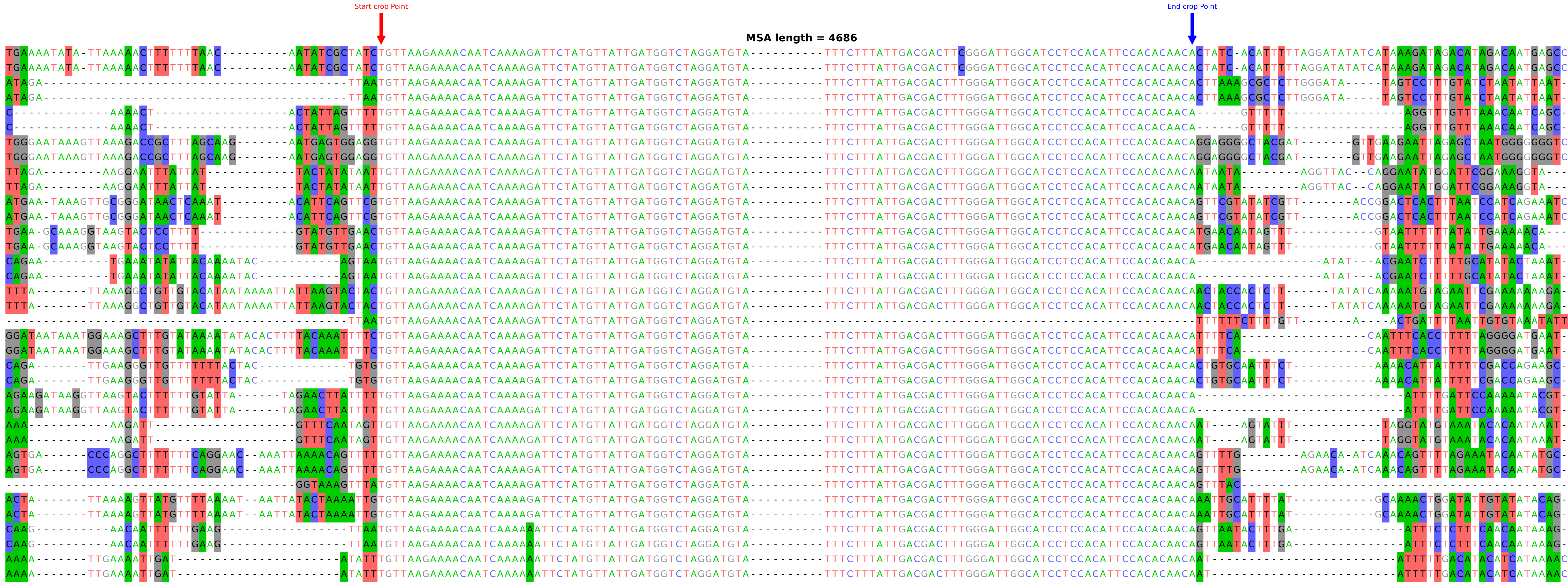
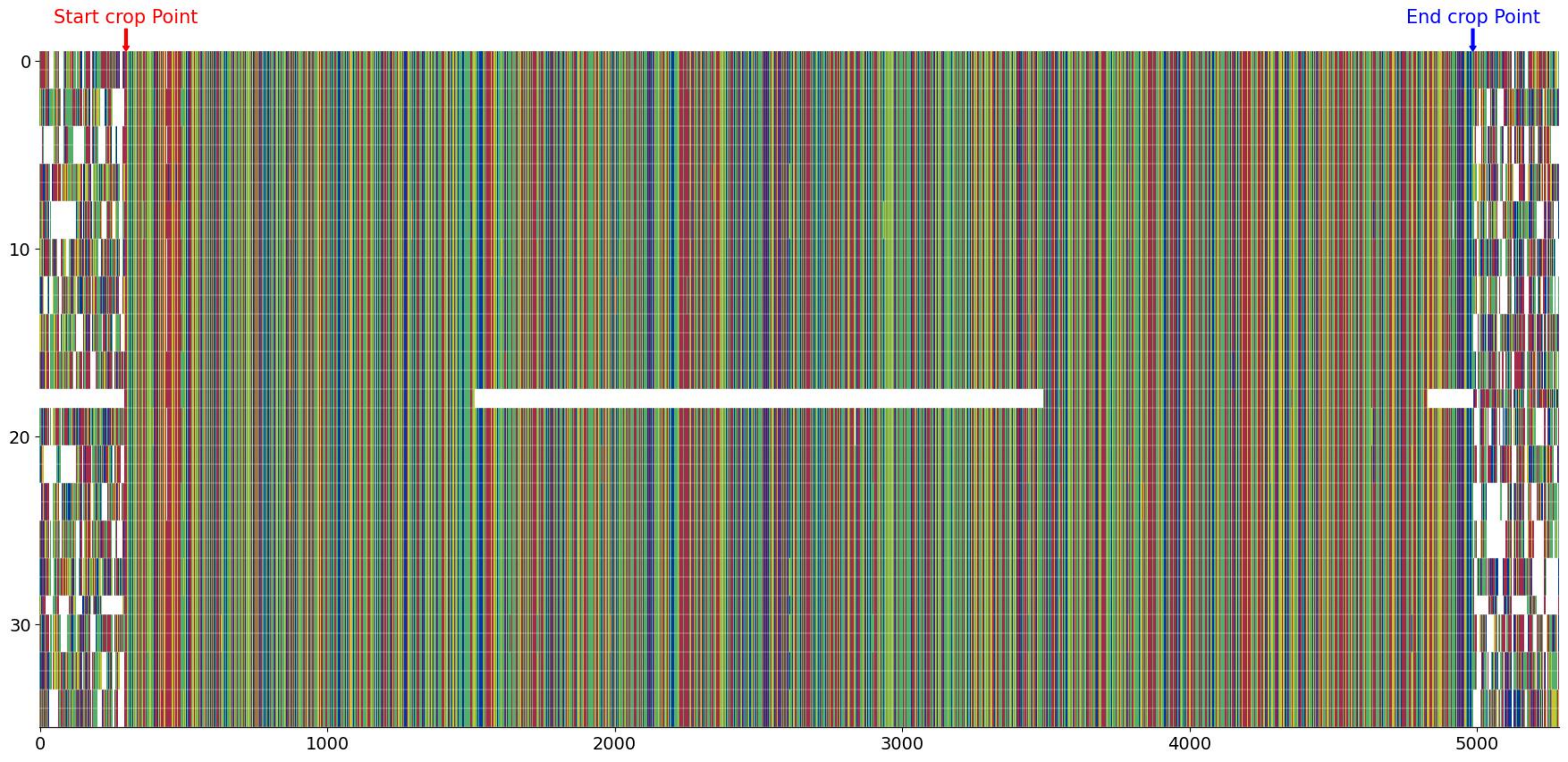
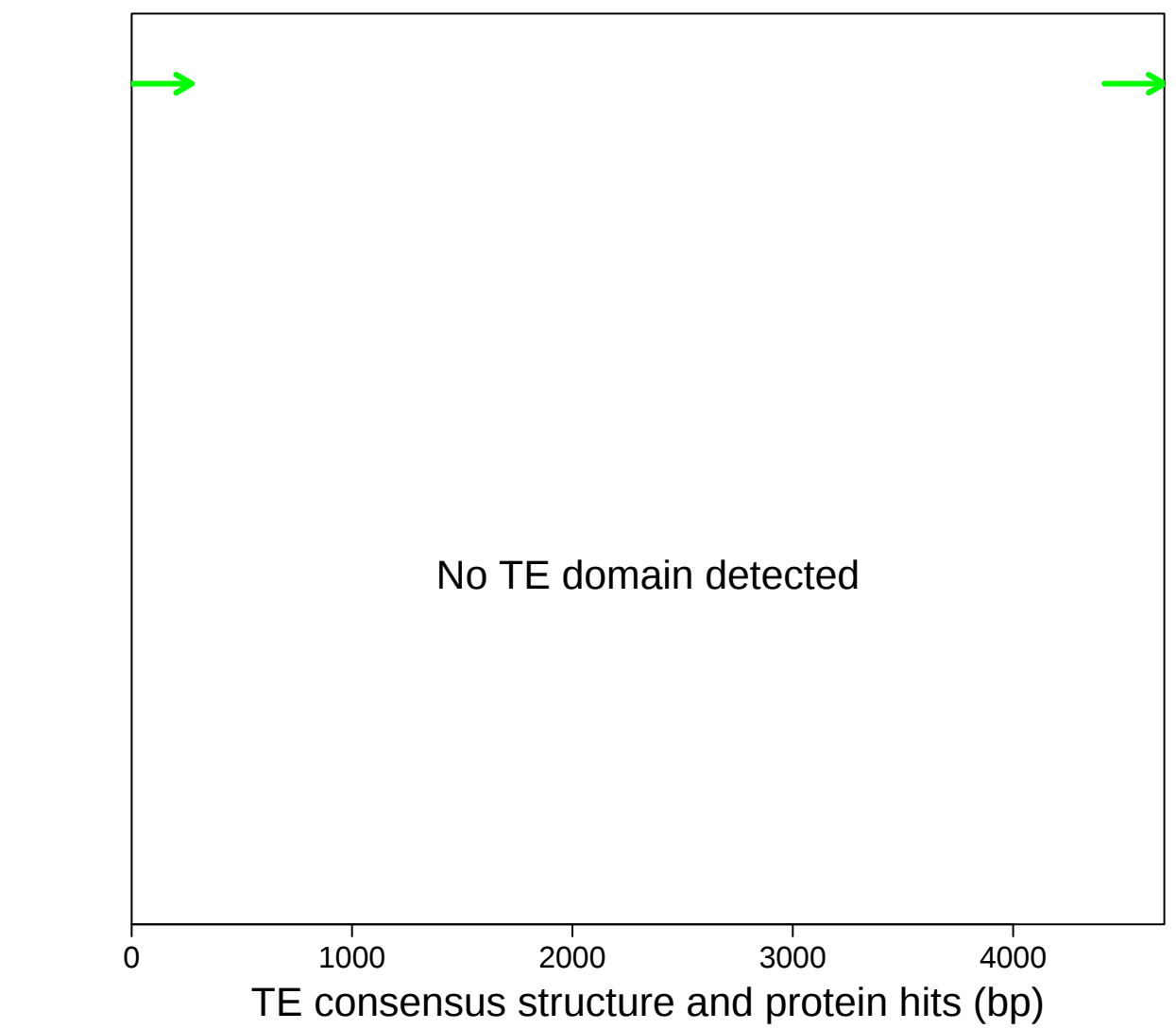
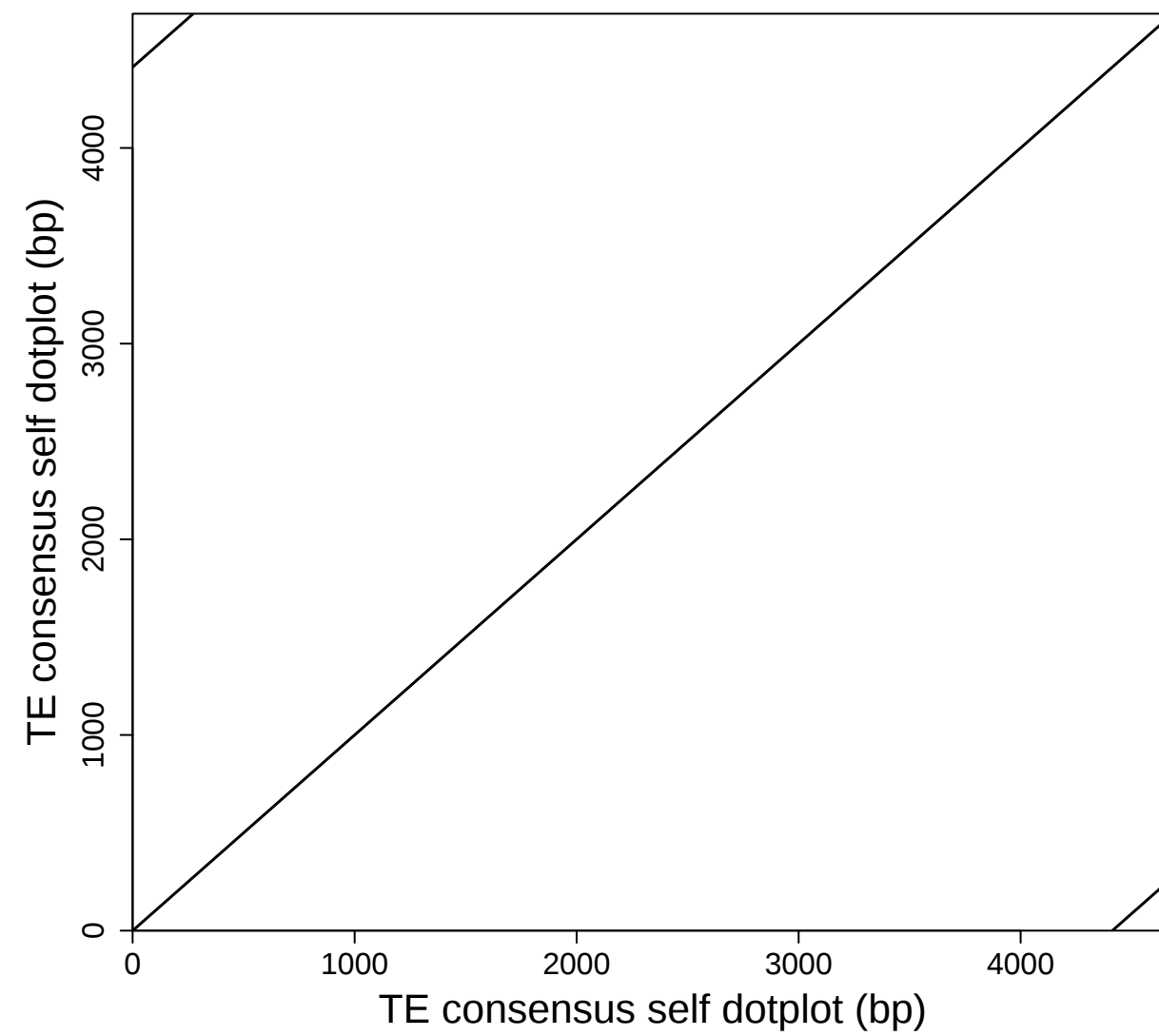
