

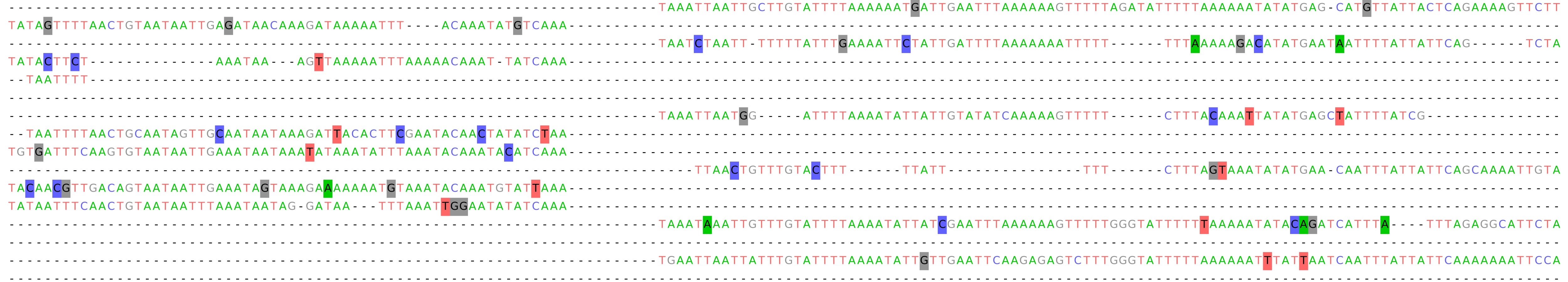
Start crop Point

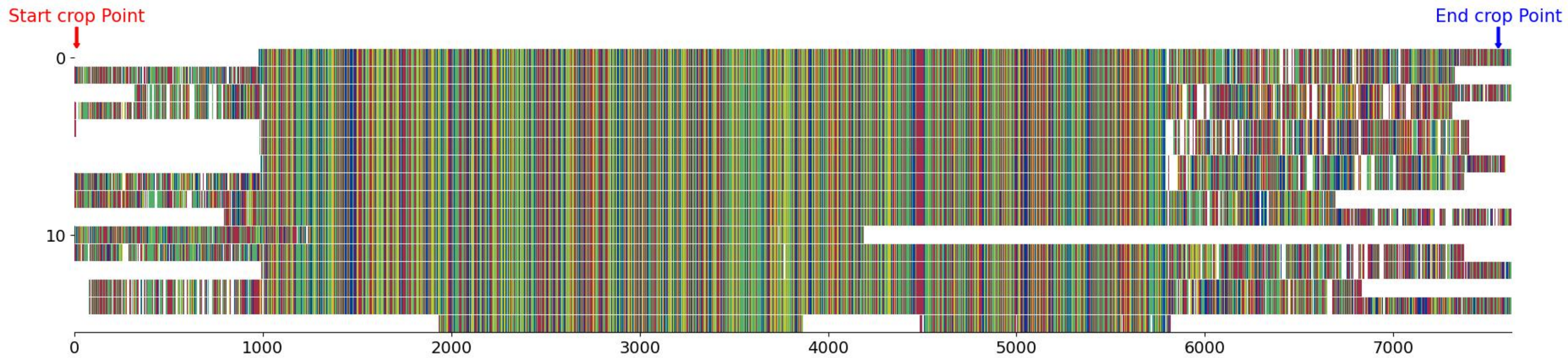


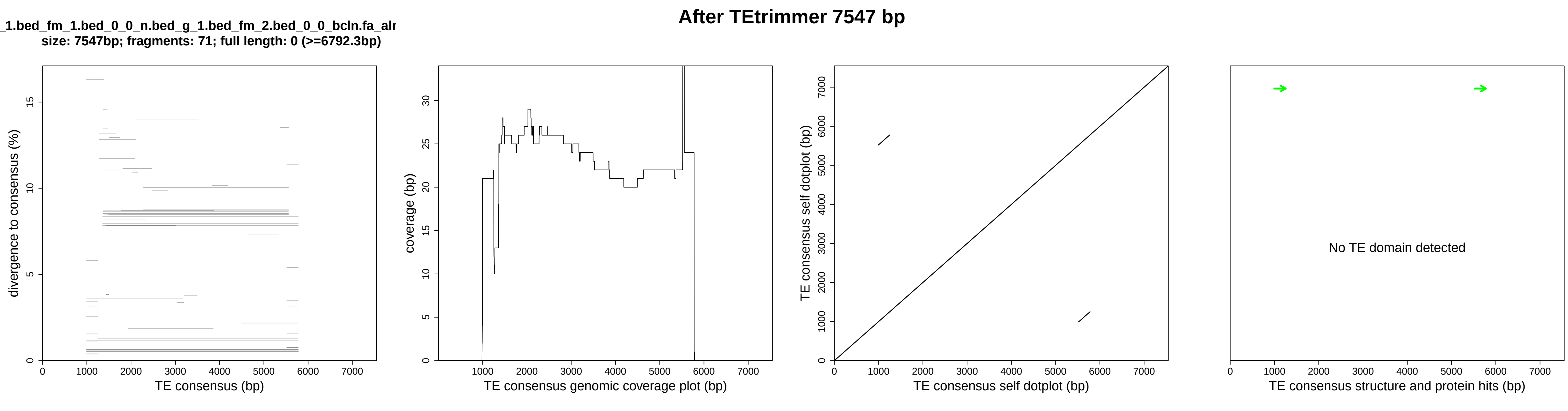
