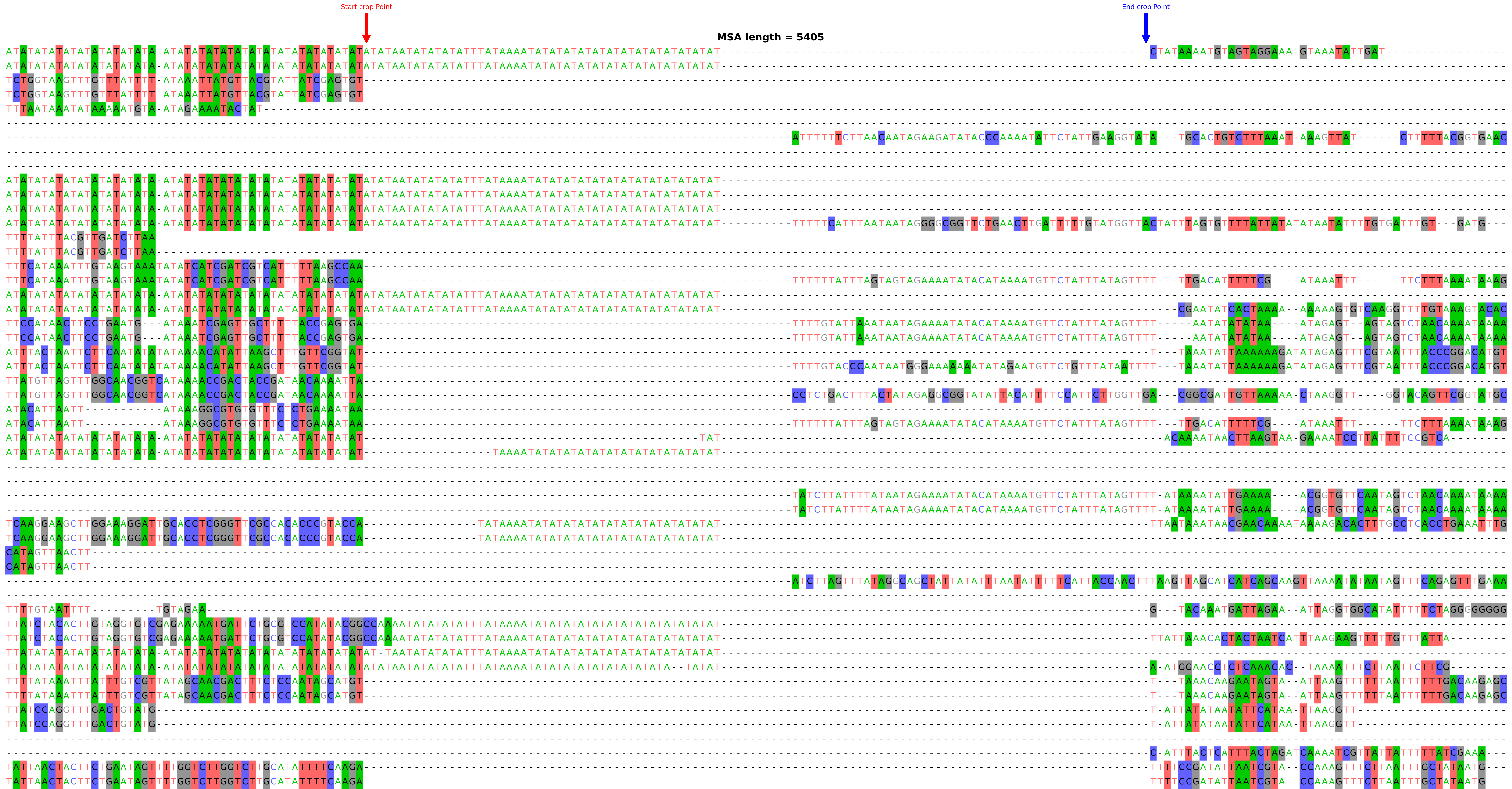
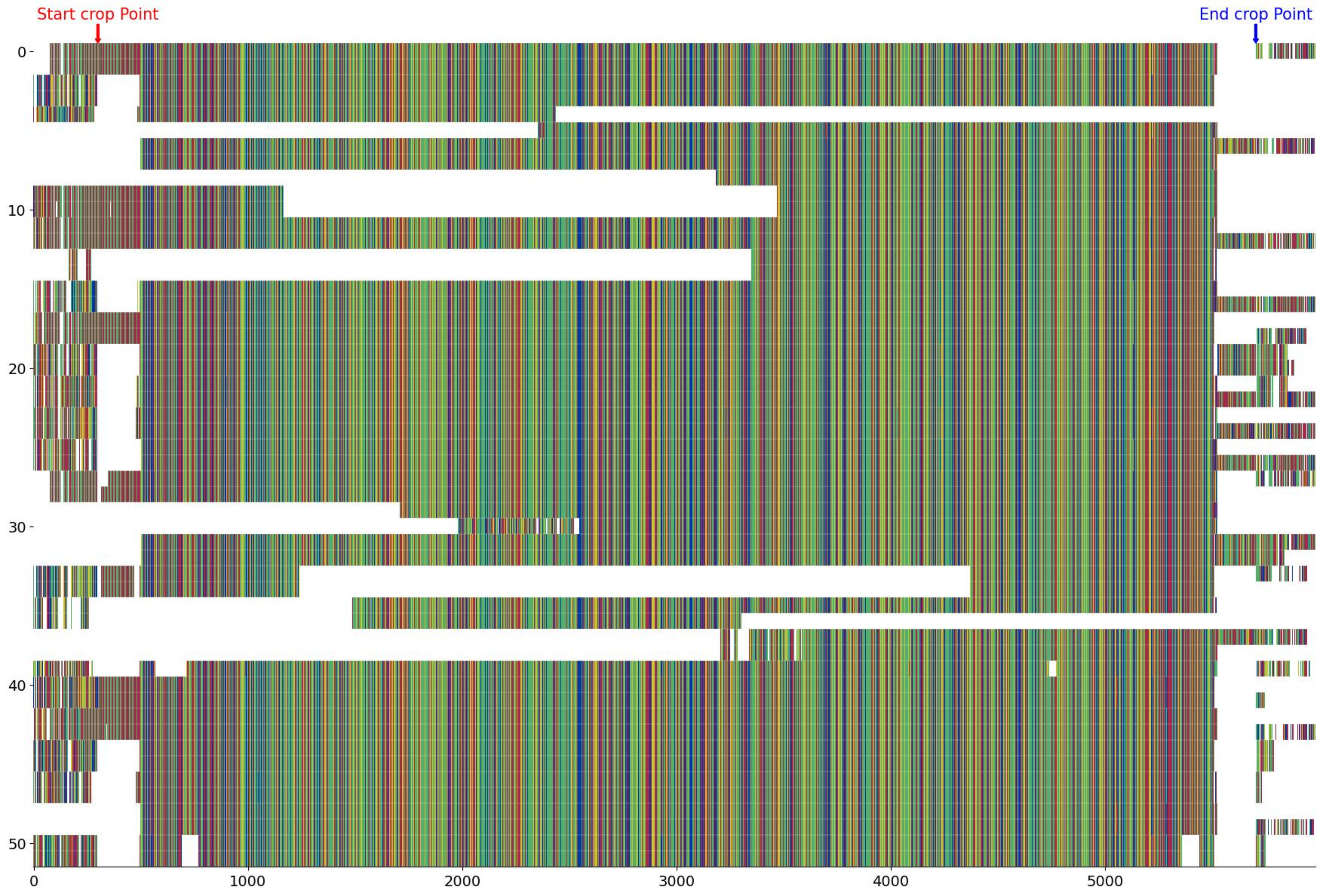
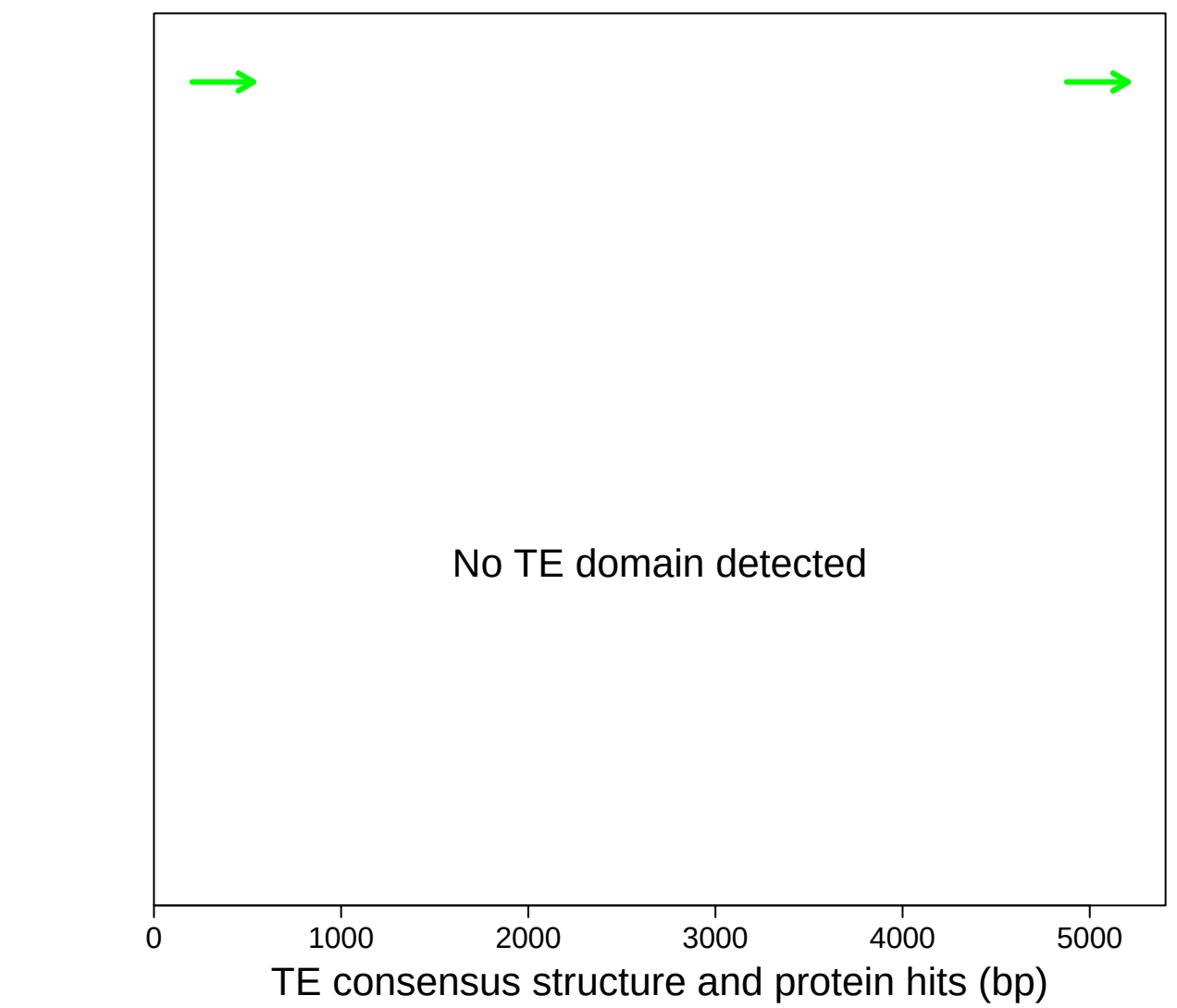
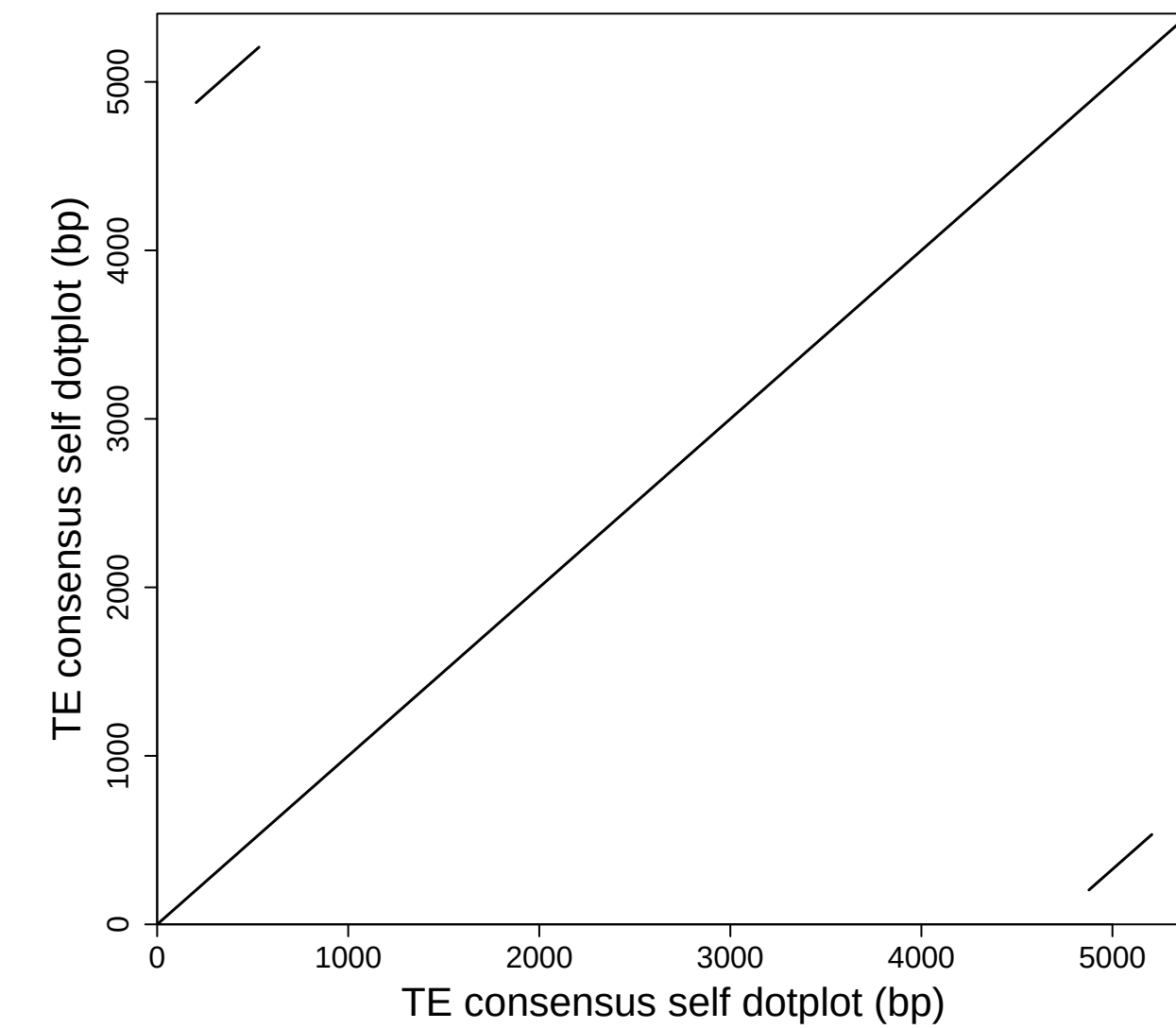