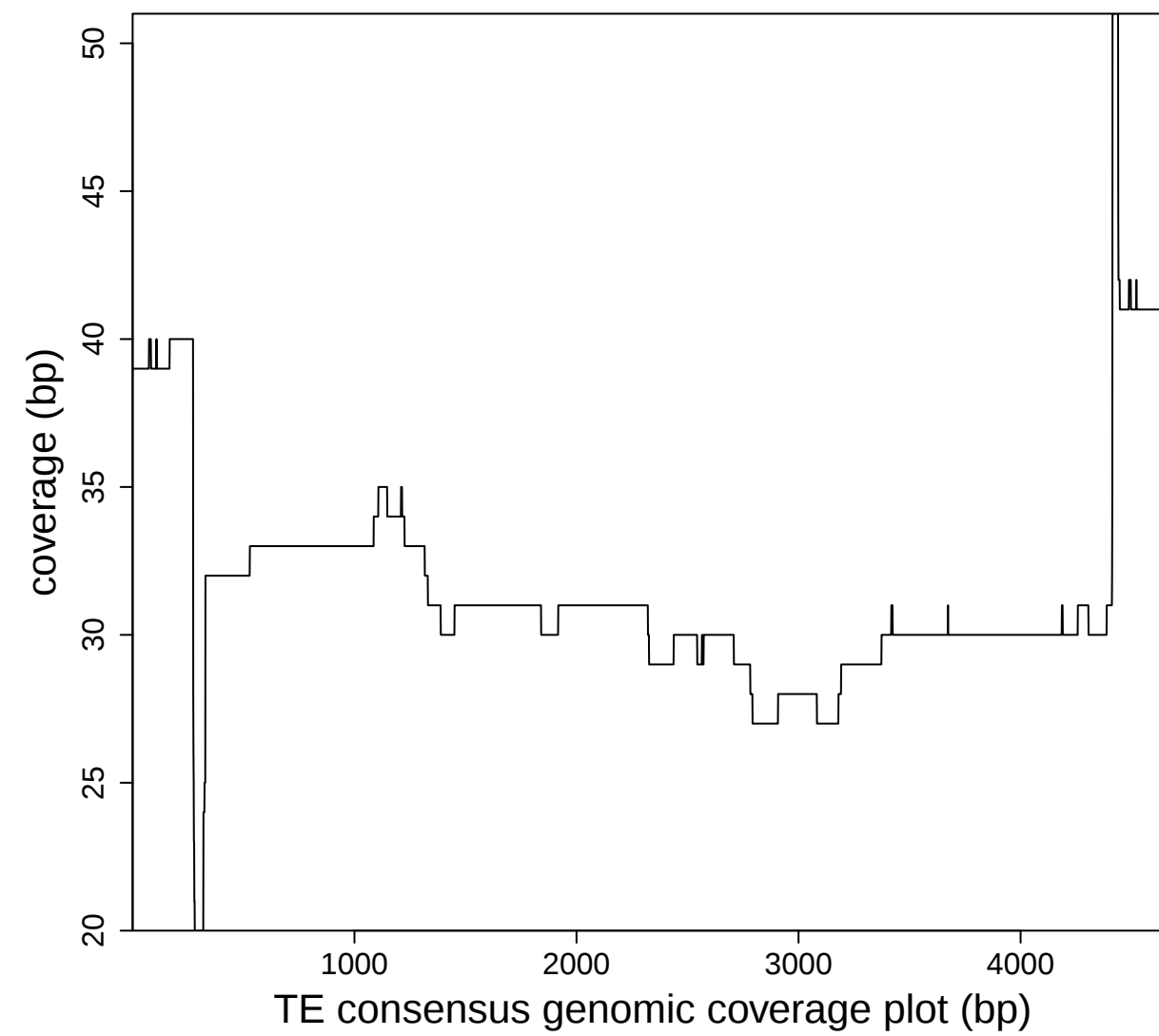
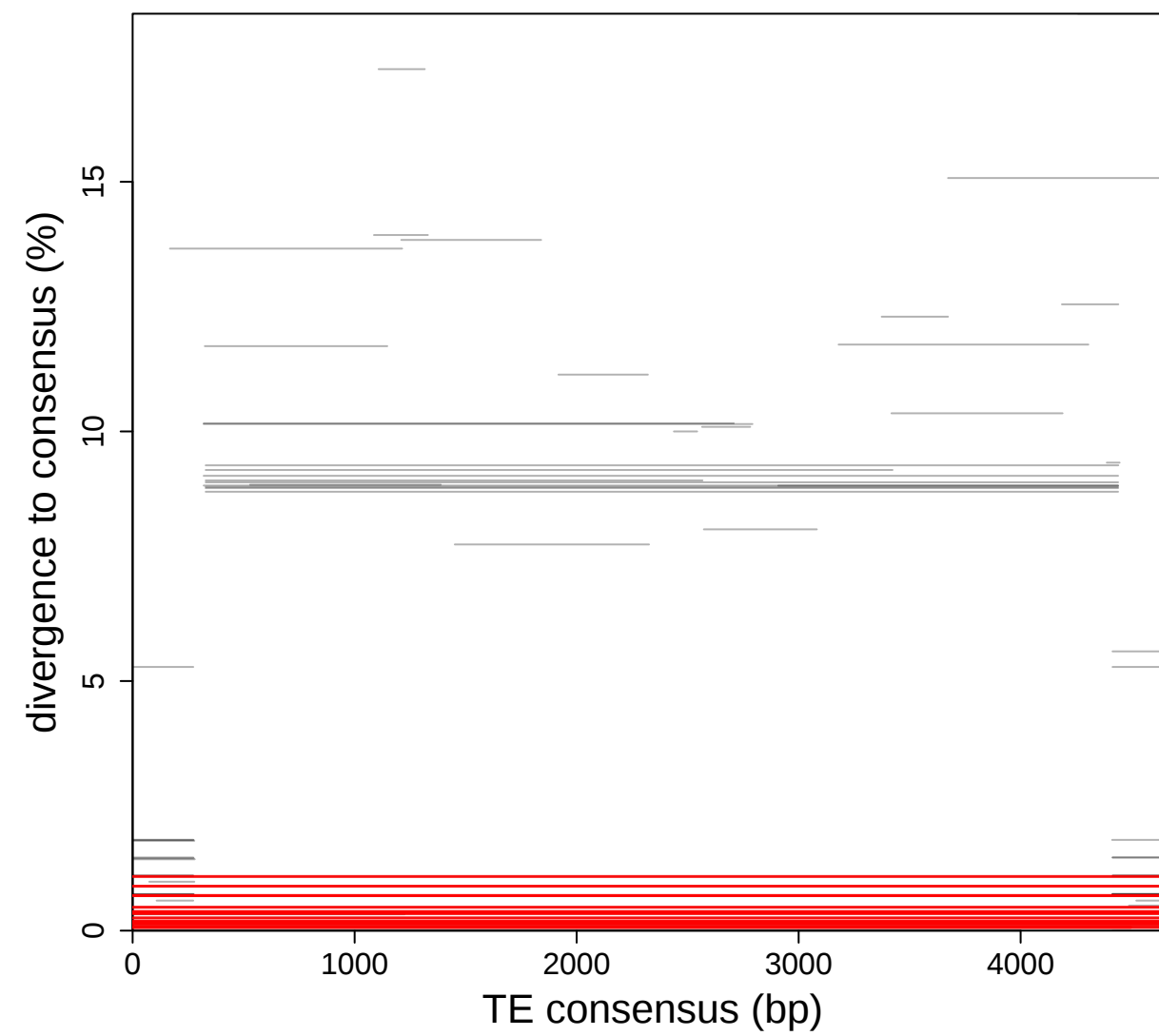






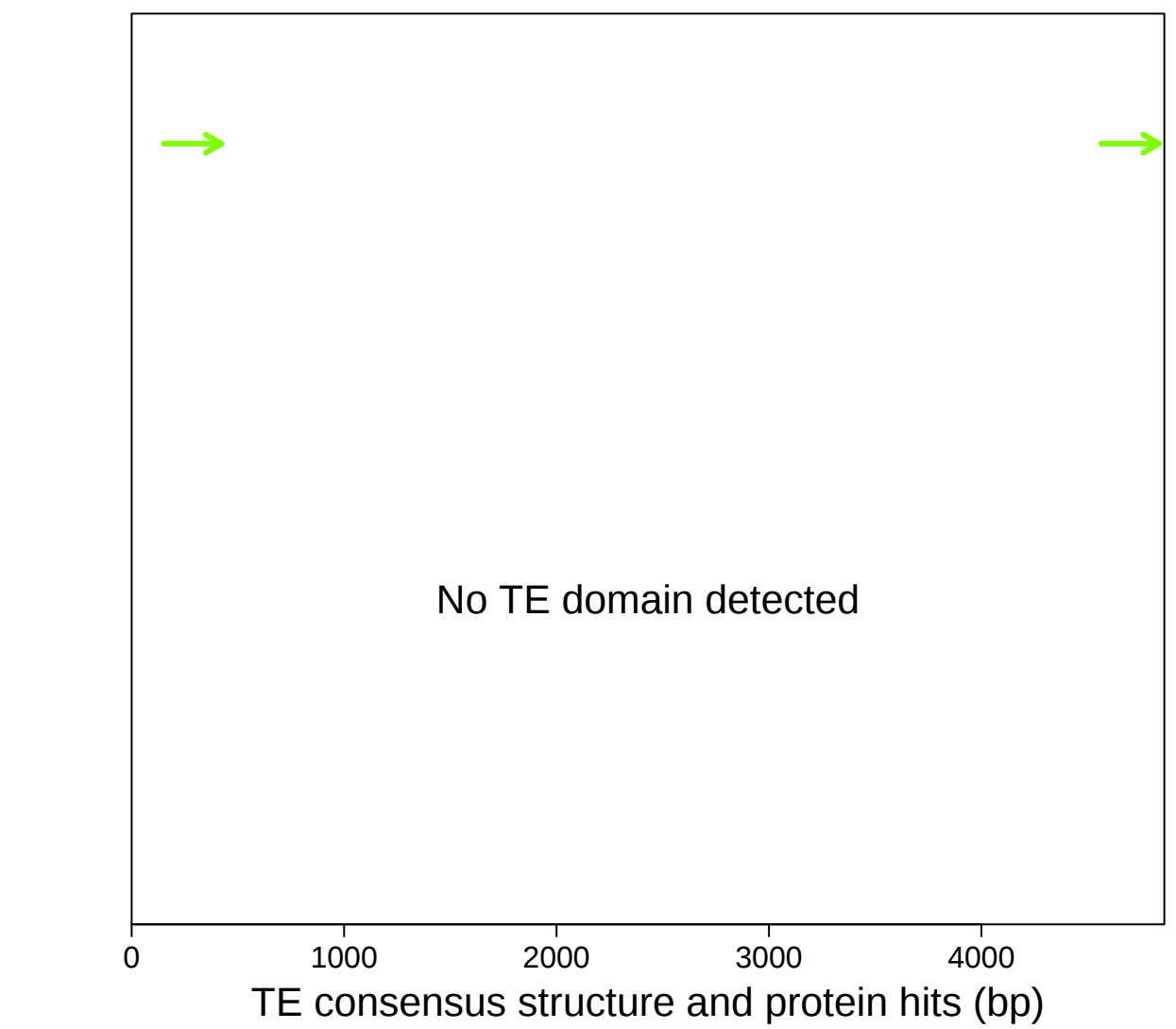
