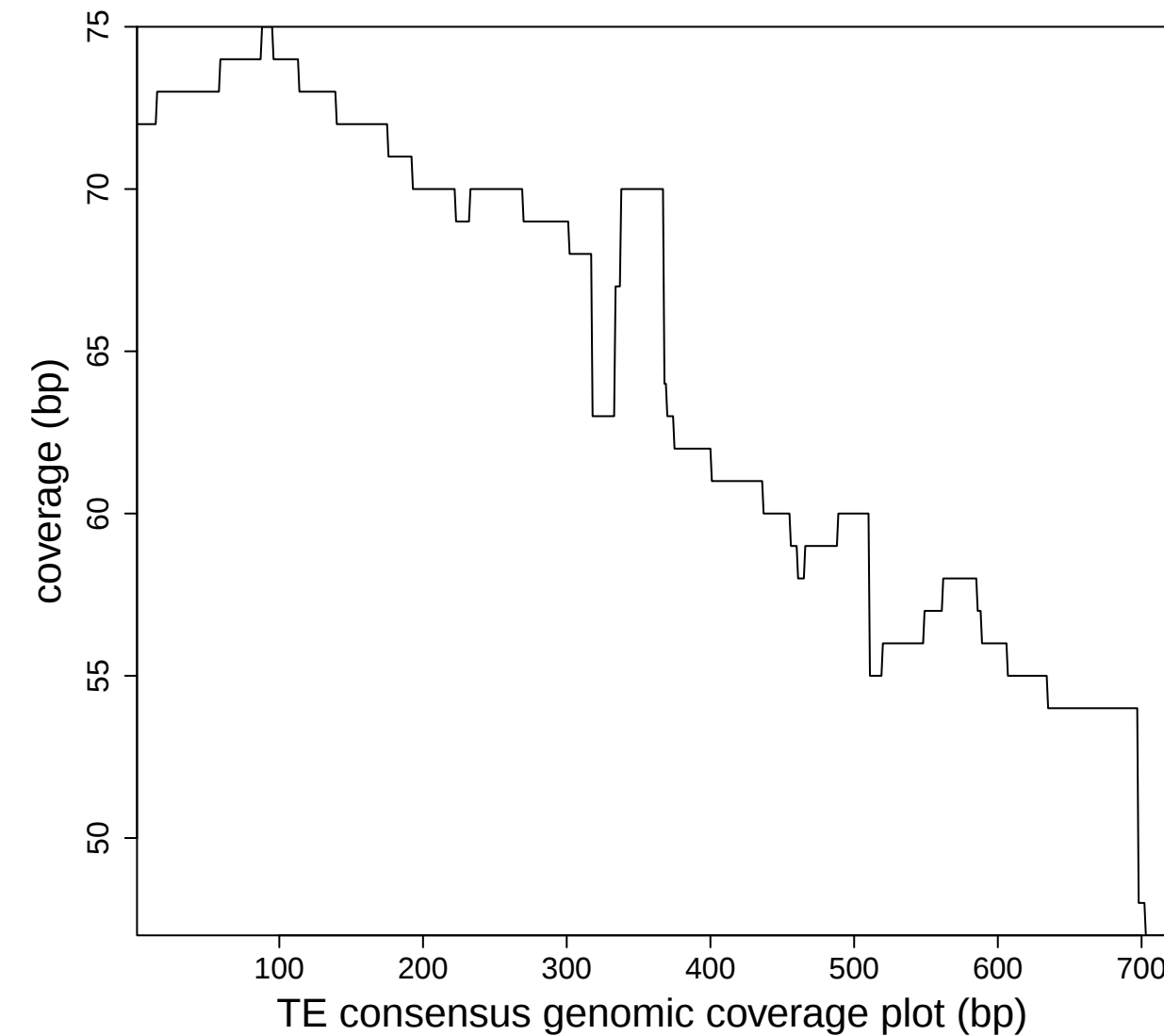
After TEtrimmer Extended plot Blue lines are boundaries