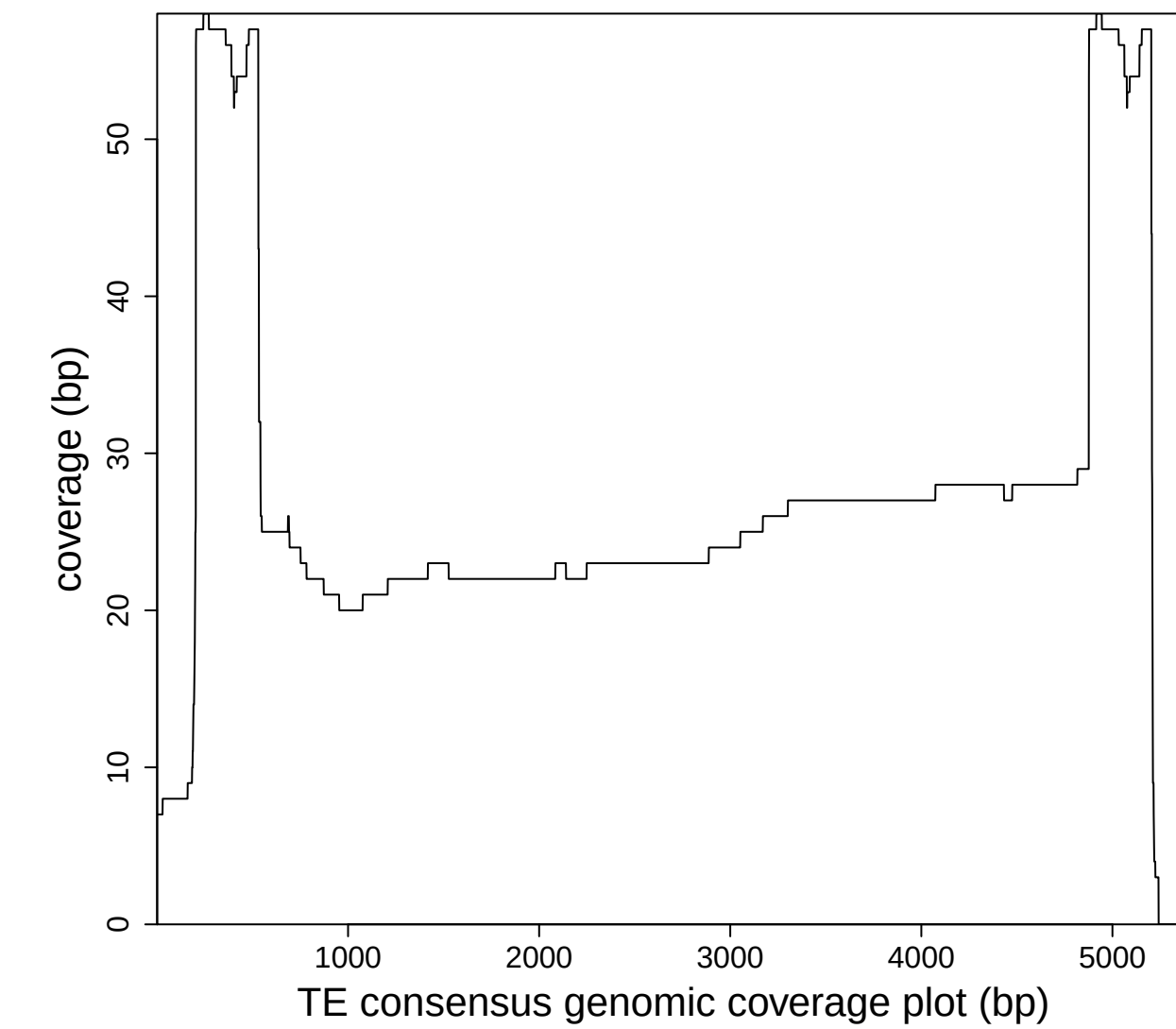
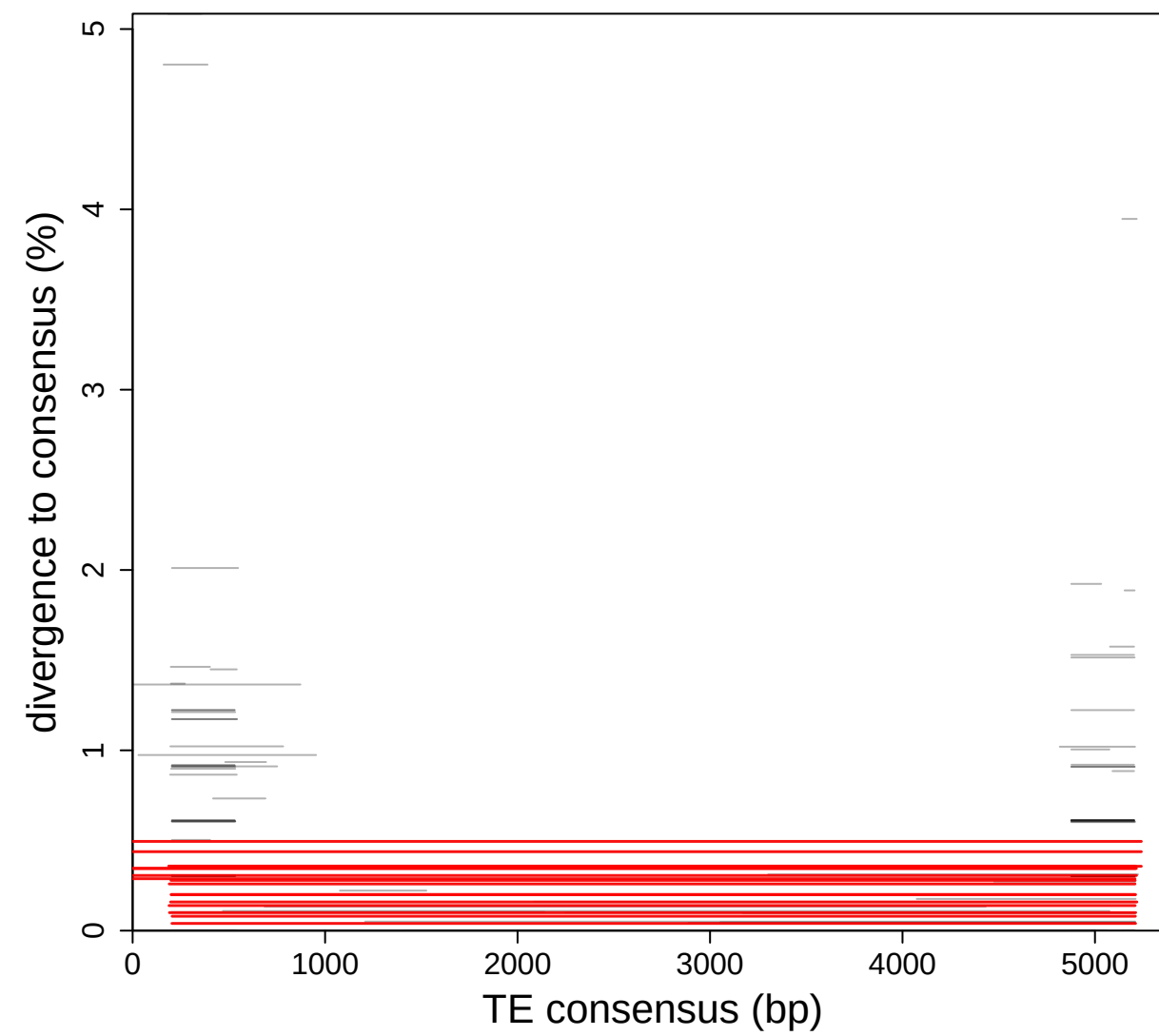






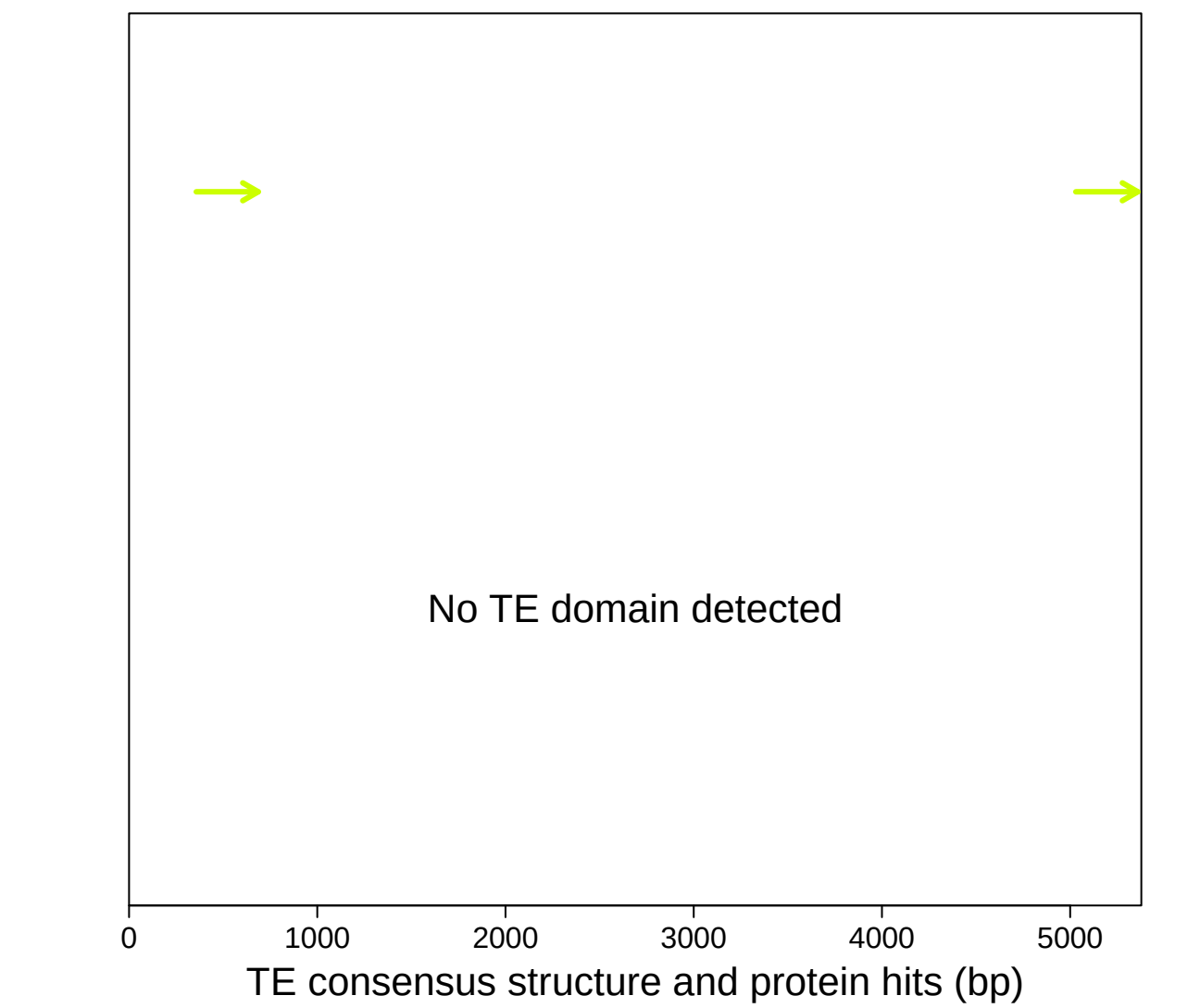
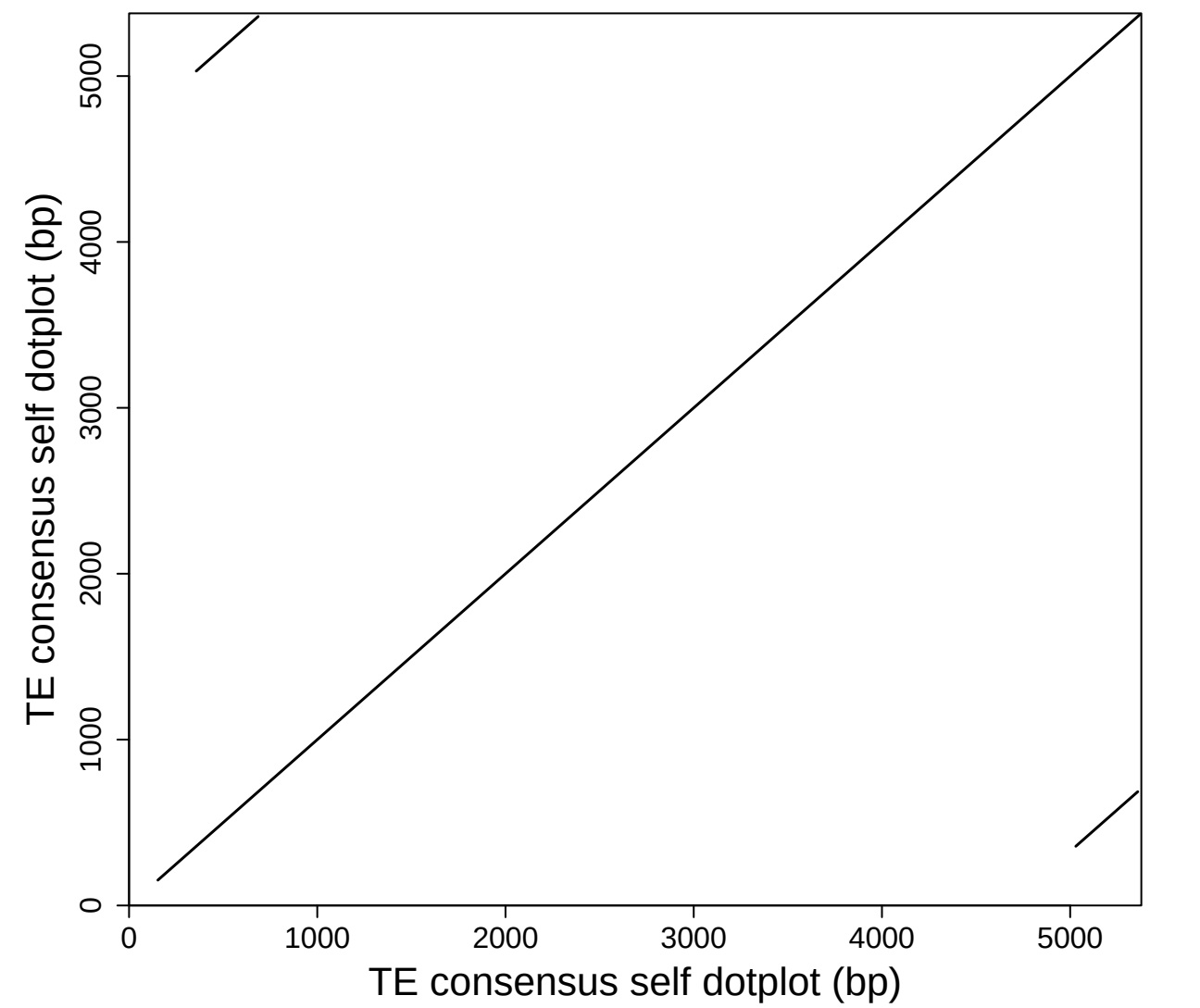