End crop Point



MSA length = 7547

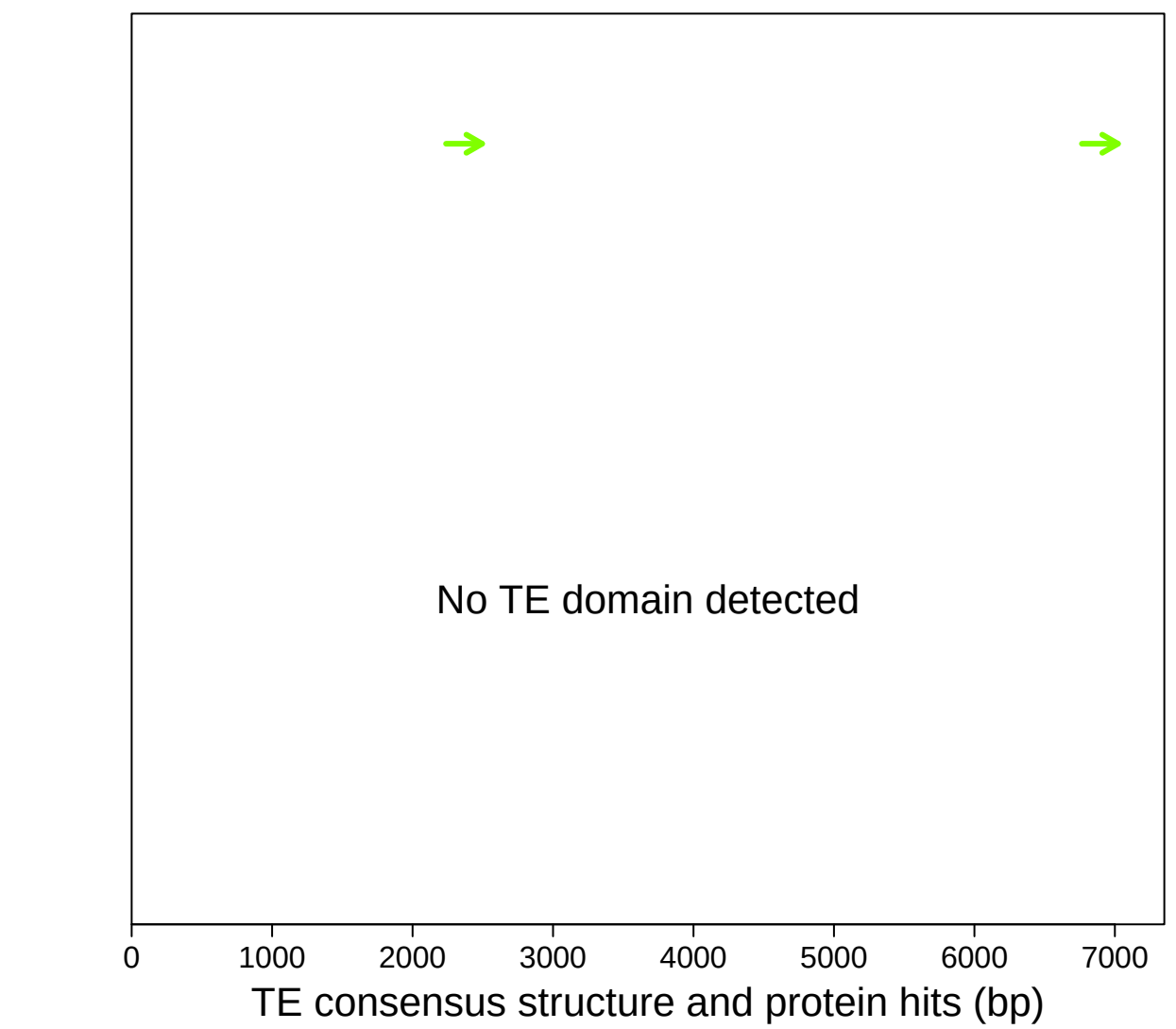
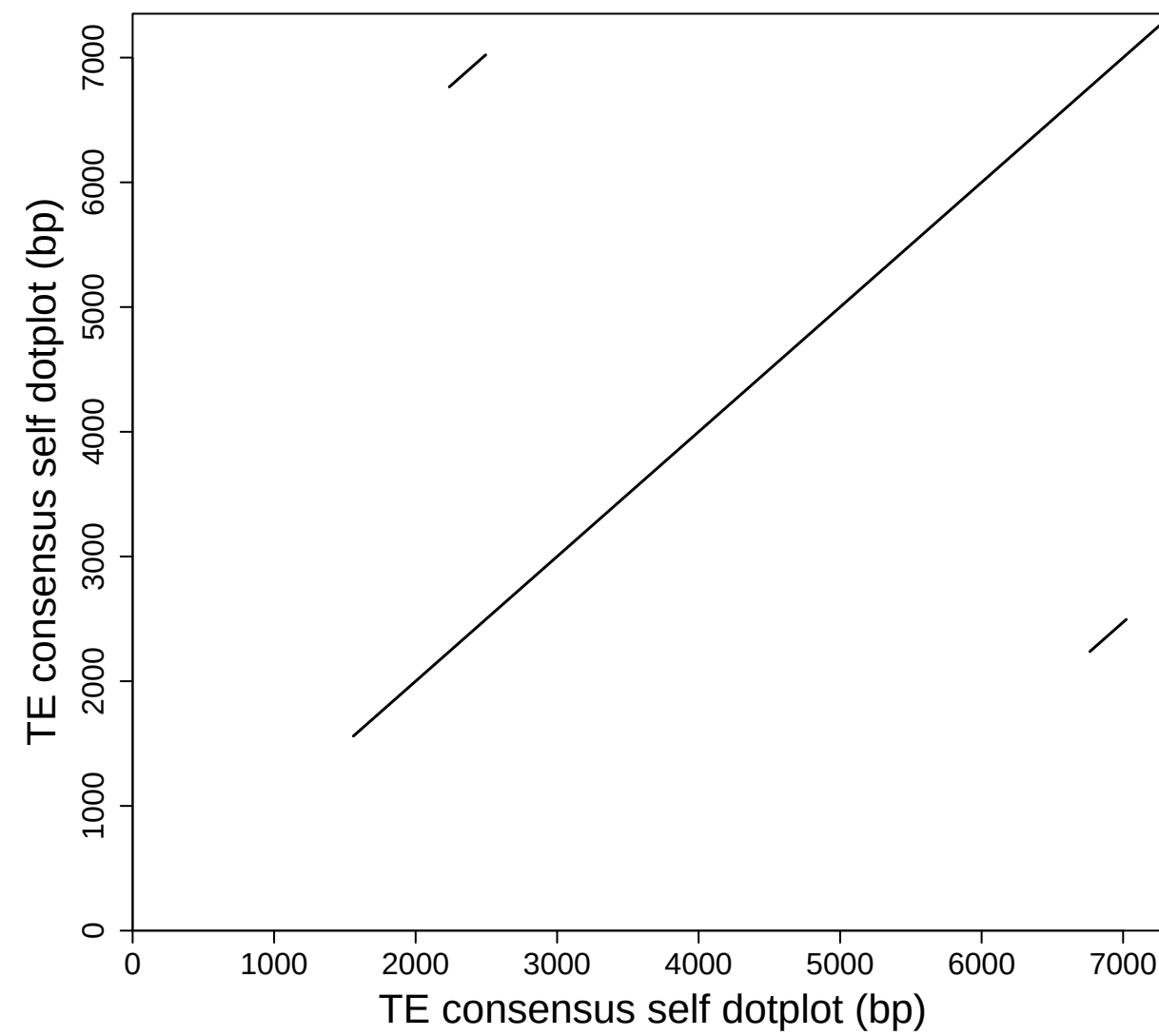