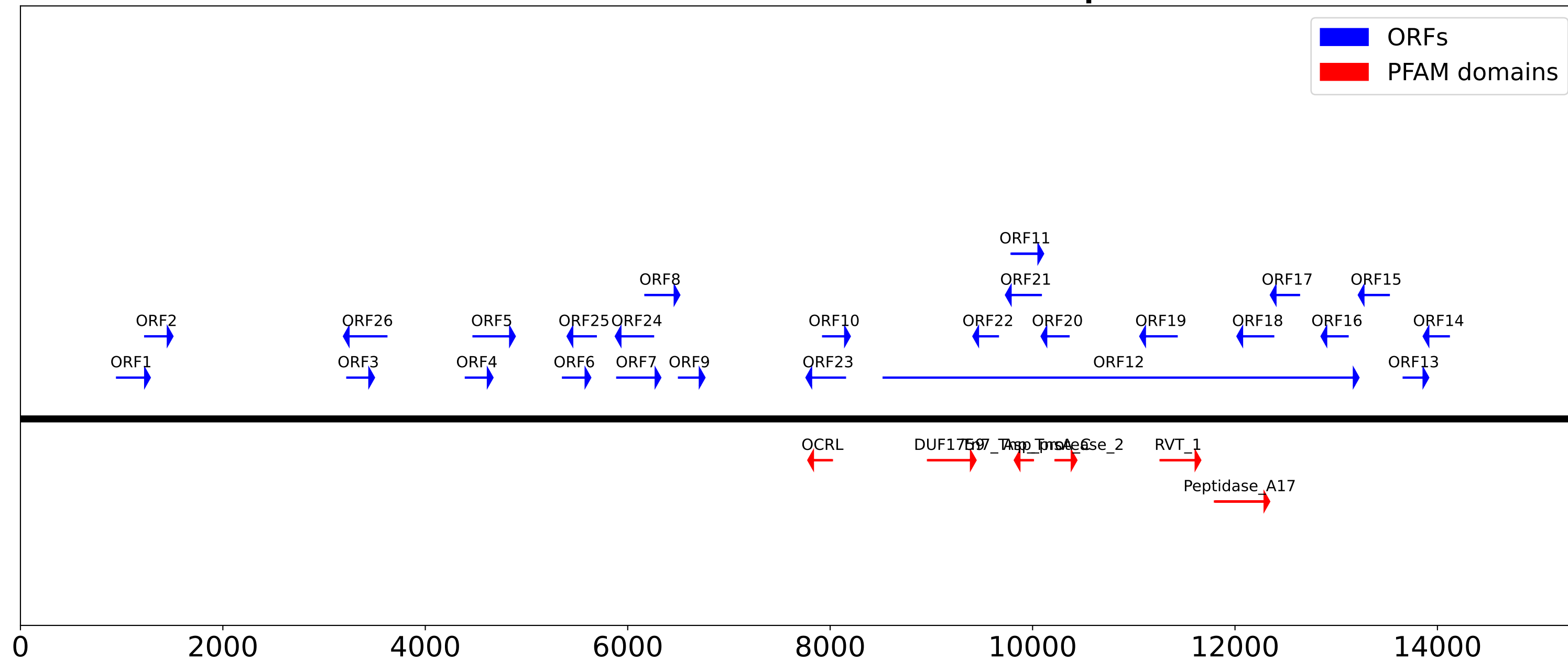
TE: ltr_1_family_304
size: 718bp; fragments: 89; full length: 41 (≥ 646.2 bp)