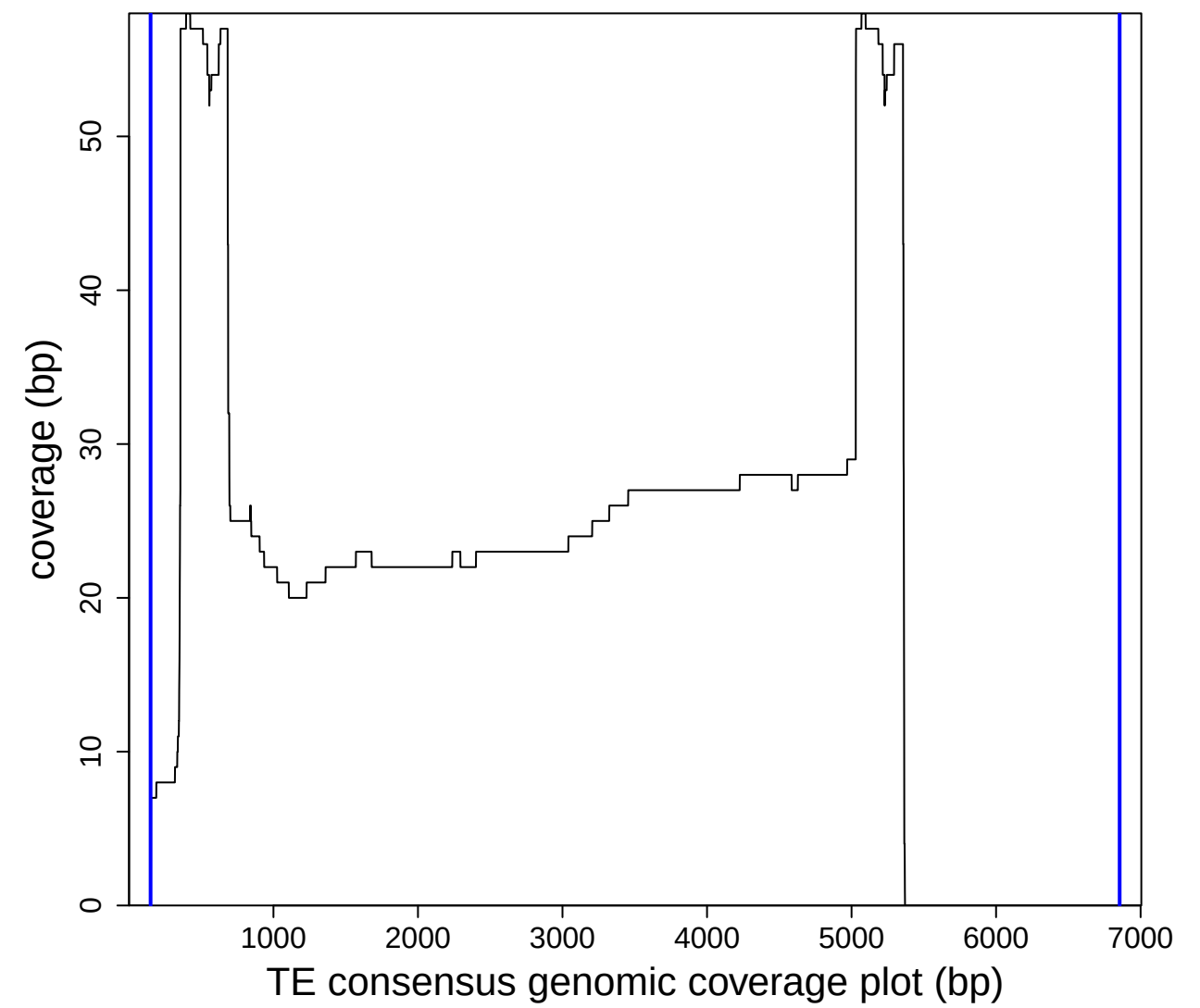
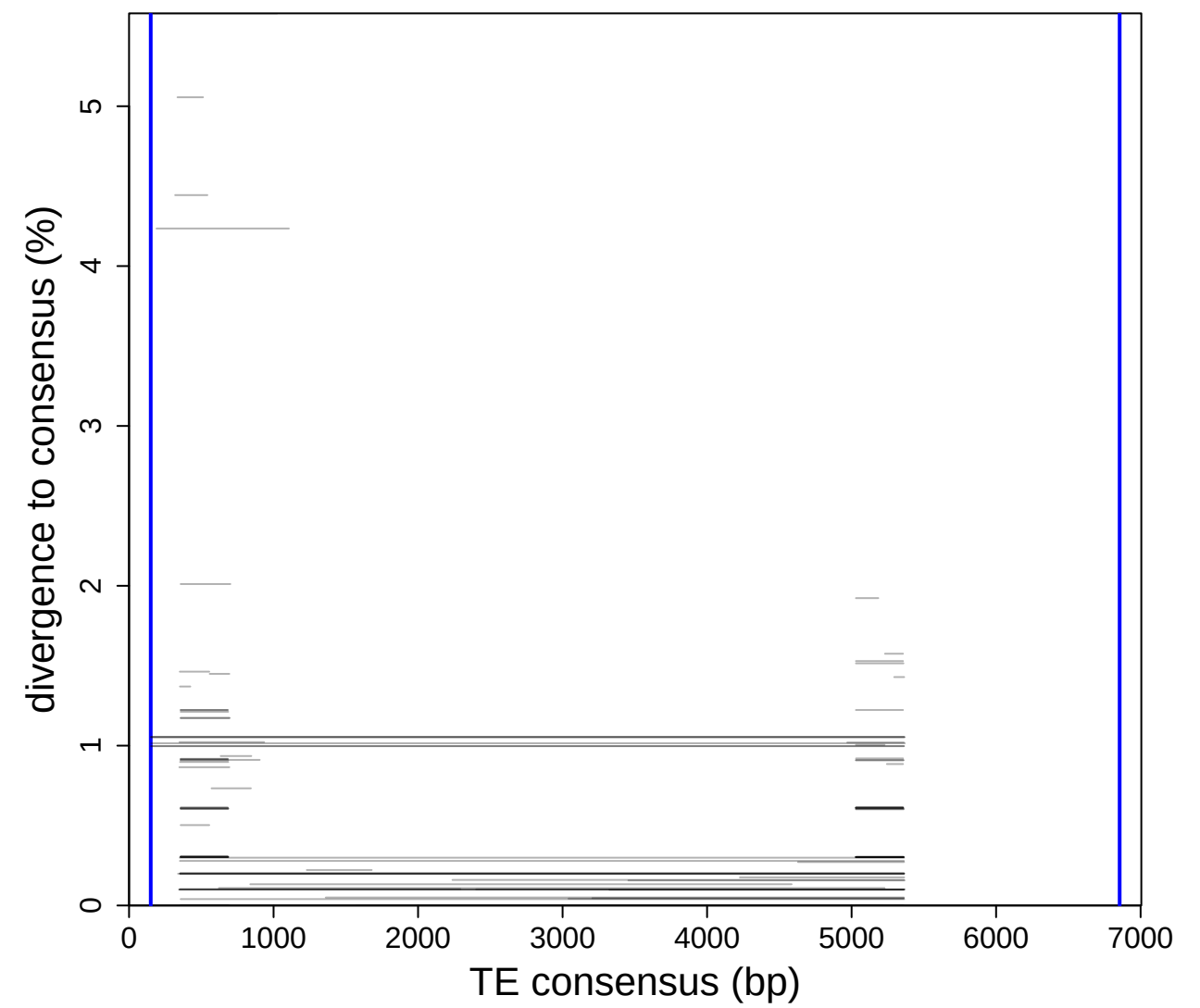
_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcln.fa al
