

Start crop Point

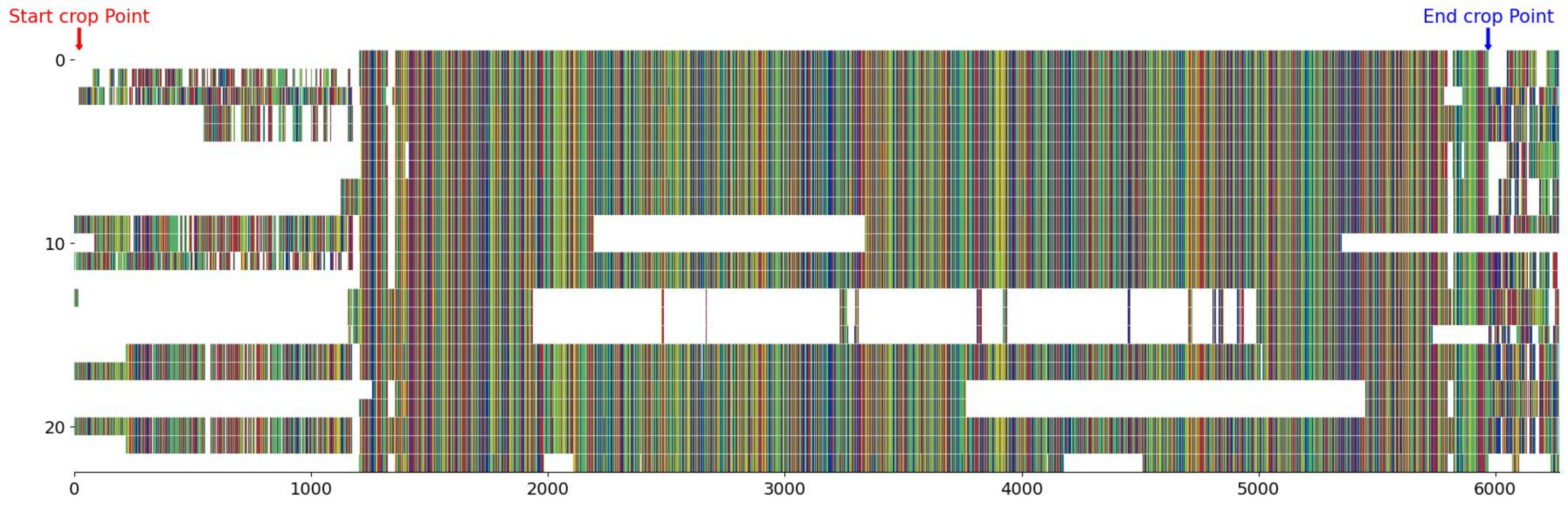


End crop Point



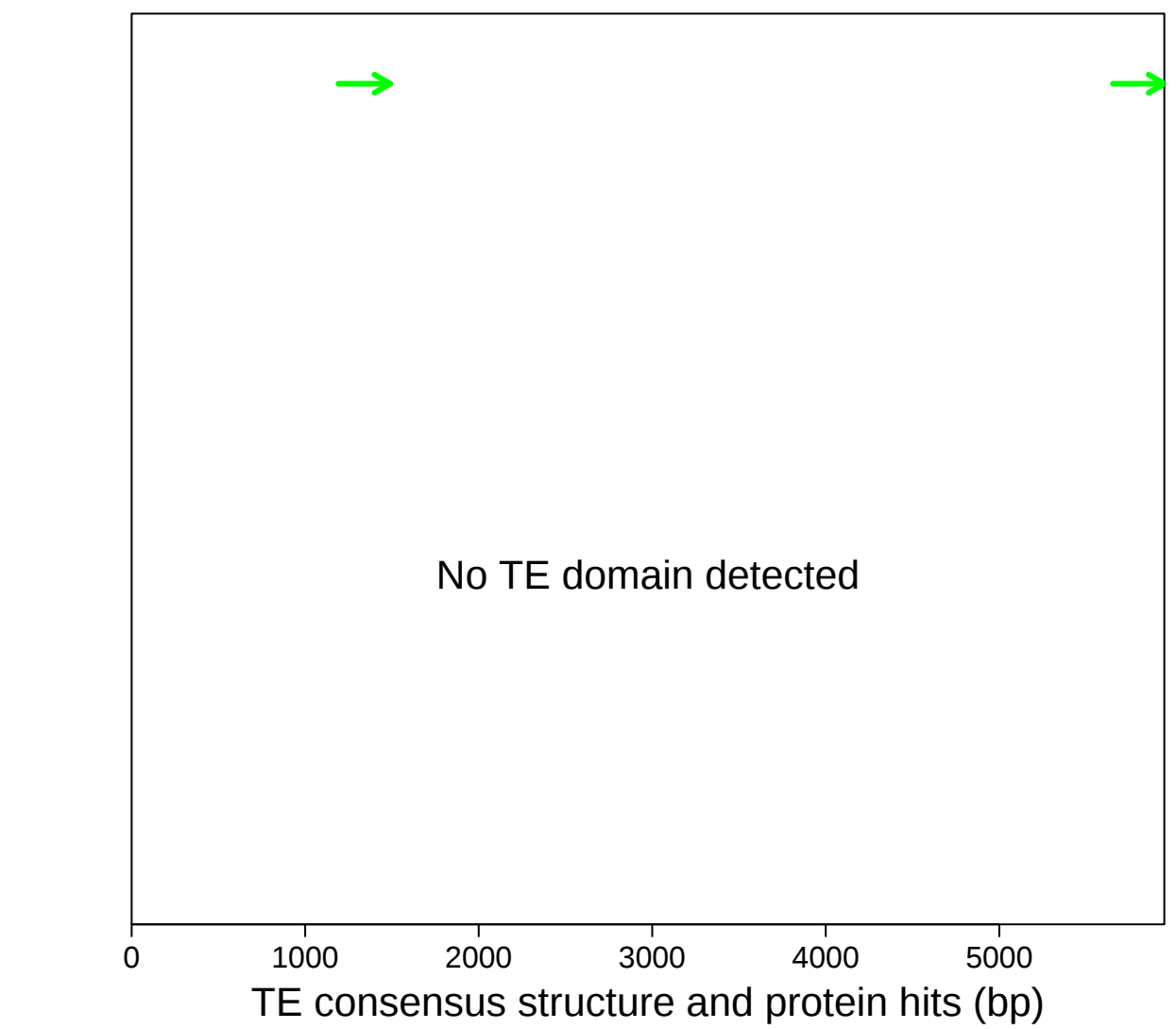