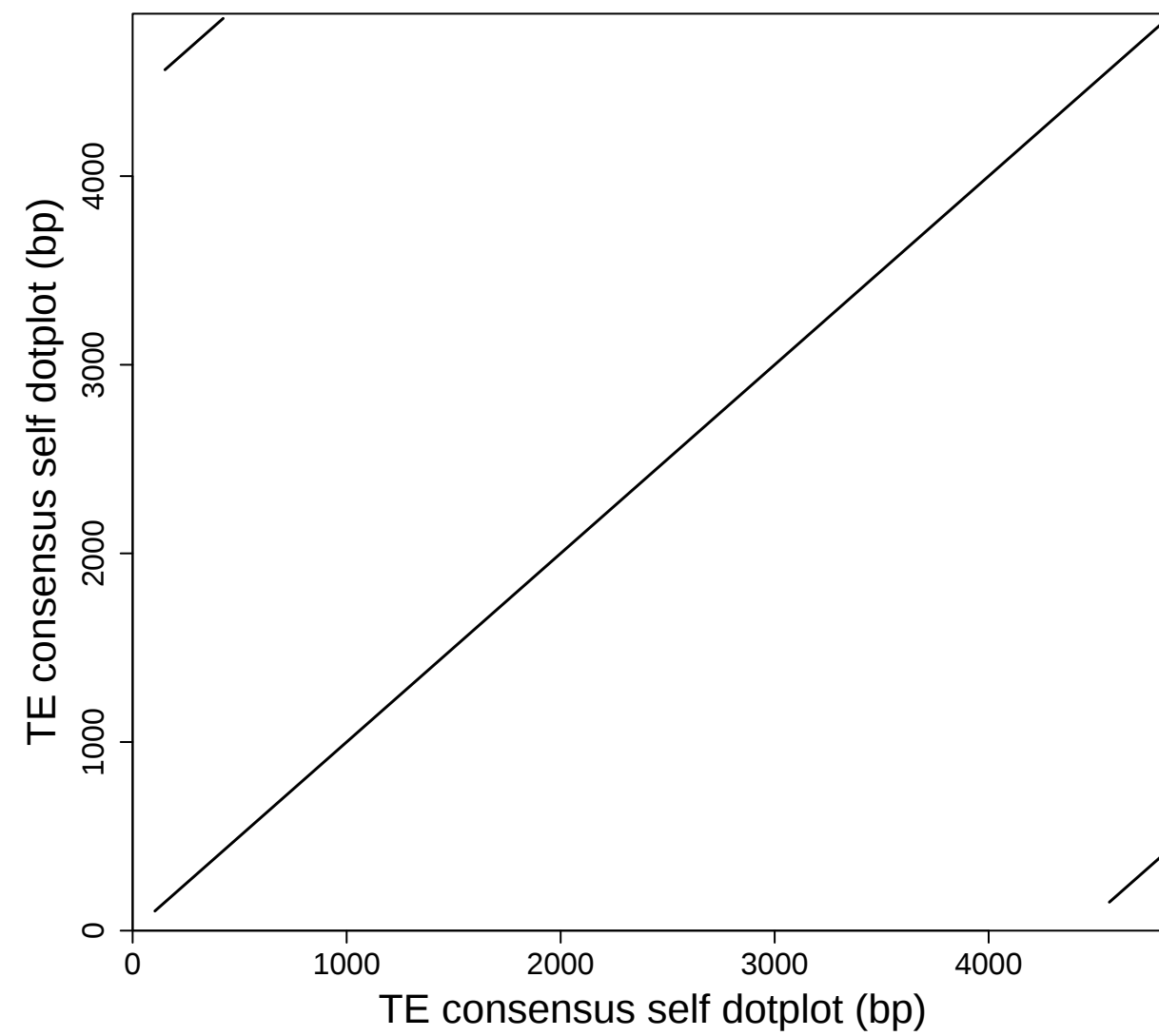
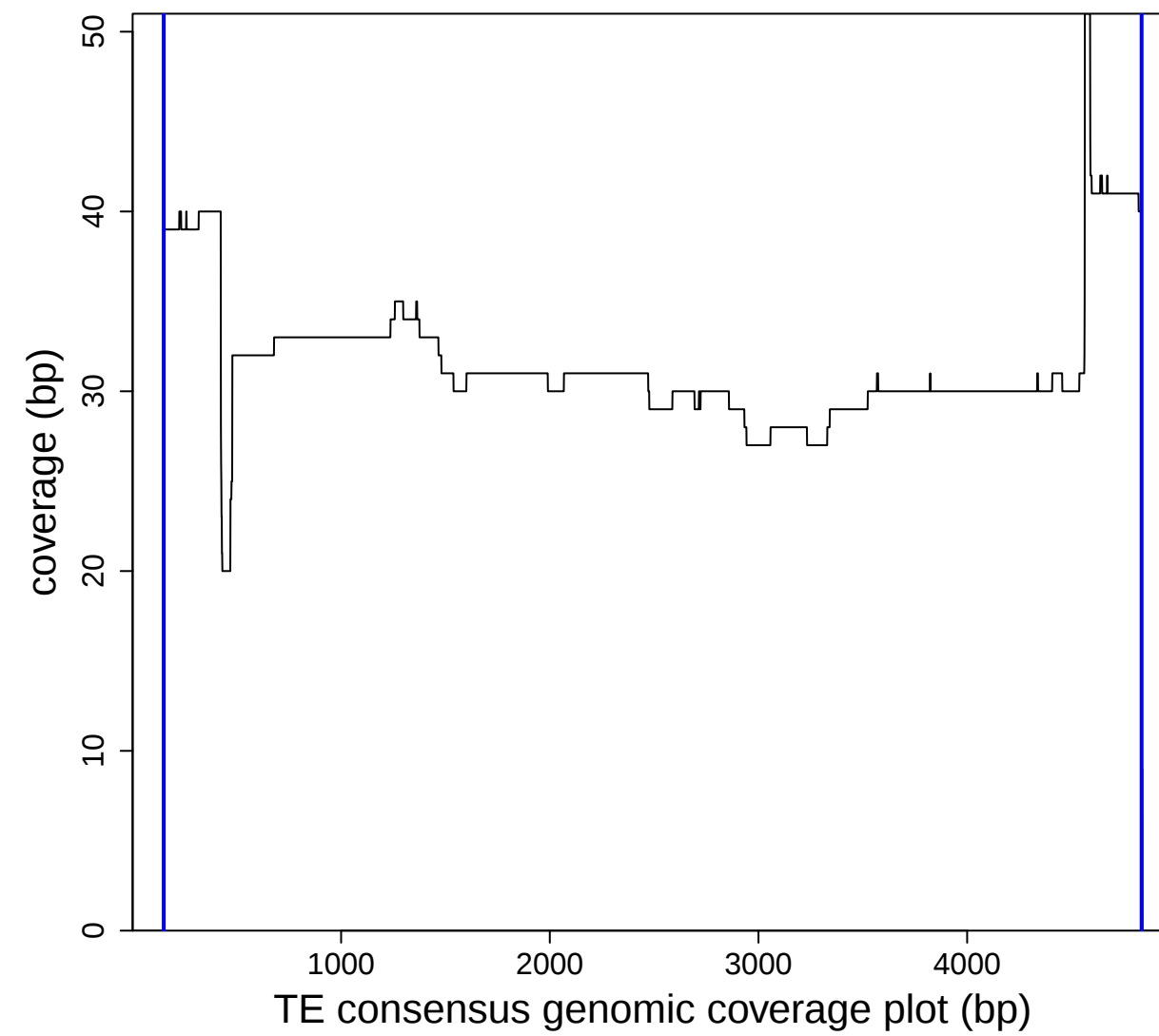
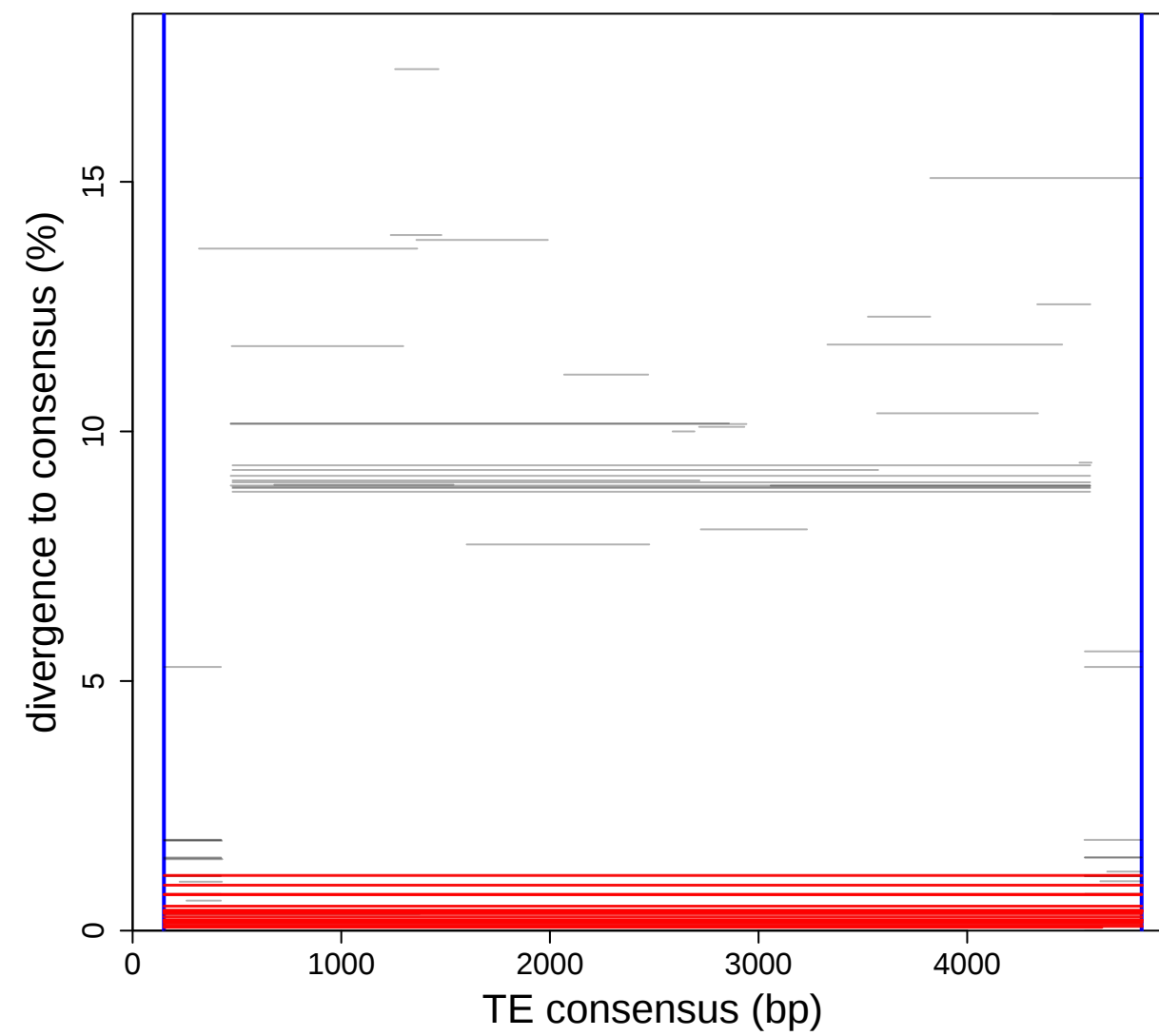