size: 5405bp; fragments: 110; full length: 17 (≥ 4864.5 bp)

After TEtrimmer 5405 bp



bed_g_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bclr
size: 7005bp; fragments: 109; full length: 0 (>=6304.5bp)

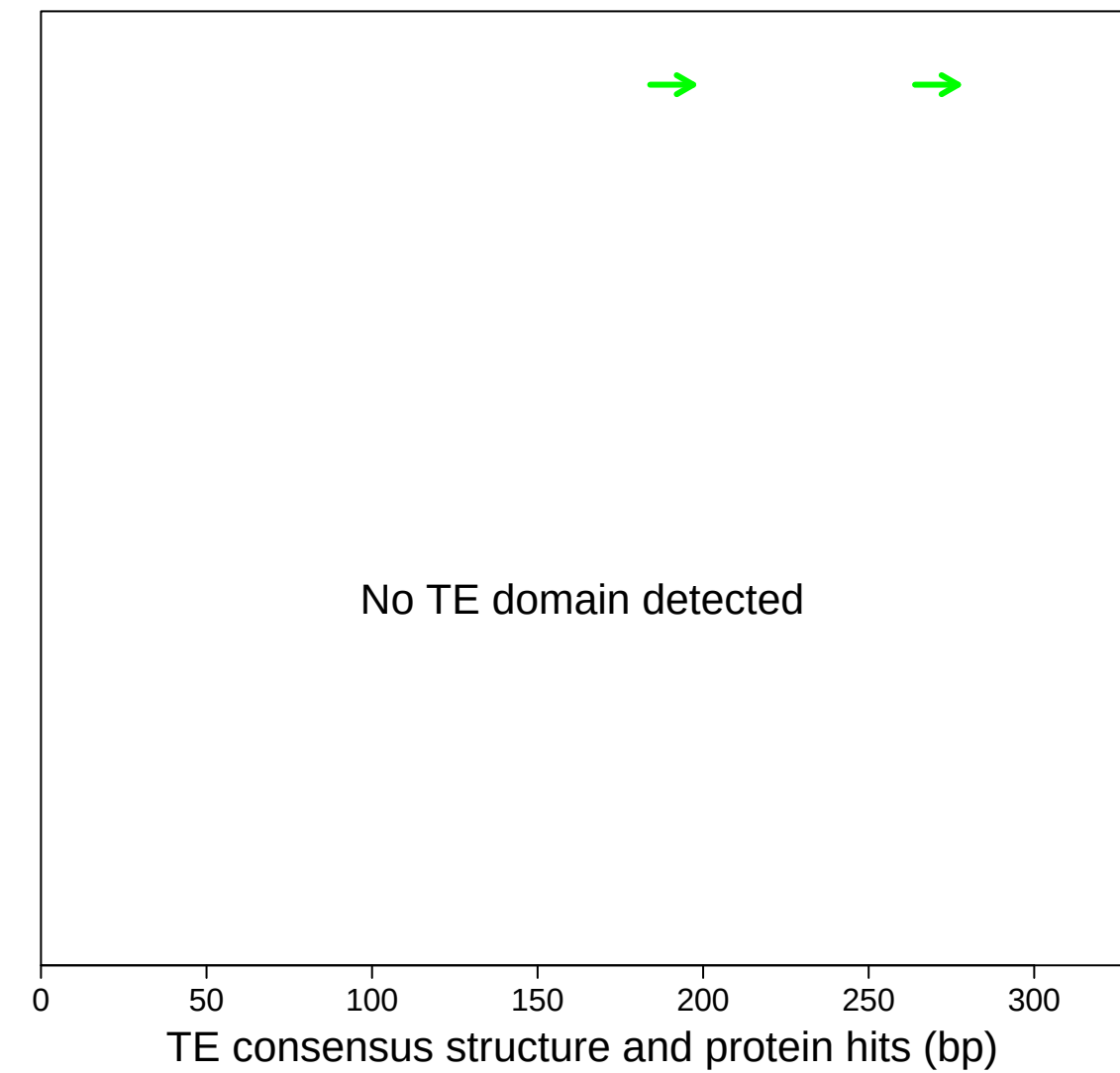
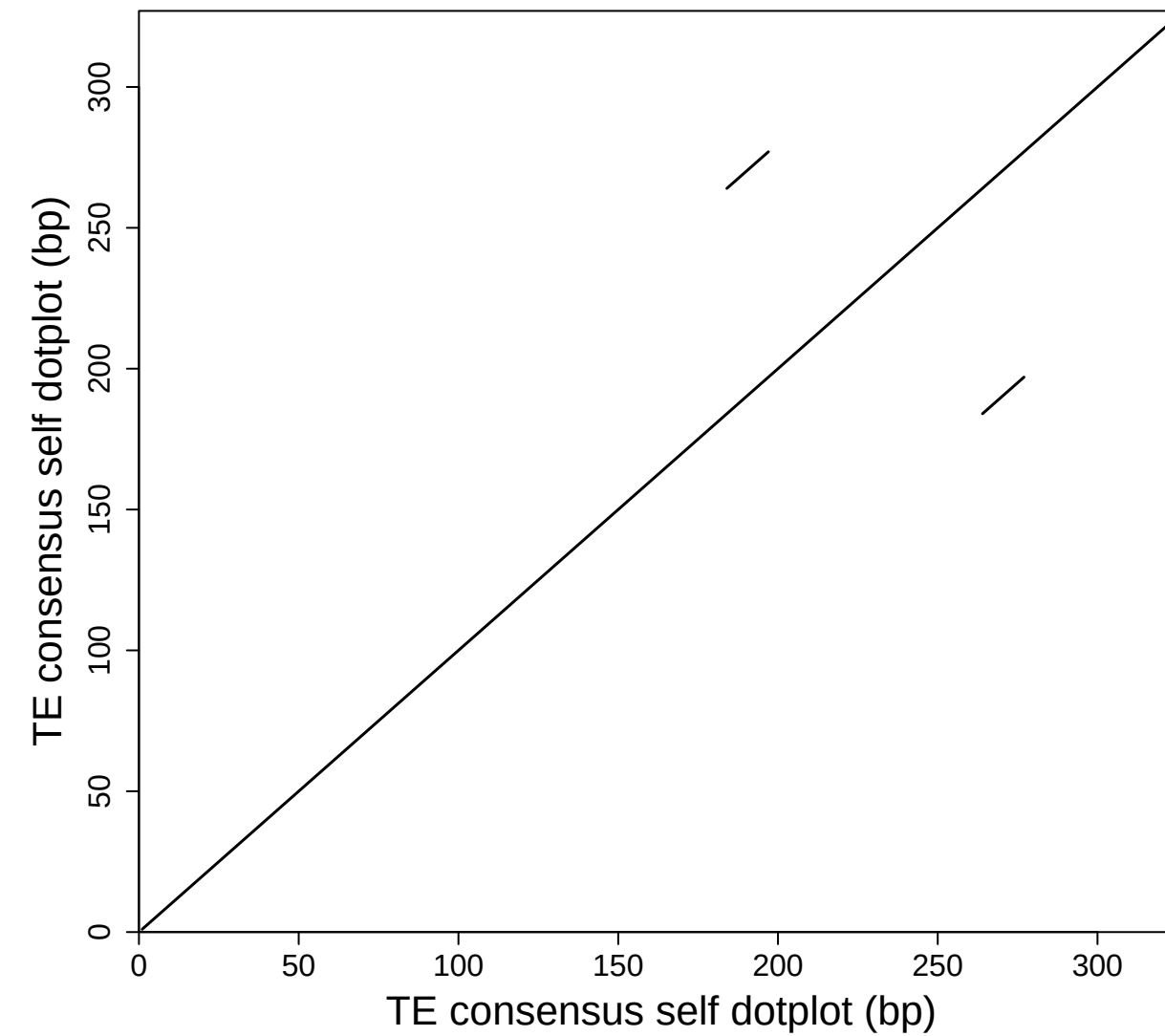