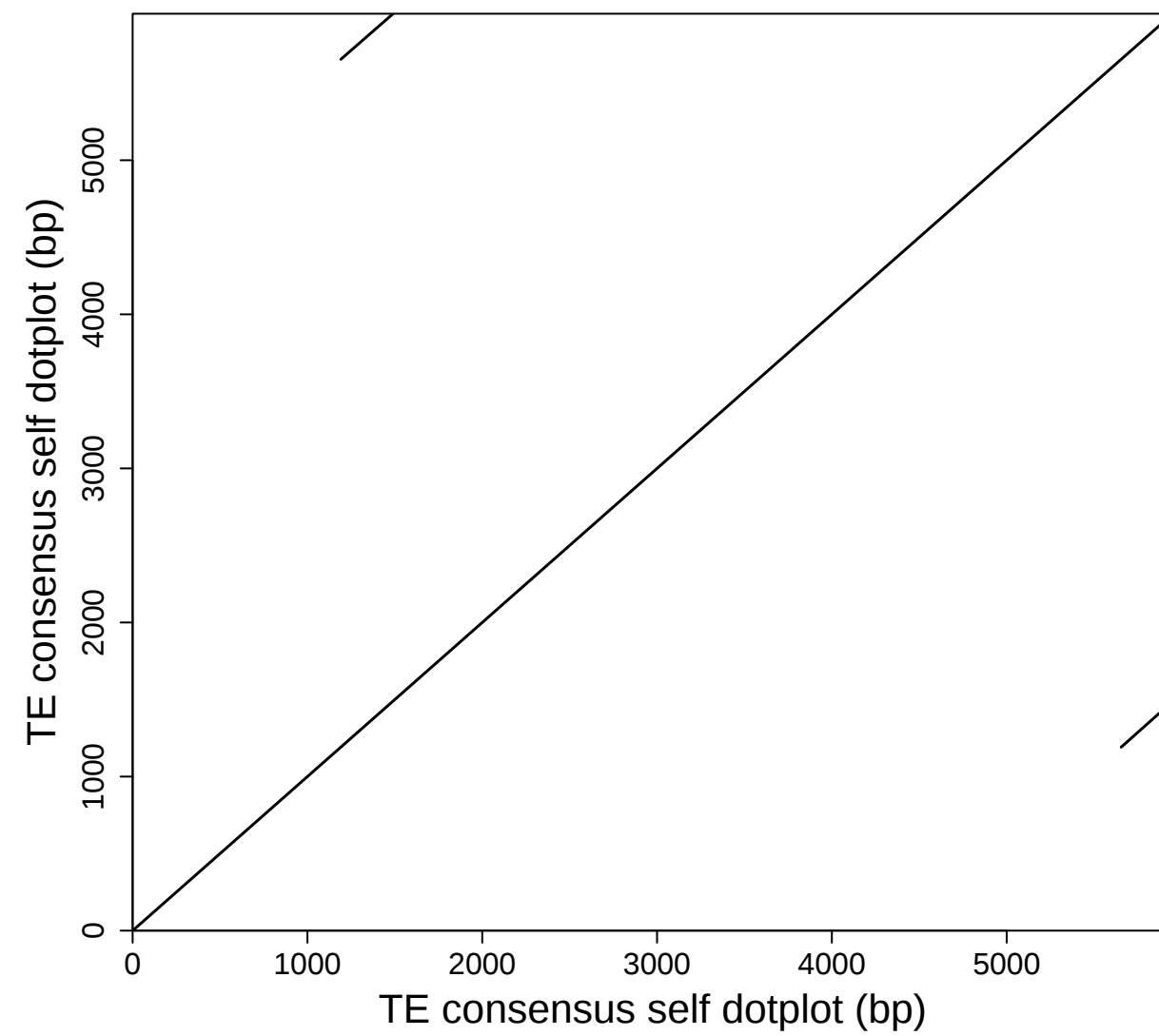
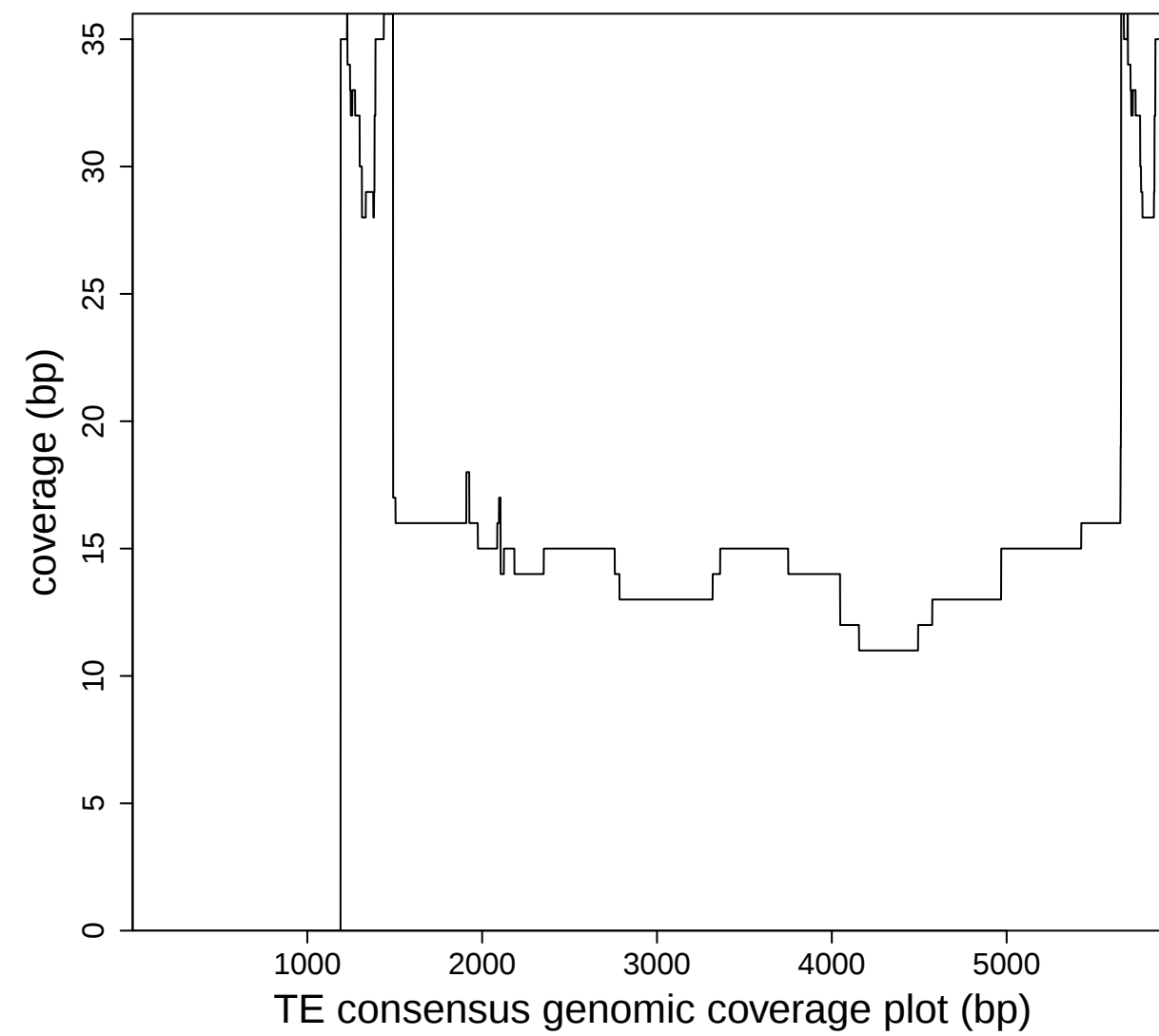
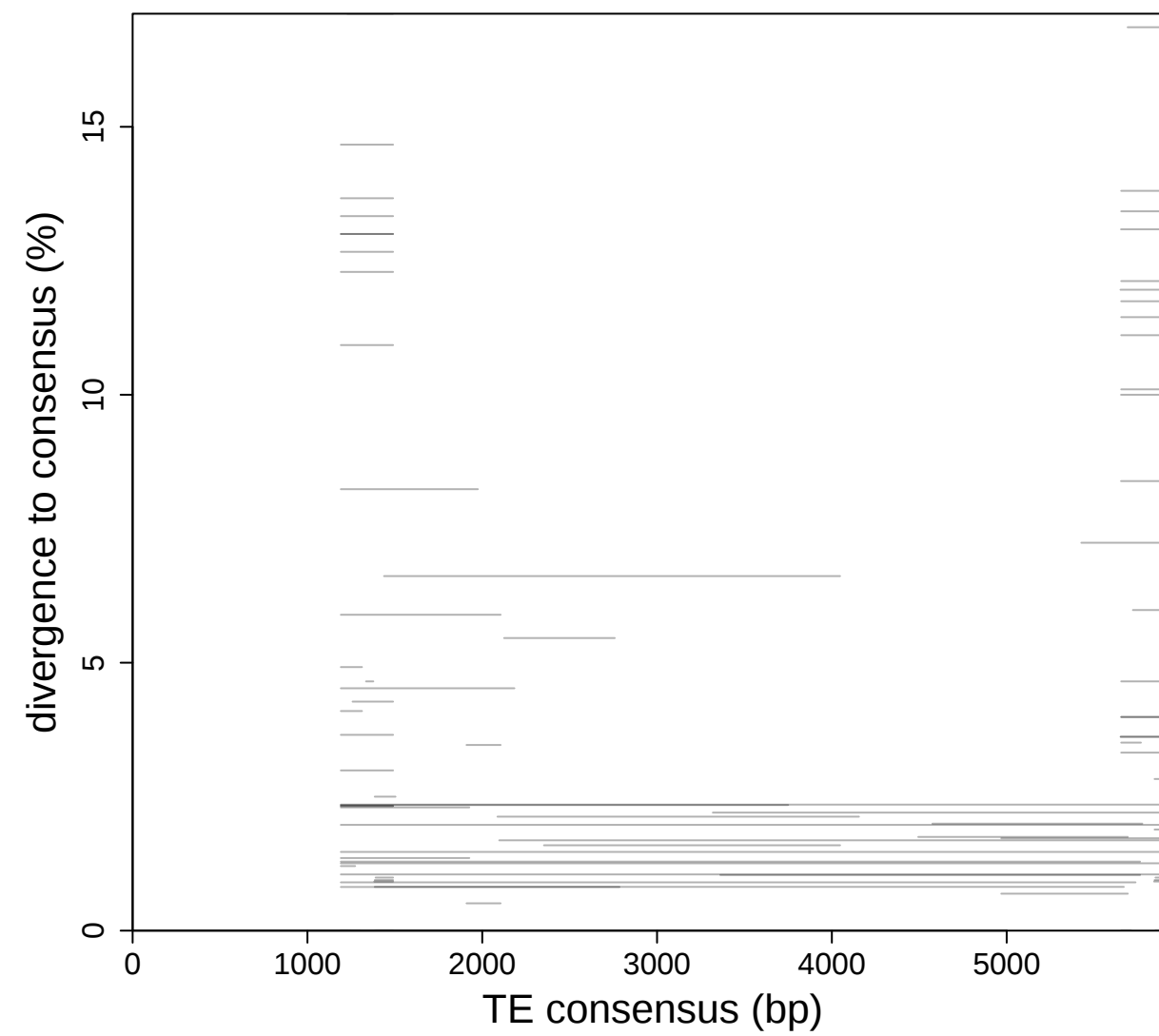