_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcIn.fa_al
size: 4686bp; fragments: 94; full length: 18 (>=4217.4bp)

After TEtrimmer 4686 bp

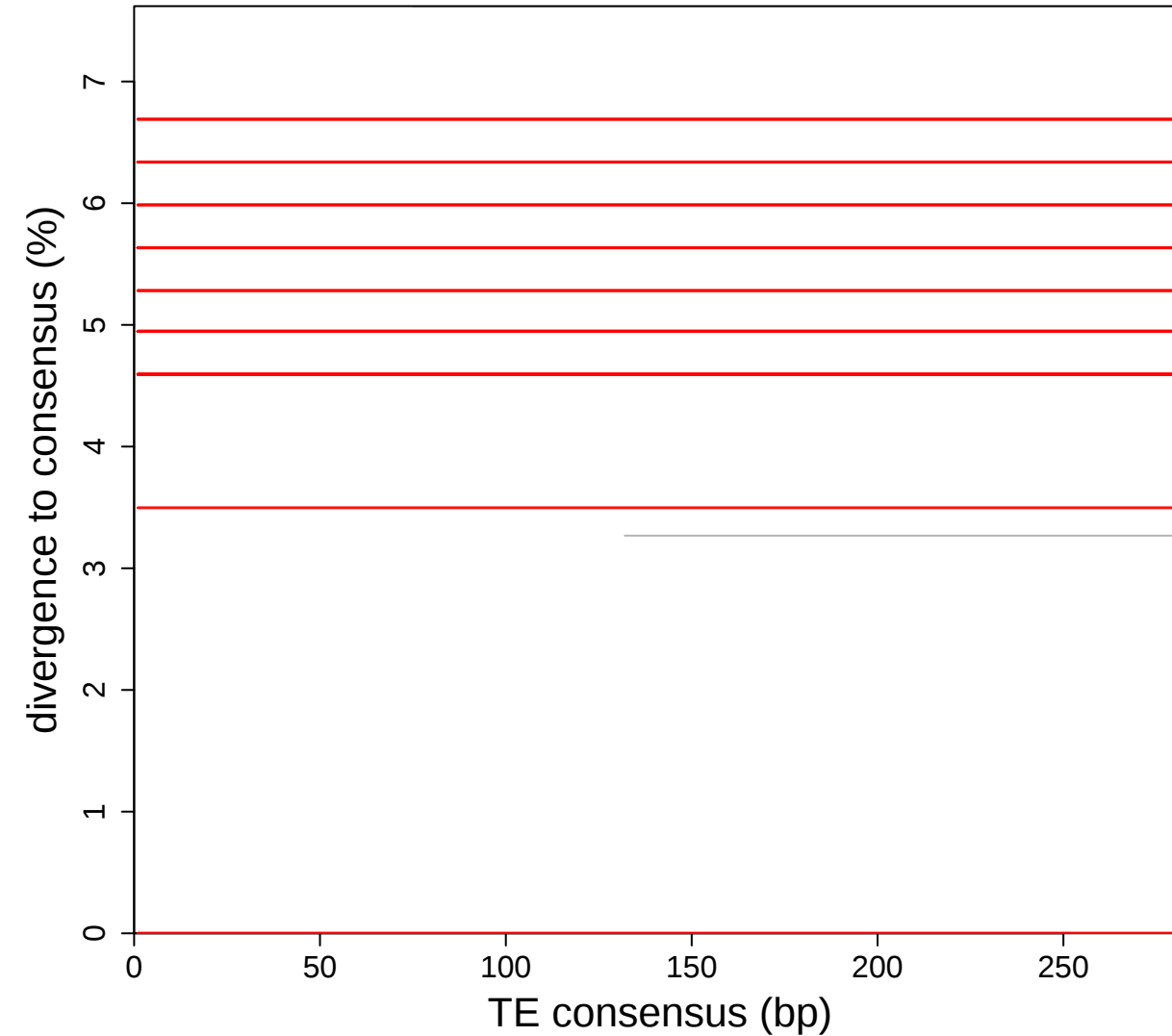


bed_g_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bclr
size: 4986bp; fragments: 94; full length: 18 (≥ 4487.4 bp)