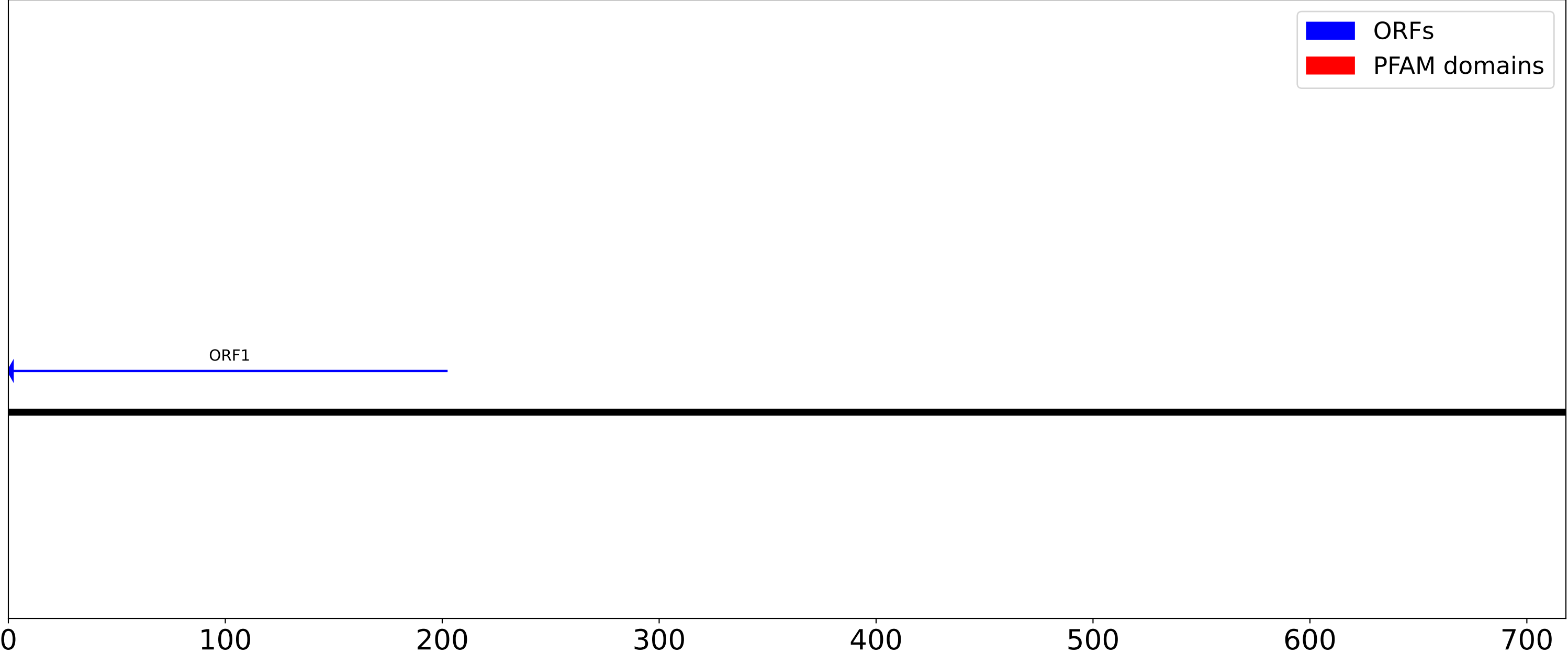
Before TEtrimmer 718 bp



After TEtrimmer ORF and PFAM domain plot



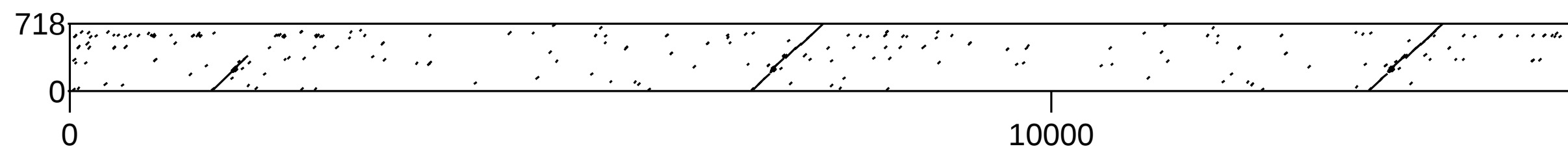
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

nsus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)