MSA length = 5951





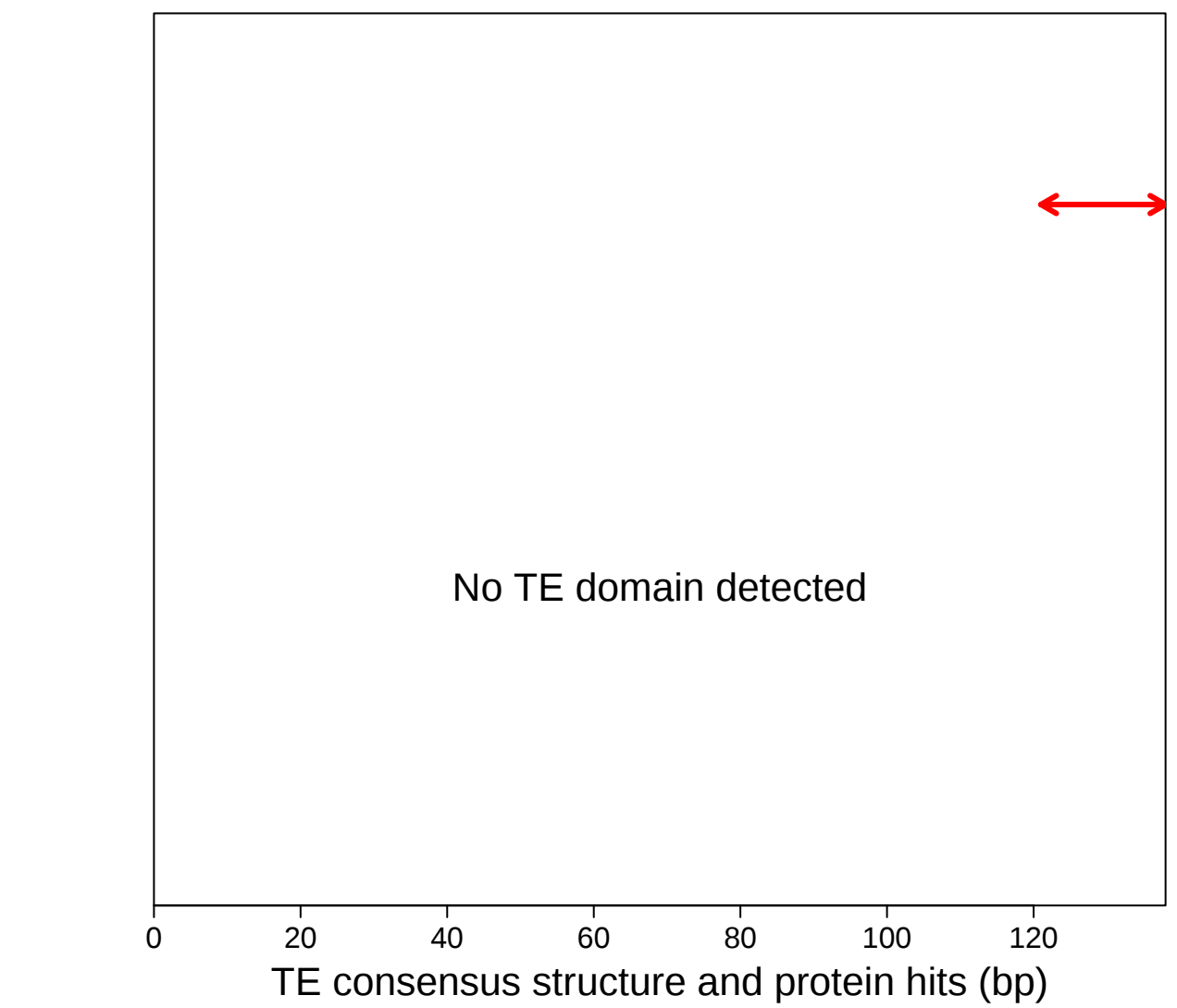
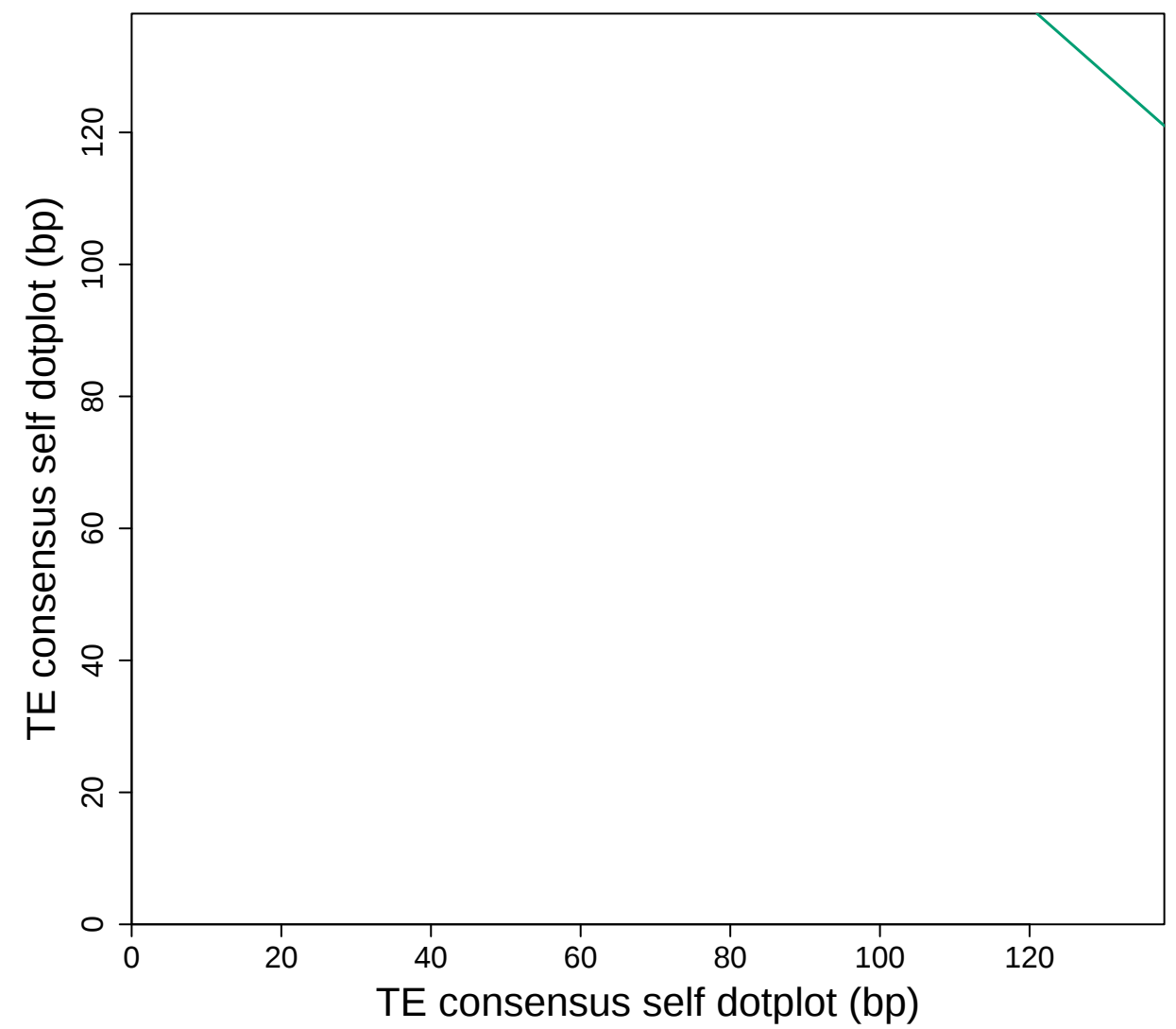