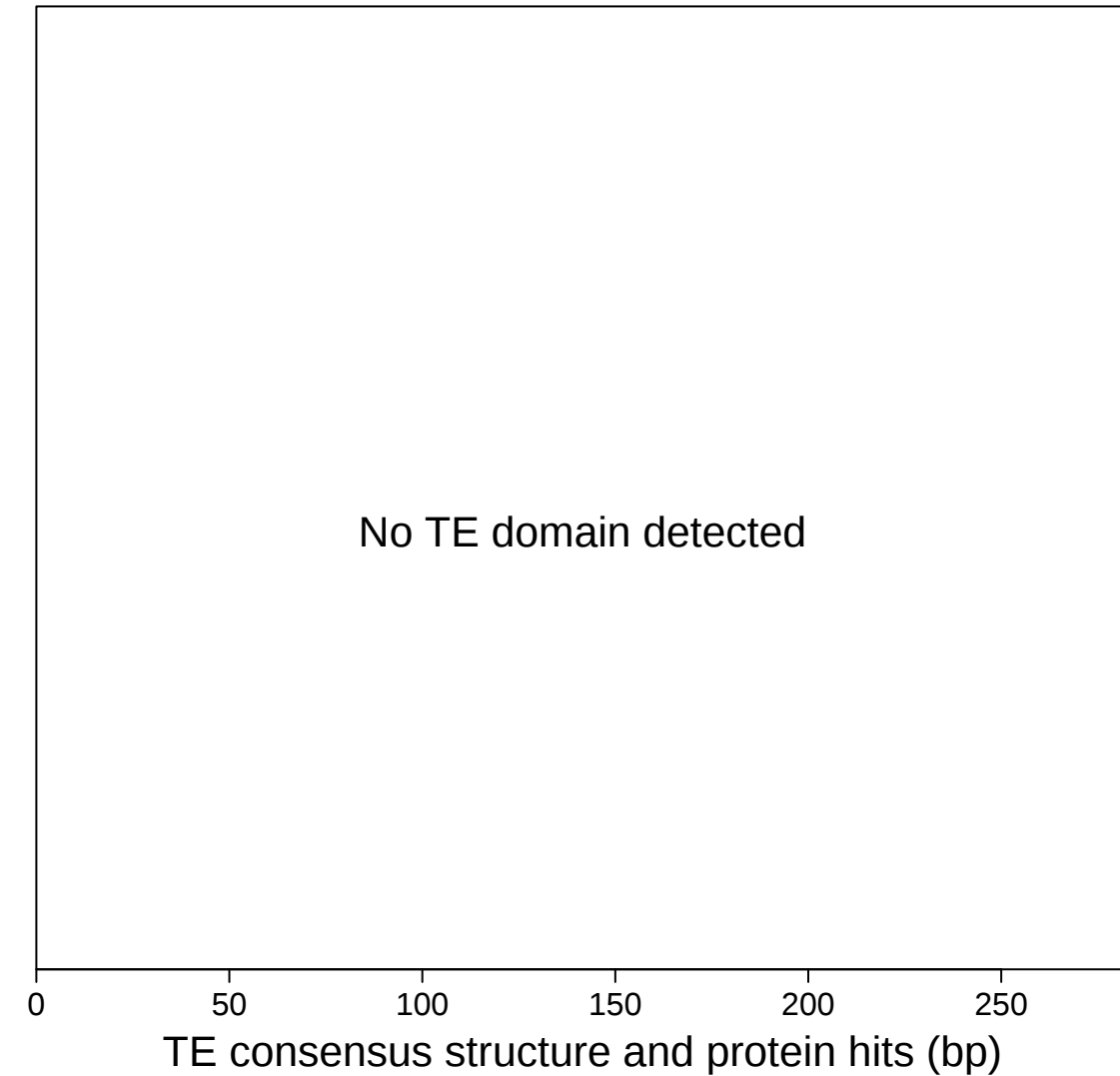
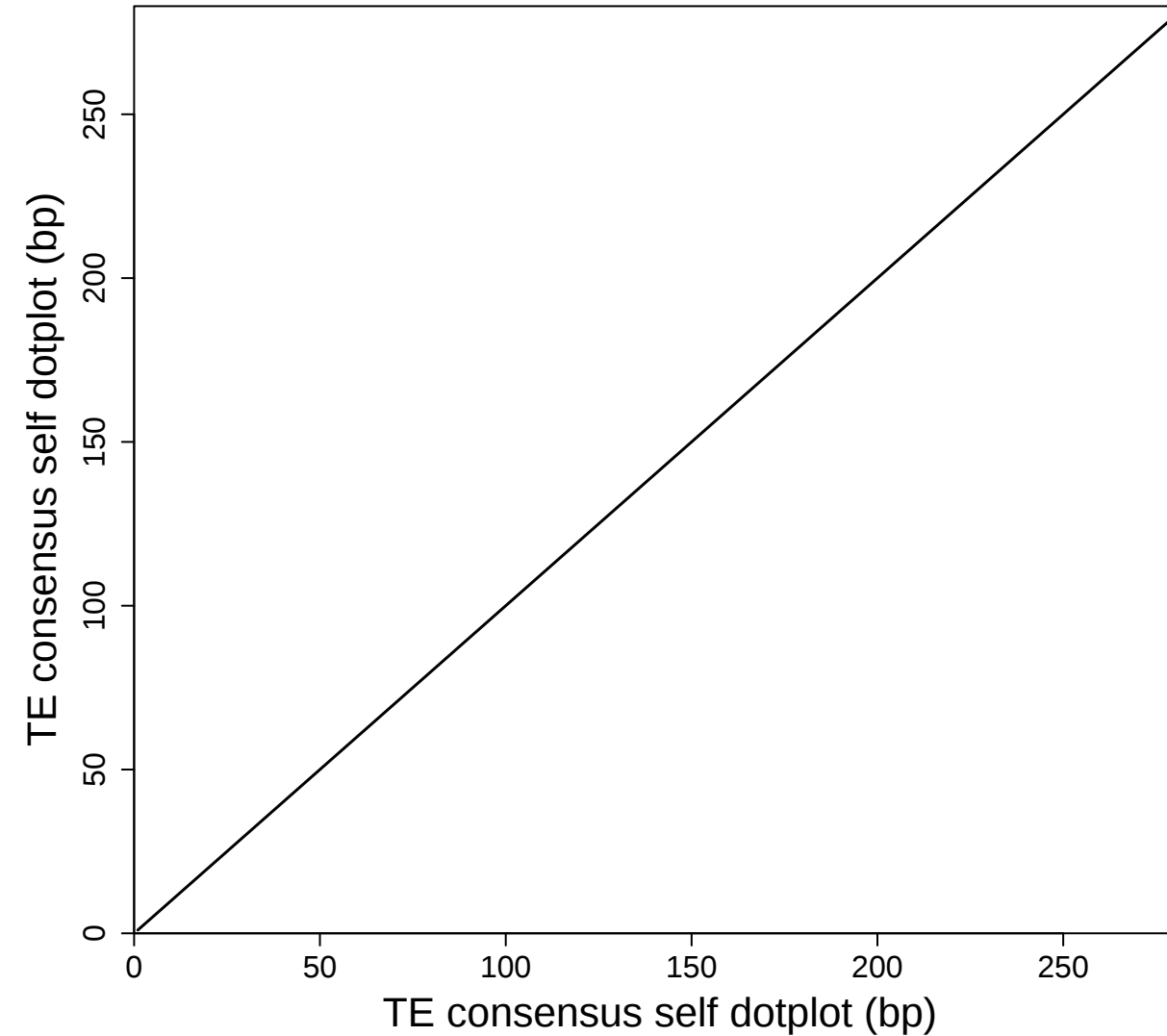
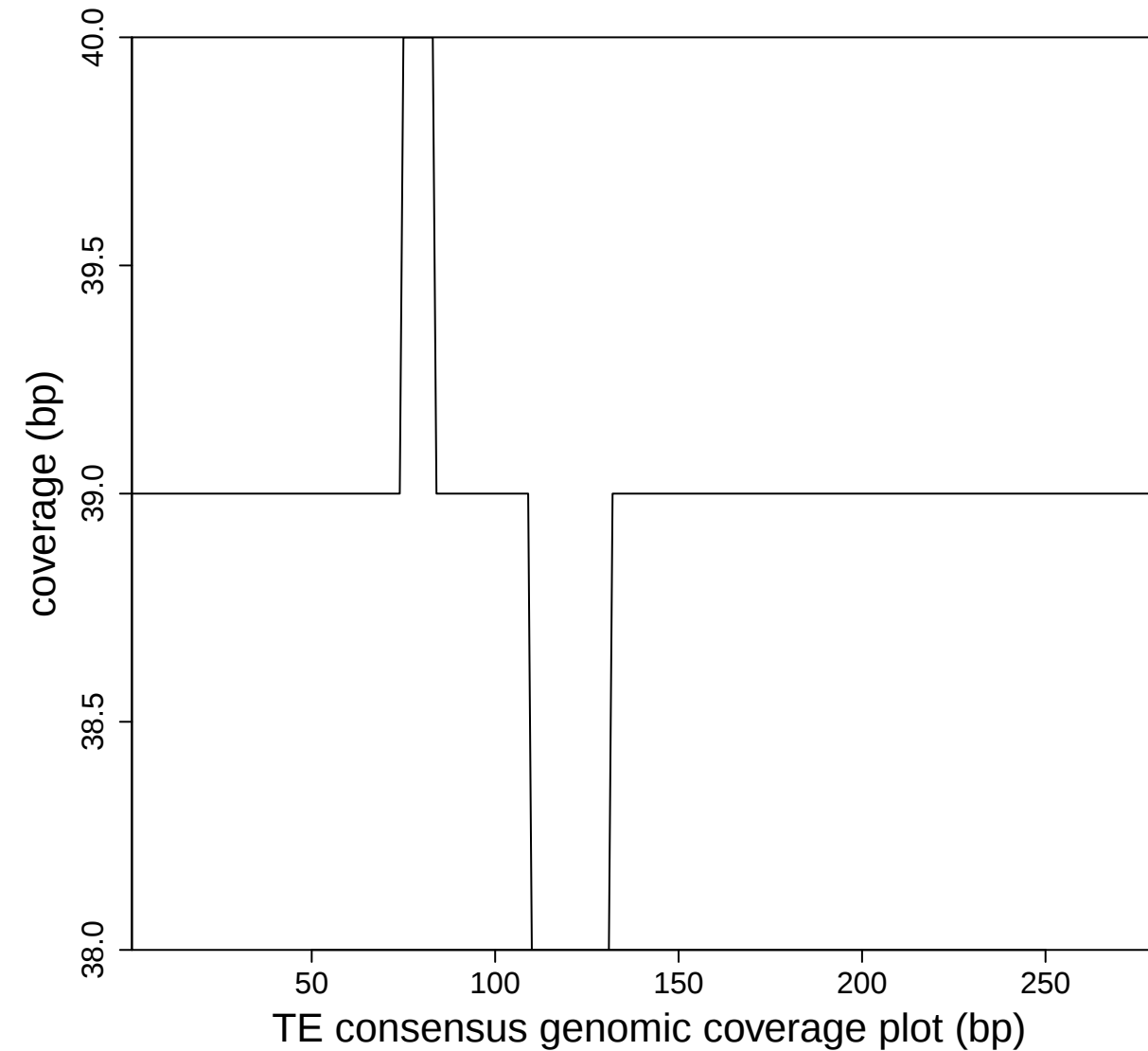