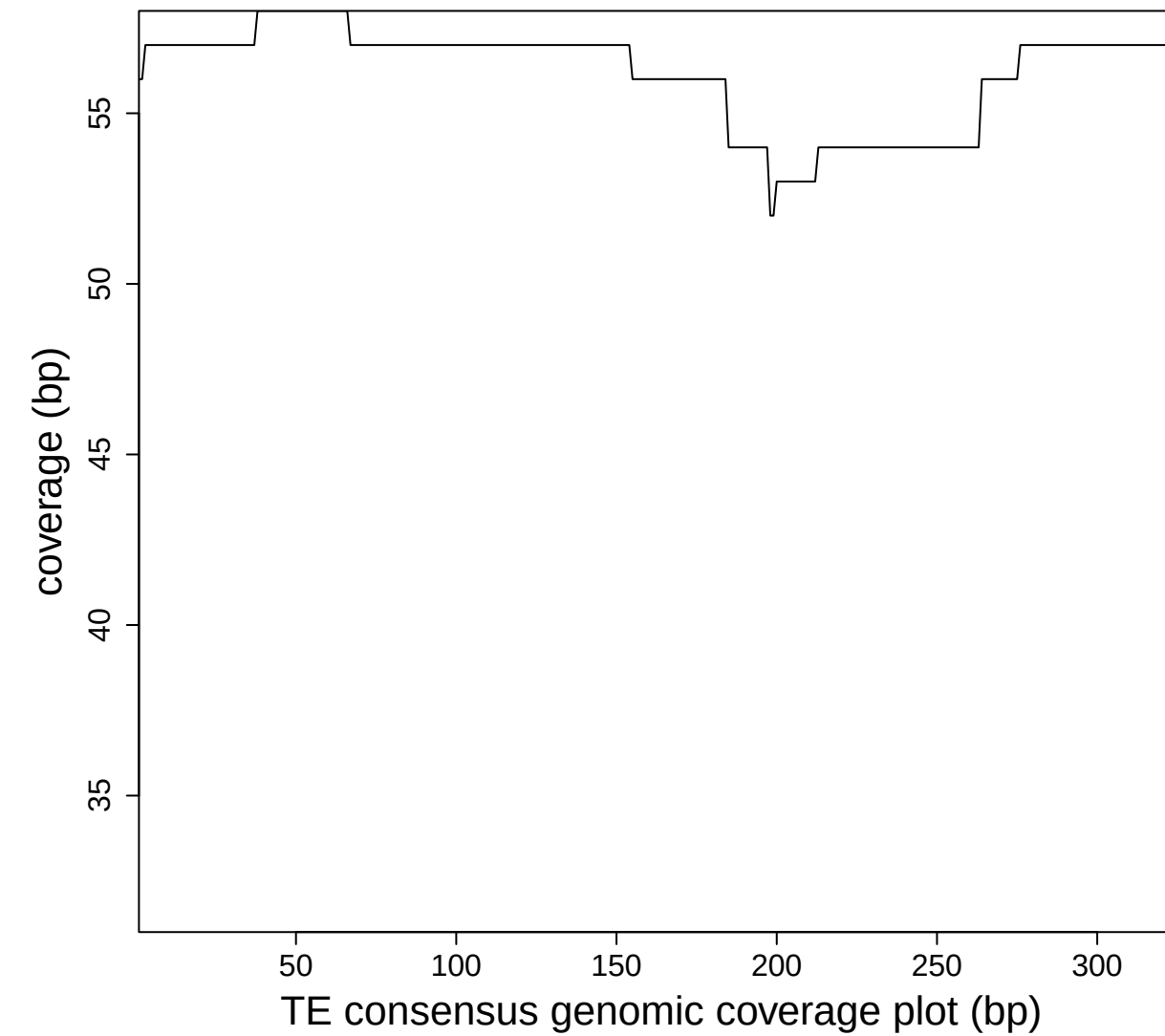
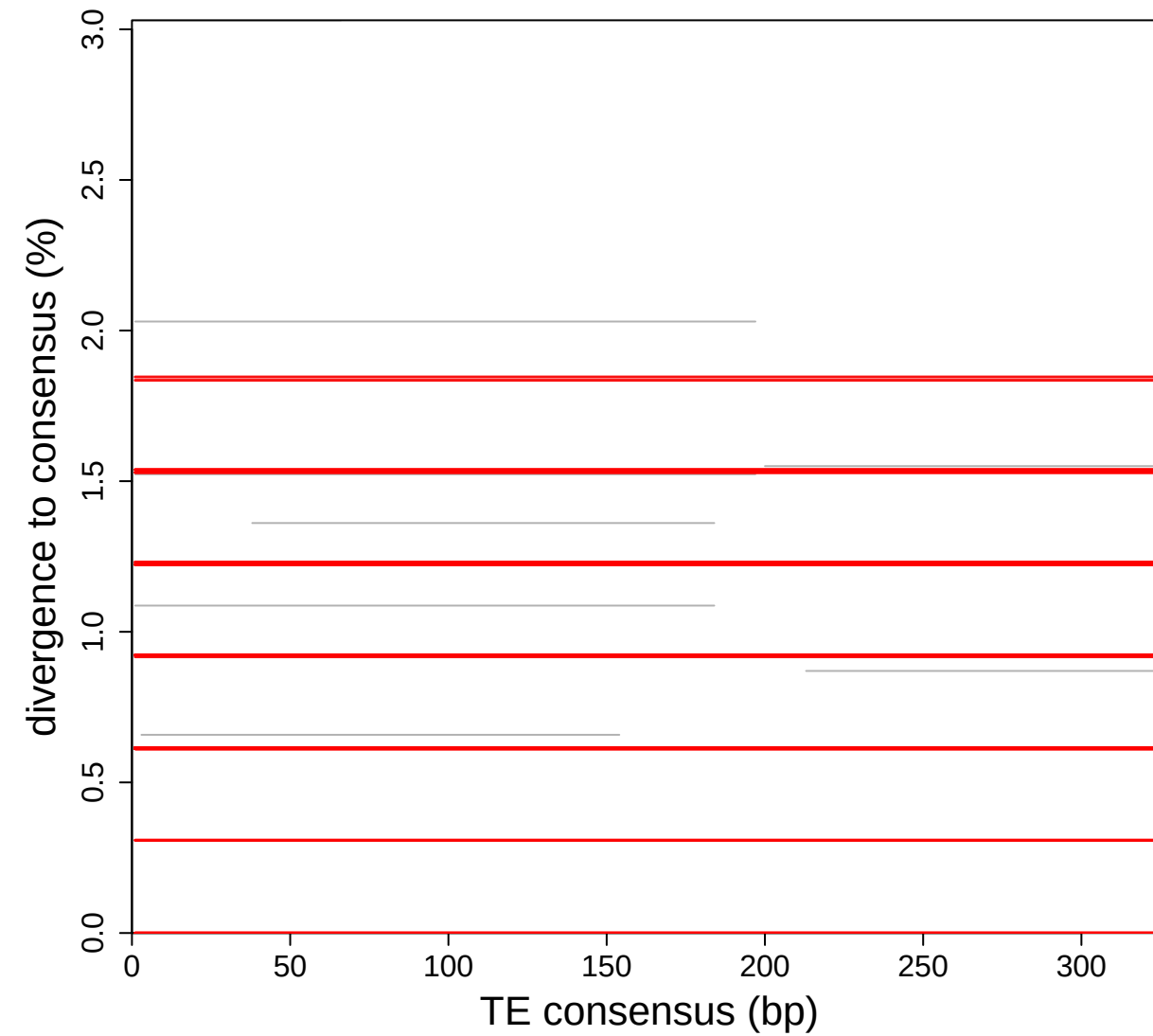
After TEtrimmer Extended plot Blue lines are boundaries



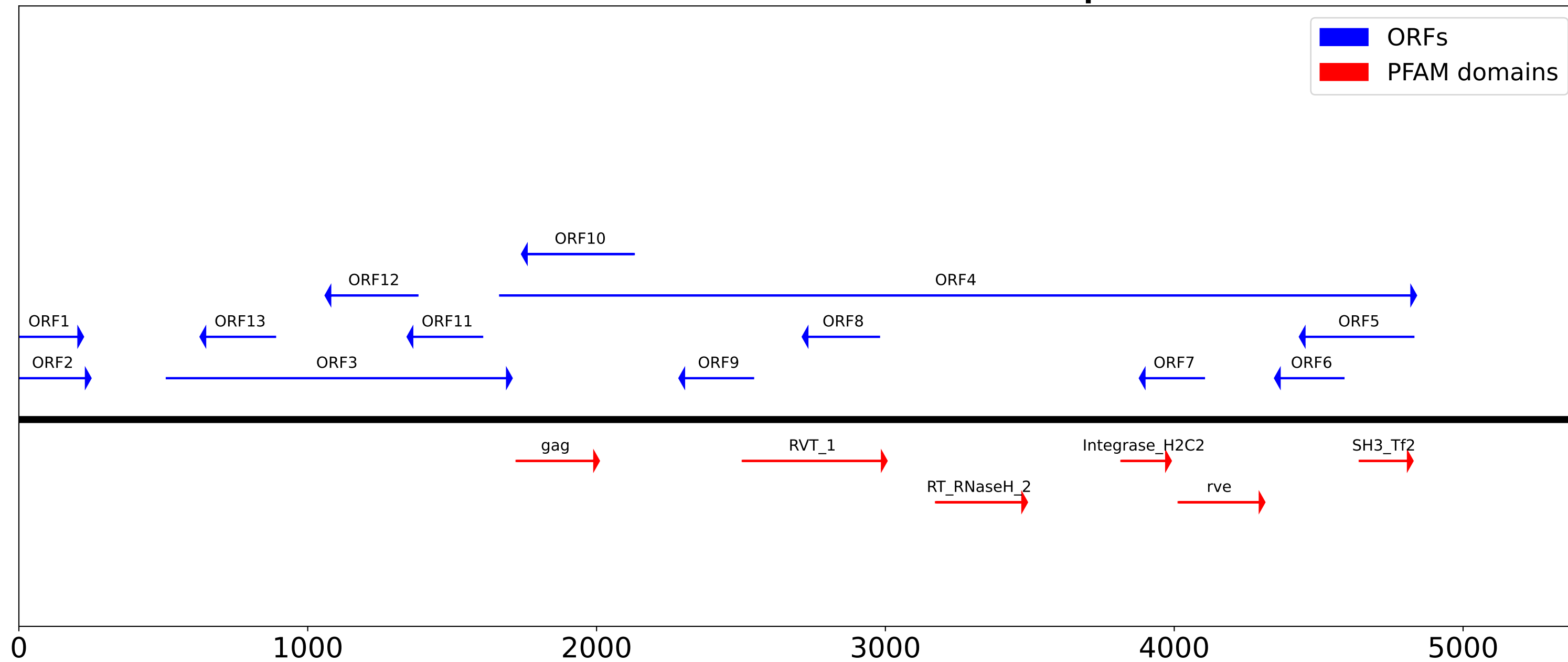
Before TEtrimmer 327 bp

TE: ltr_1_family_414

size: 327bp; fragments: 63; full length: 52 (≥ 294.3 bp)



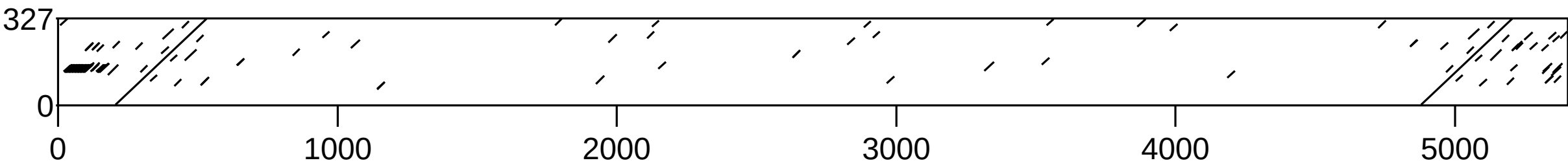
After TEtrimmer ORF and PFAM domain plot



Dotplot

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

consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)