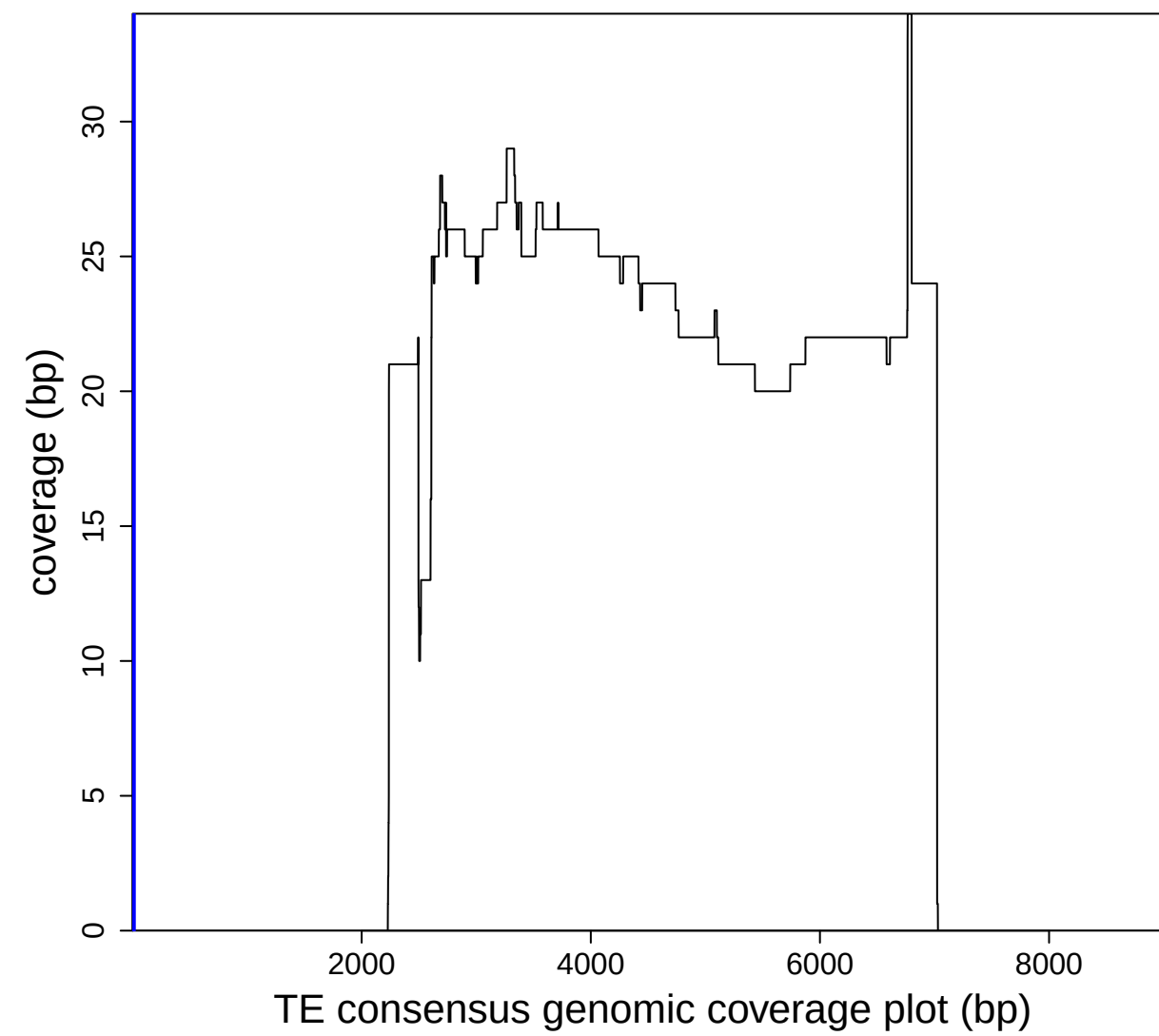
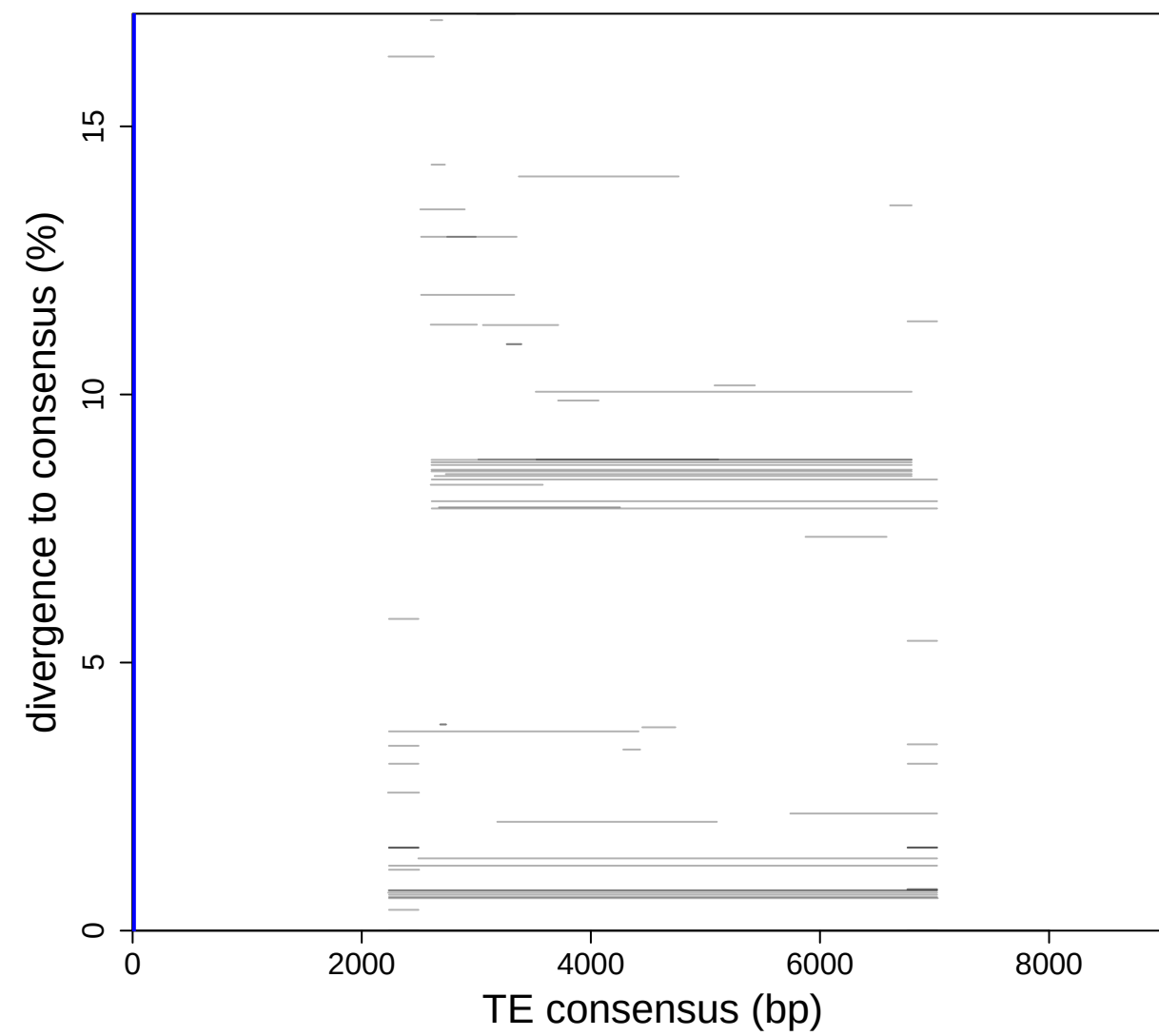






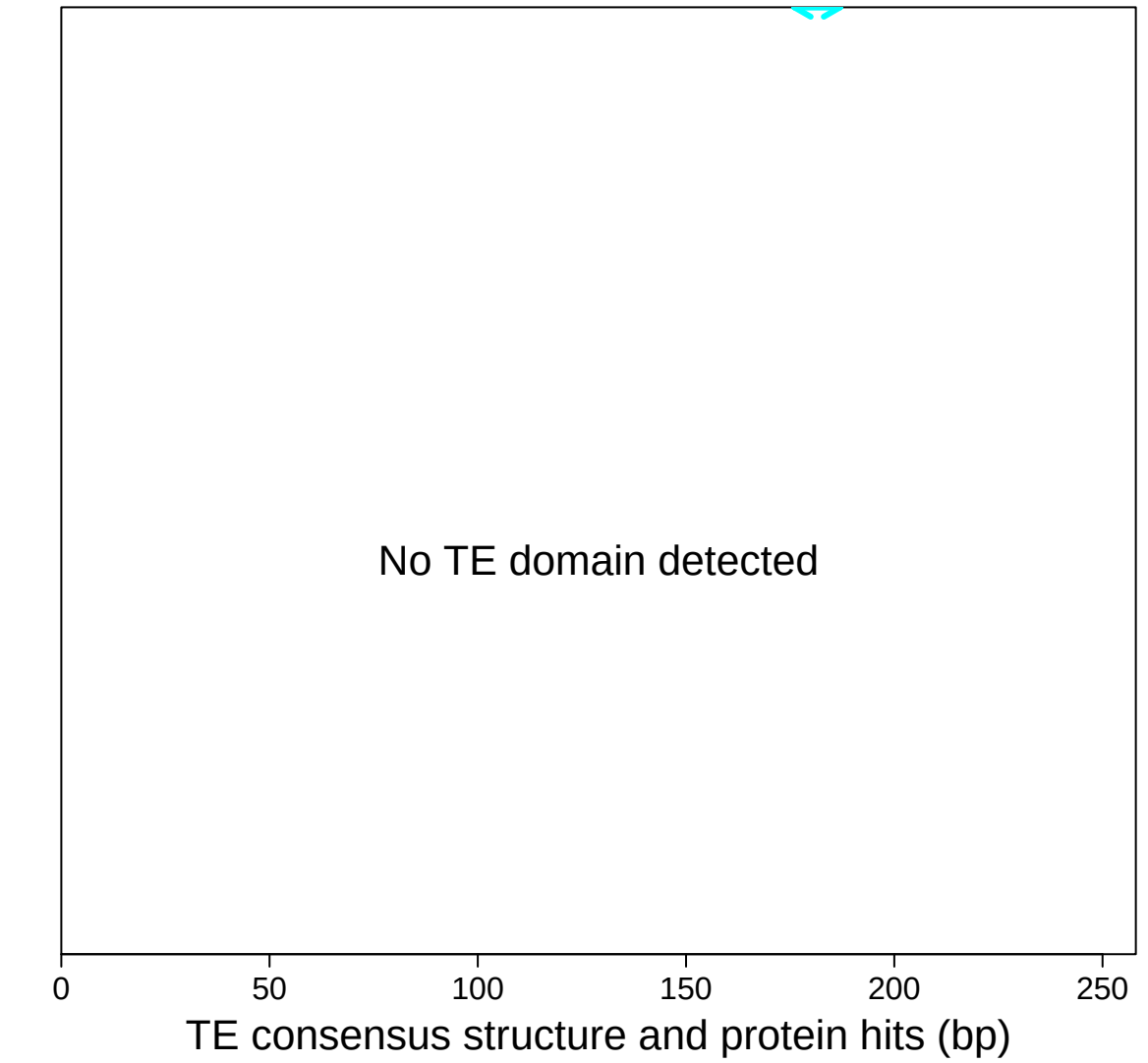