After TEtrimmer Extended plot Blue lines are boundaries



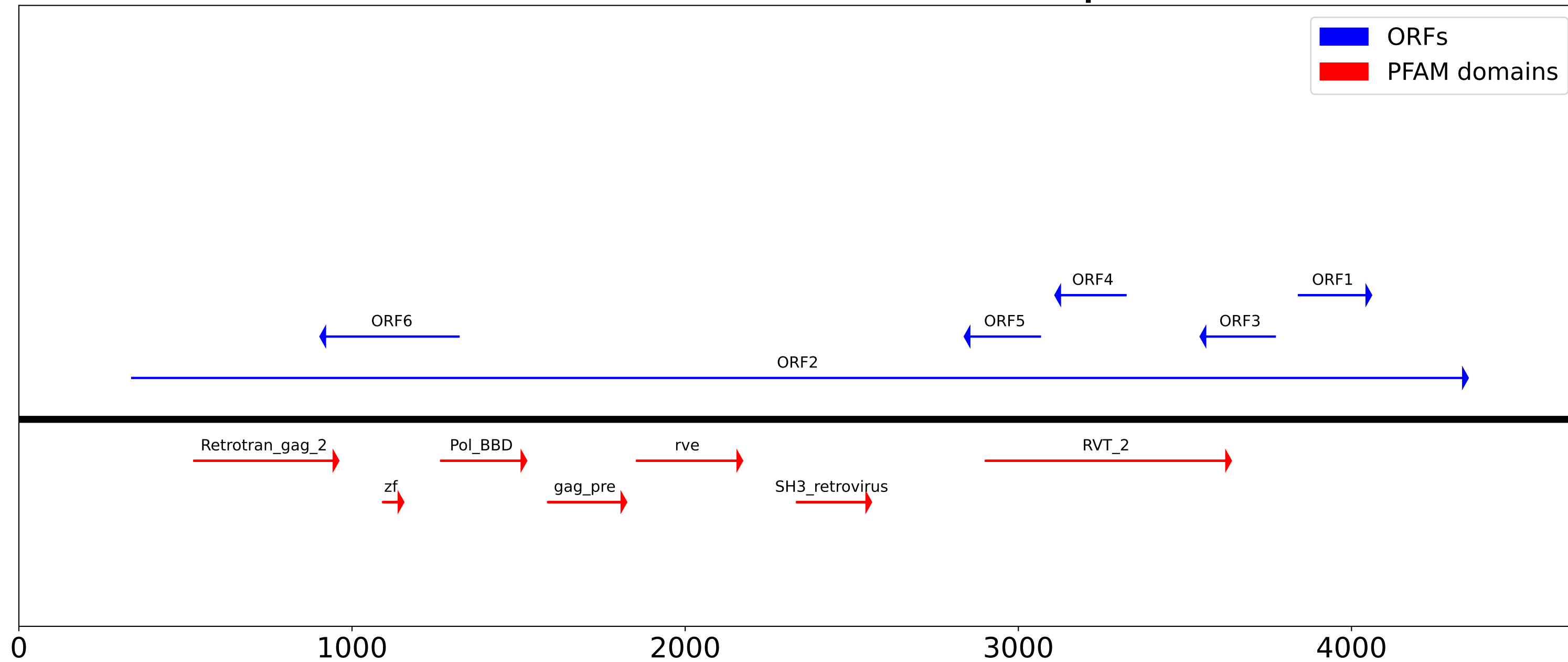
TE: ltr_1_family_155
size: 283bp; fragments: 41; full length: 37 (≥ 254.7 bp)



Before TEtrimmer 283 bp



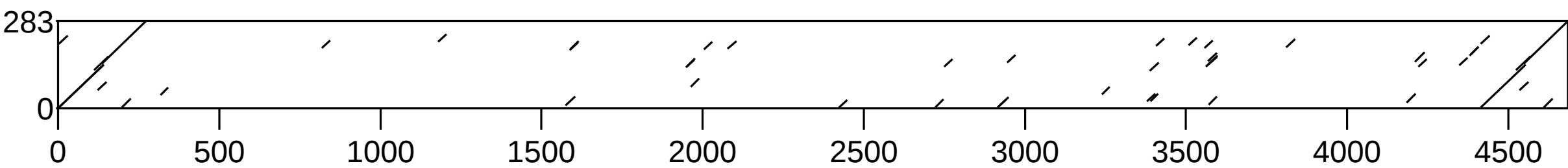
After TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)