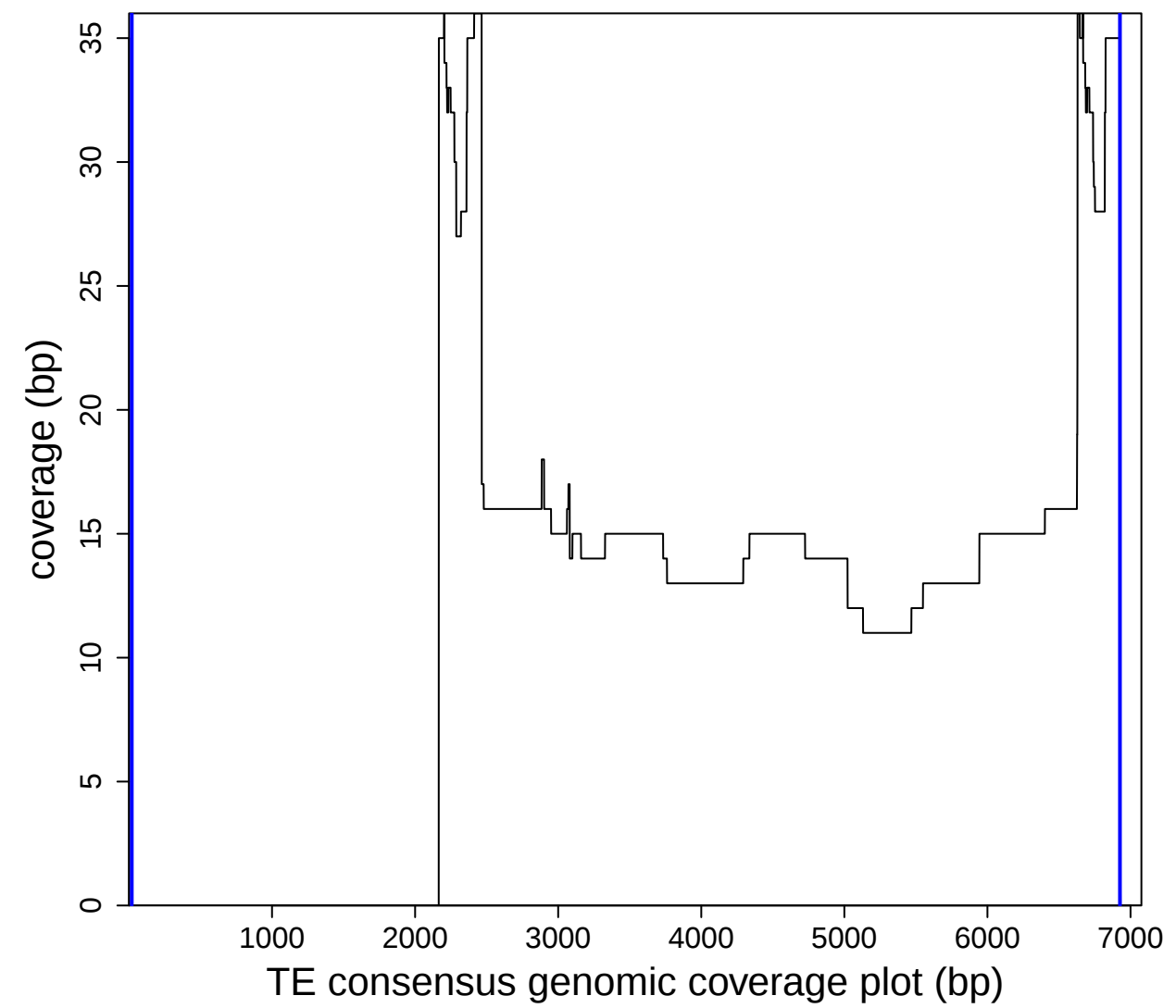
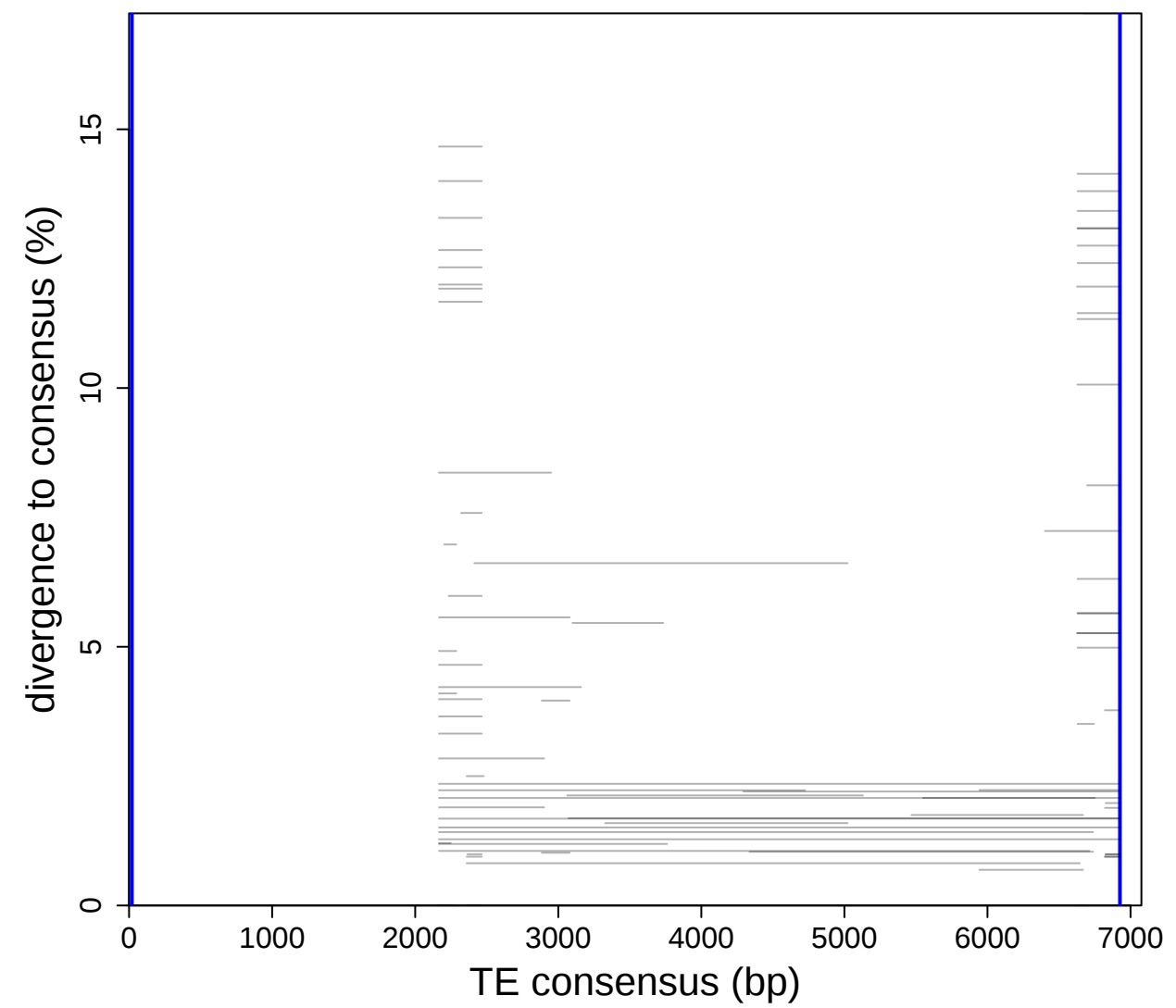
_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcIn.fa_al
size: 5951bp; fragments: 88; full length: 0 (>=5355.9bp)

After TEtrimmer 5951 bp



bed_g_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bclr
size: 7076bp; fragments: 88; full length: 0 (>=6368.4bp)

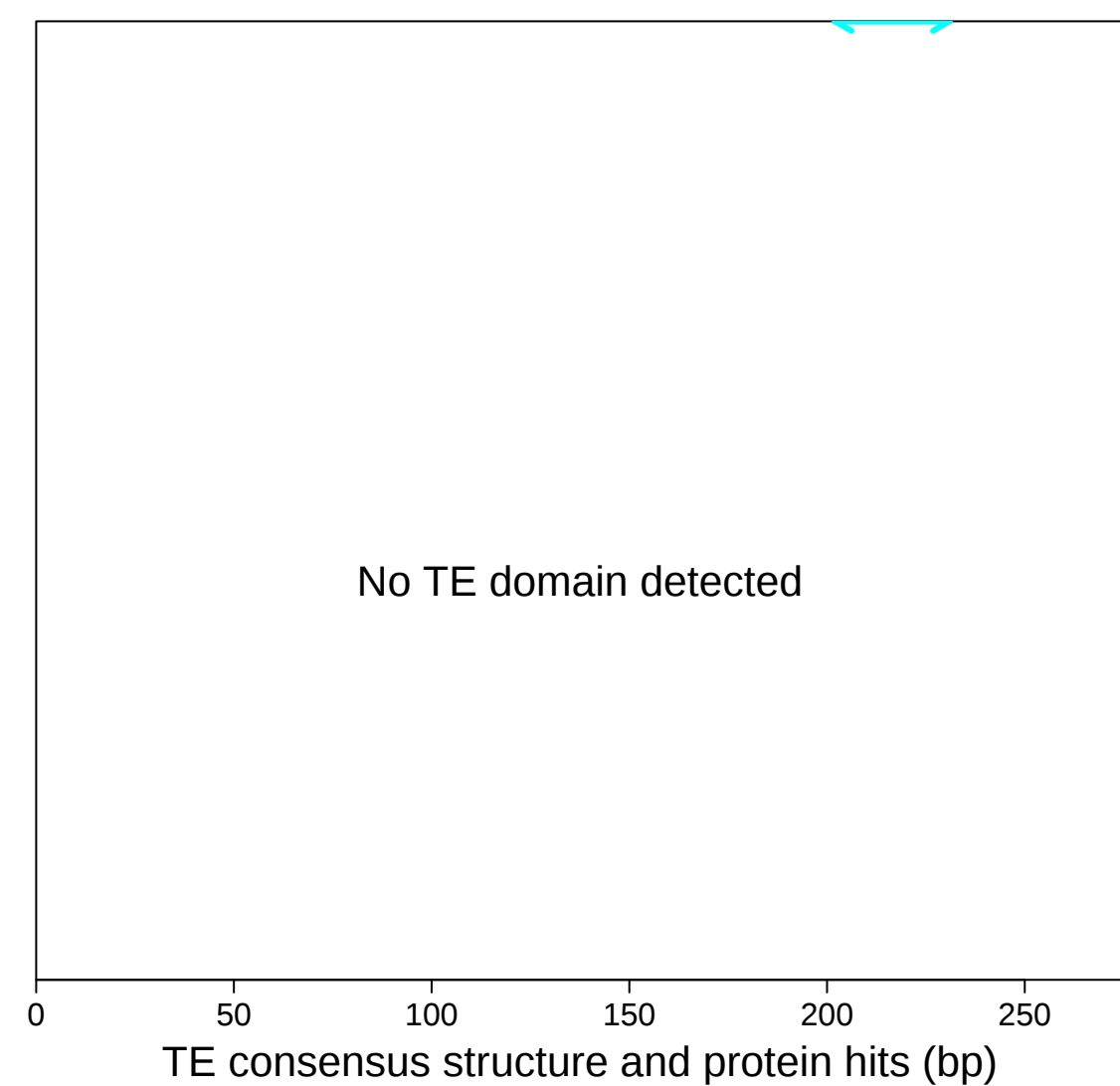
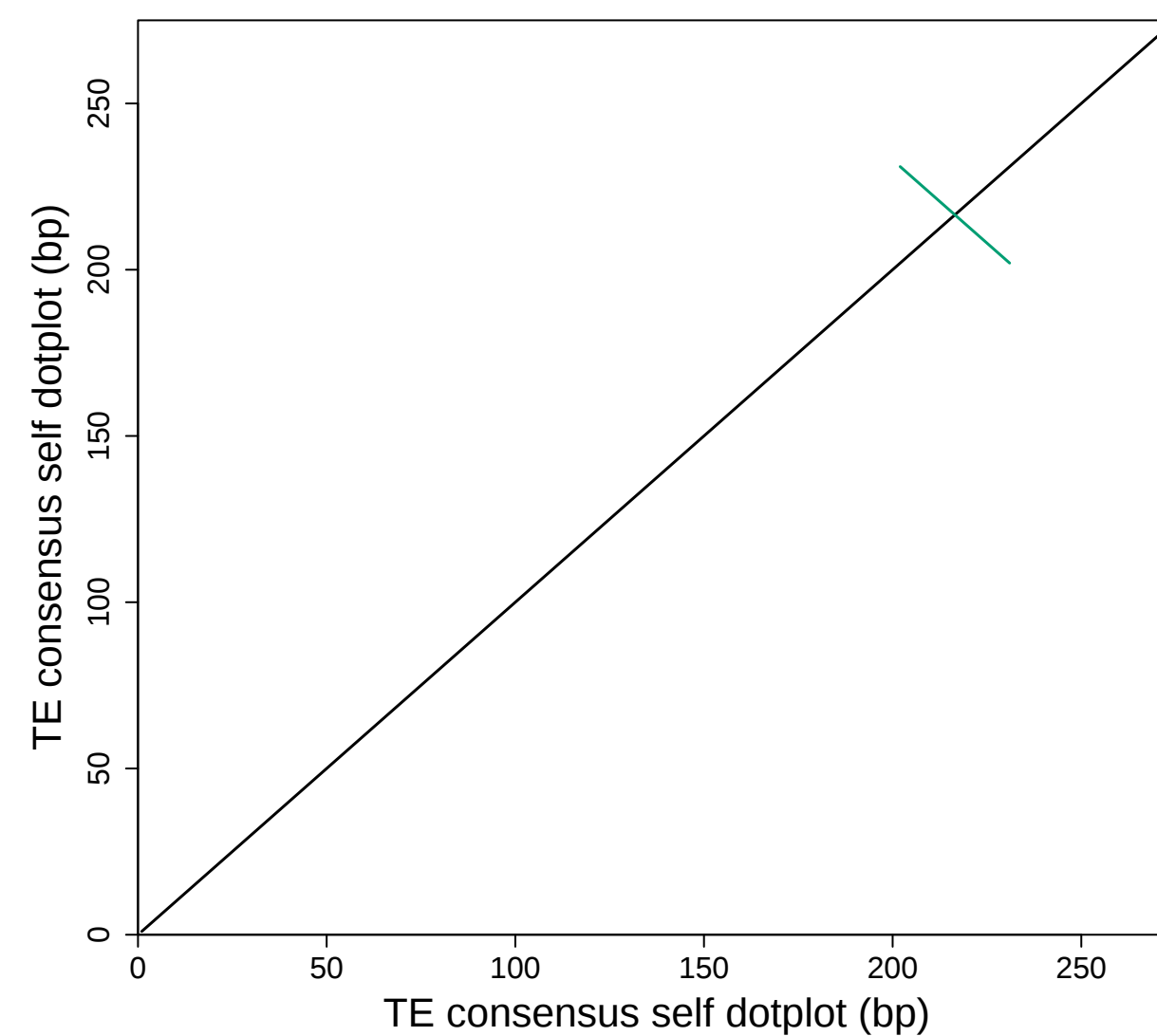
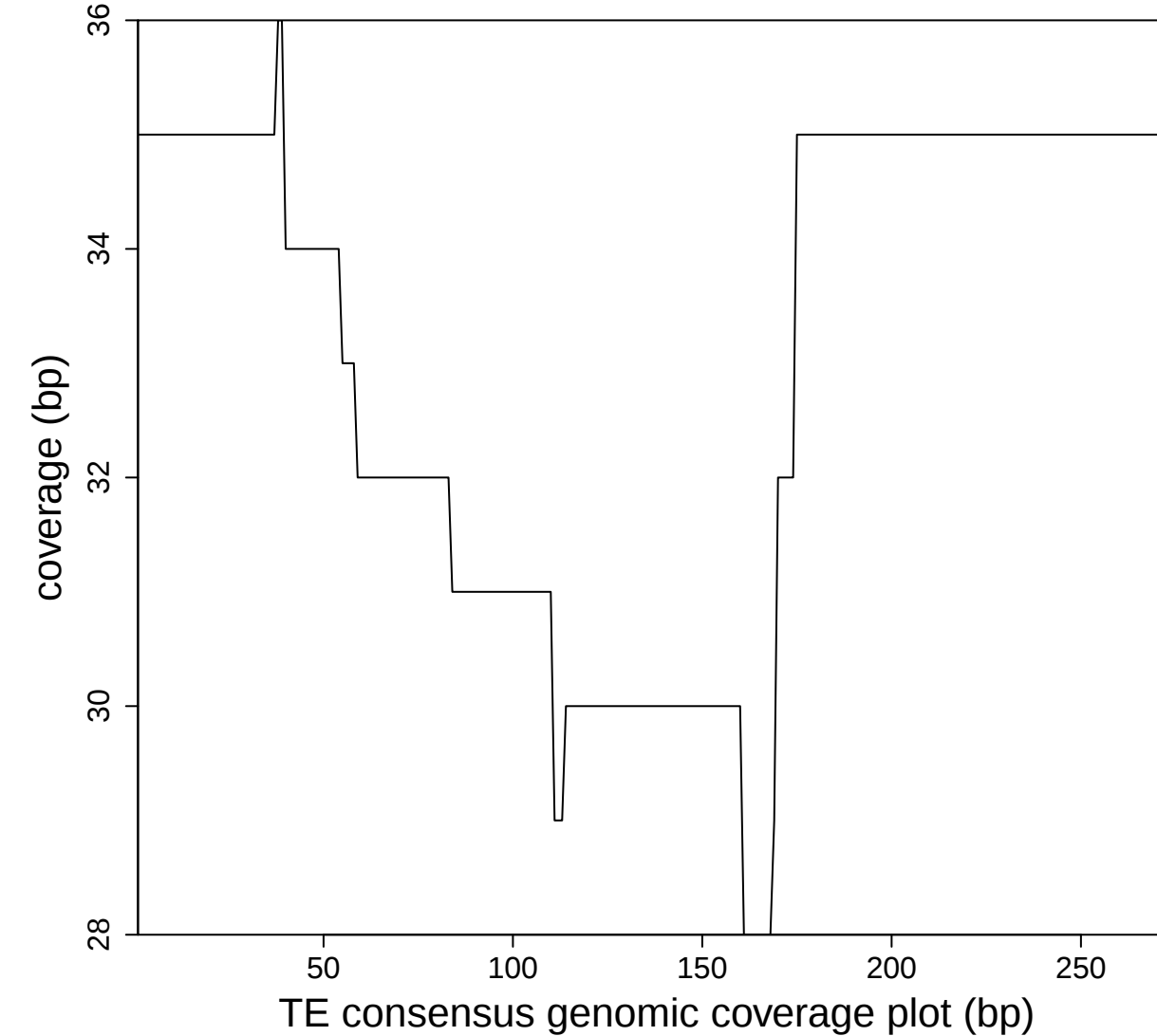