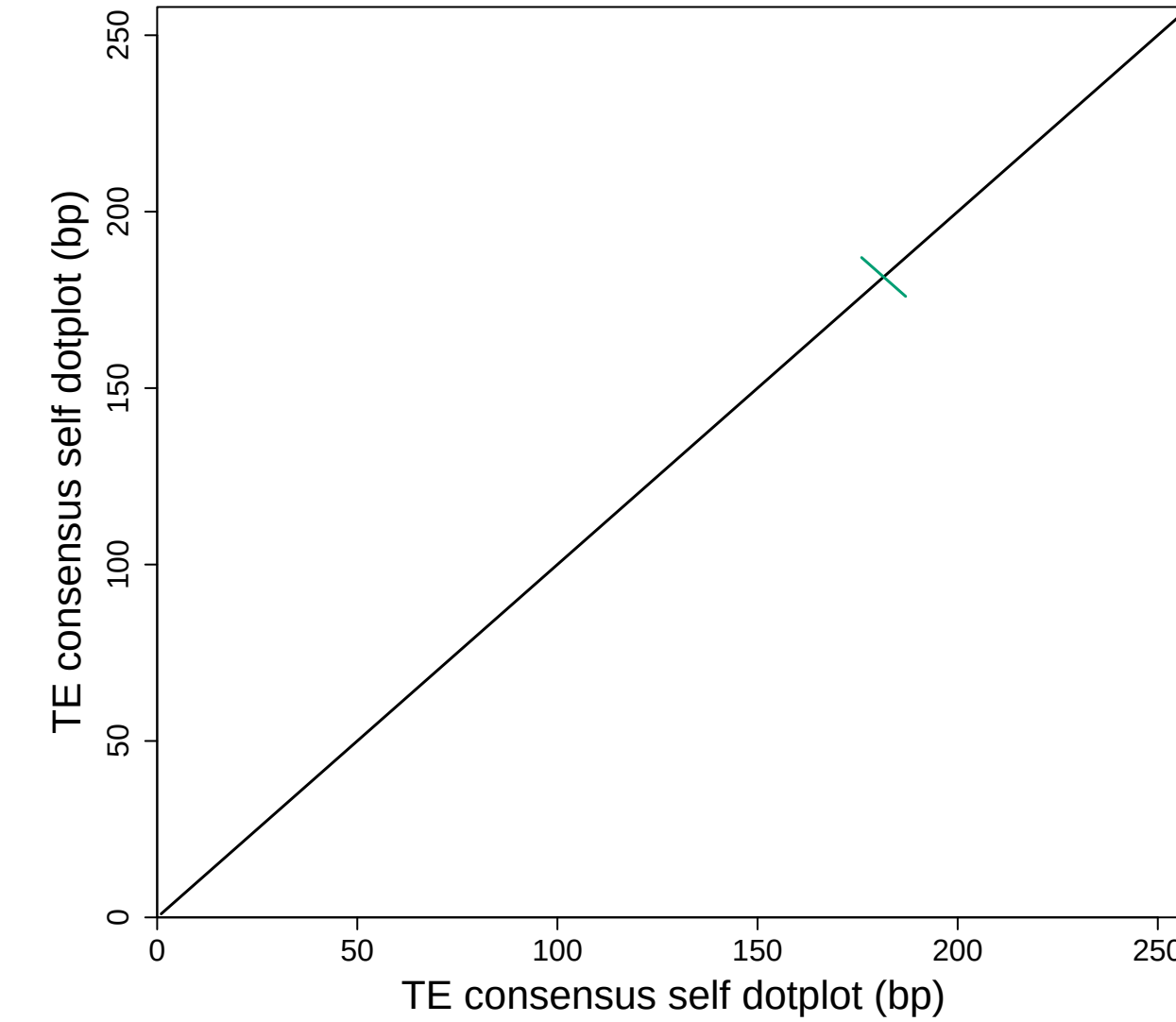
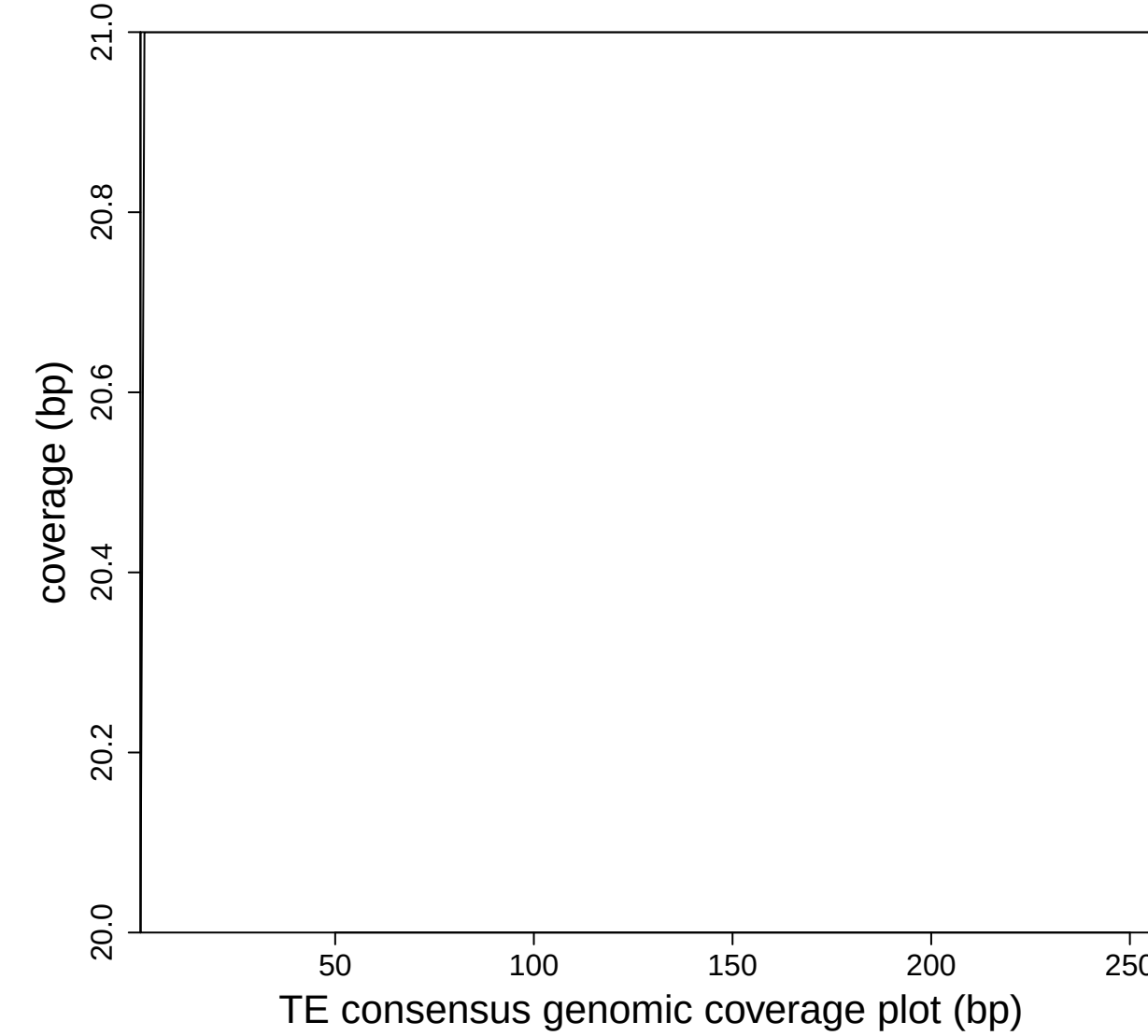
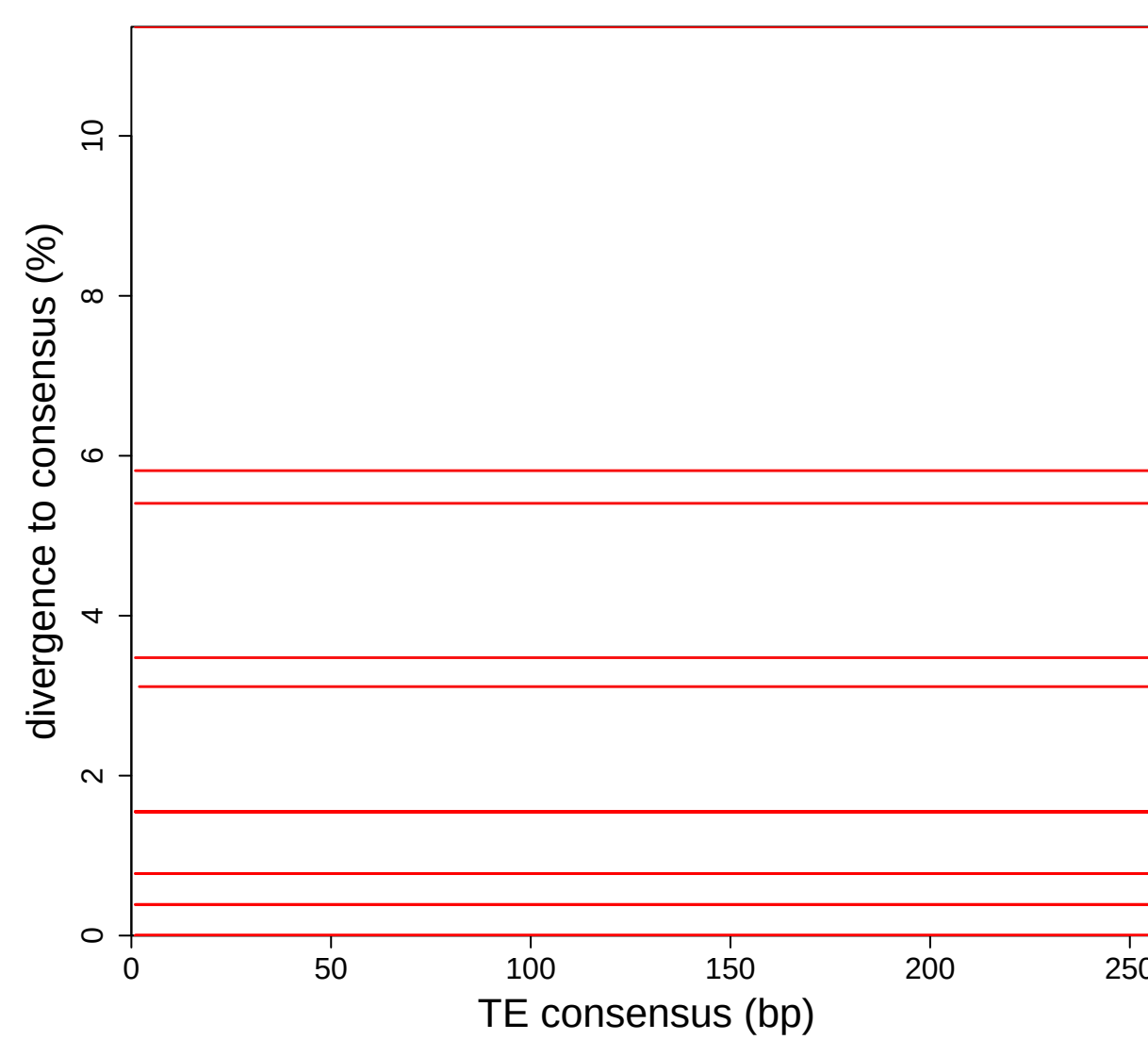
bed_g_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcln
size: 9081bp; fragments: 71; full length: 0 (>=8172.9bp)

After TEtrimmer Extended plot Blue lines are boundaries

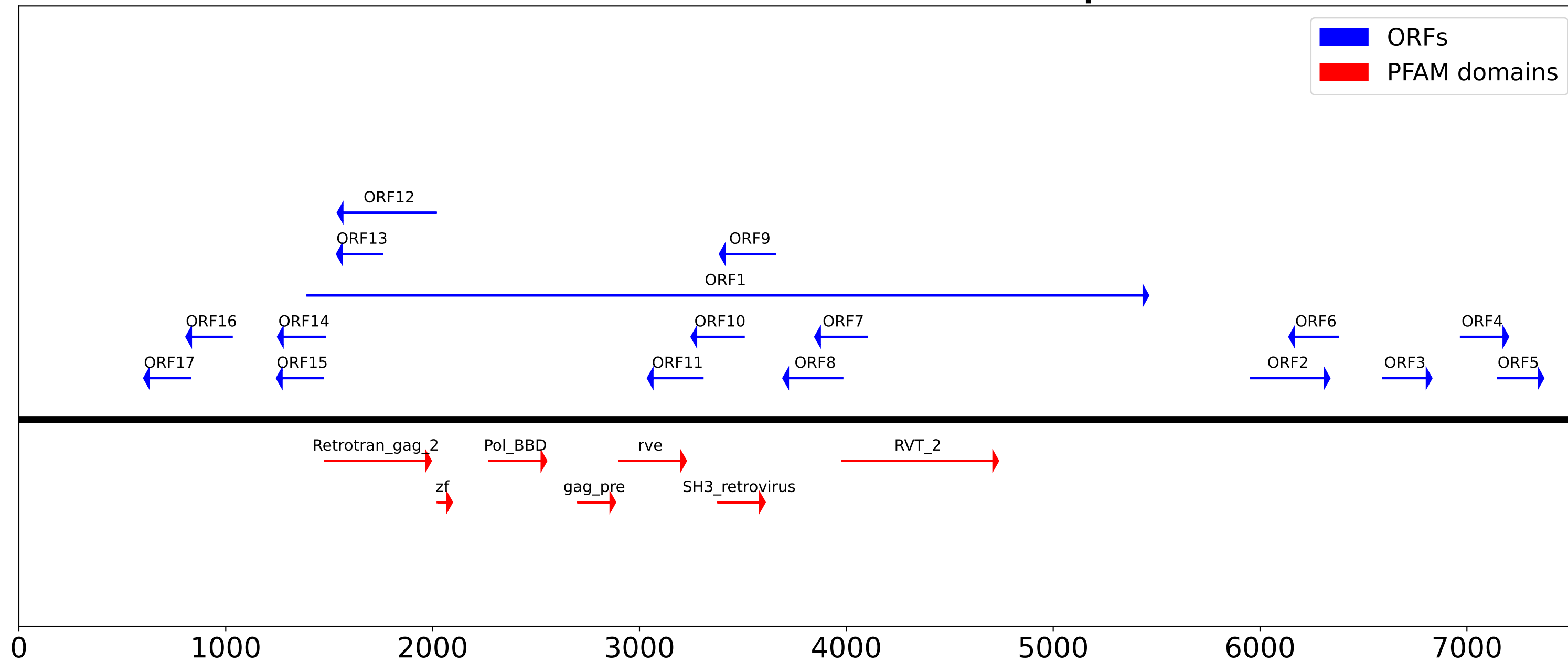


Before TEtrimmer 258 bp

TE: ltr_1_family_50
size: 258bp; fragments: 21; full length: 21 (≥ 232.2 bp)



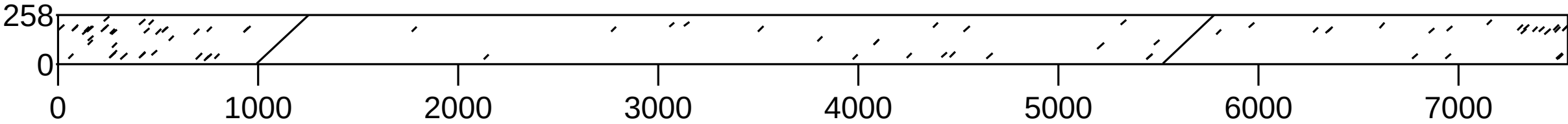
After TEtrimmer ORF and PFAM domain plot



Dotplot

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

nsus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)