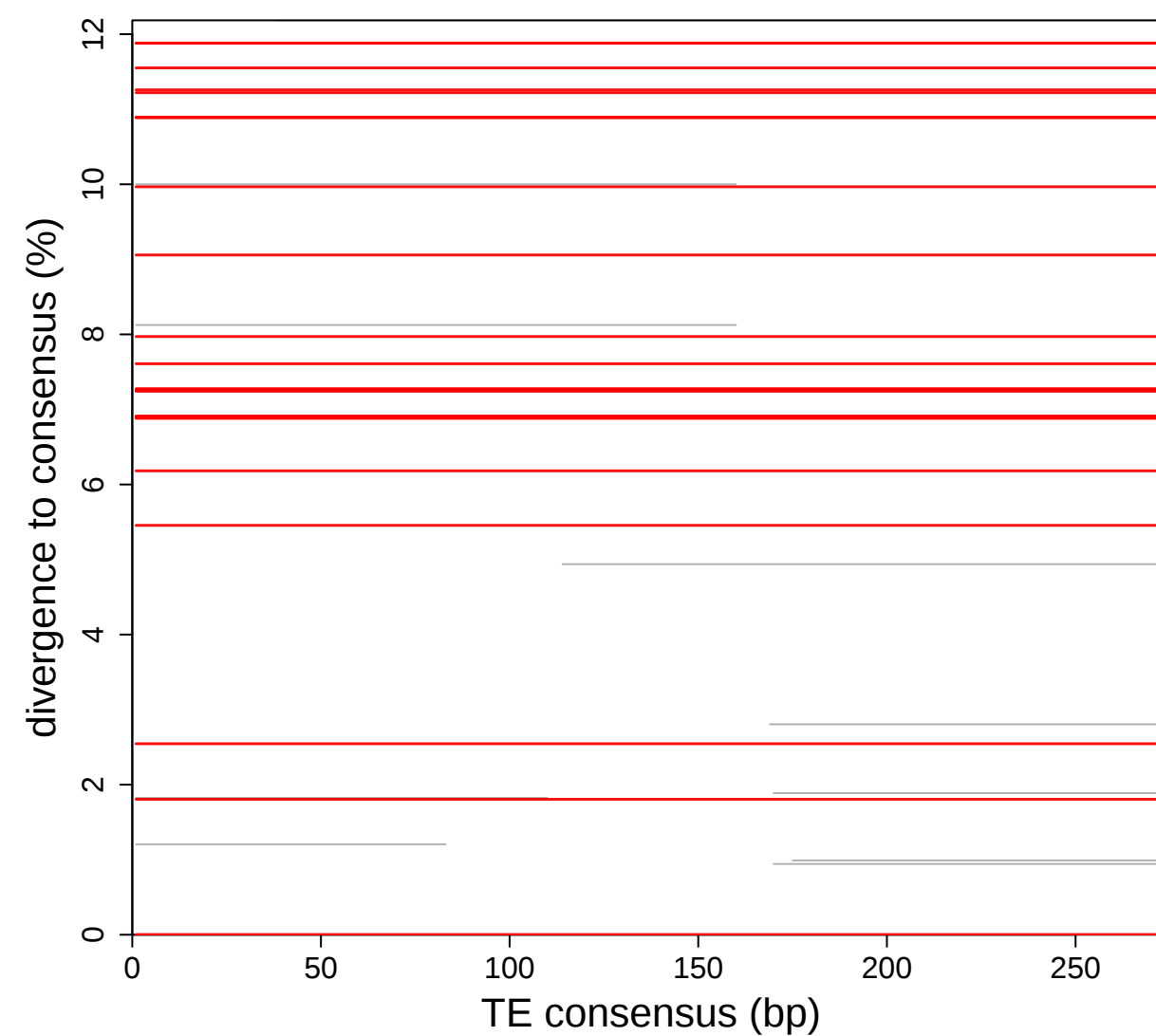
After TEtrimmer Extended plot Blue lines are boundaries



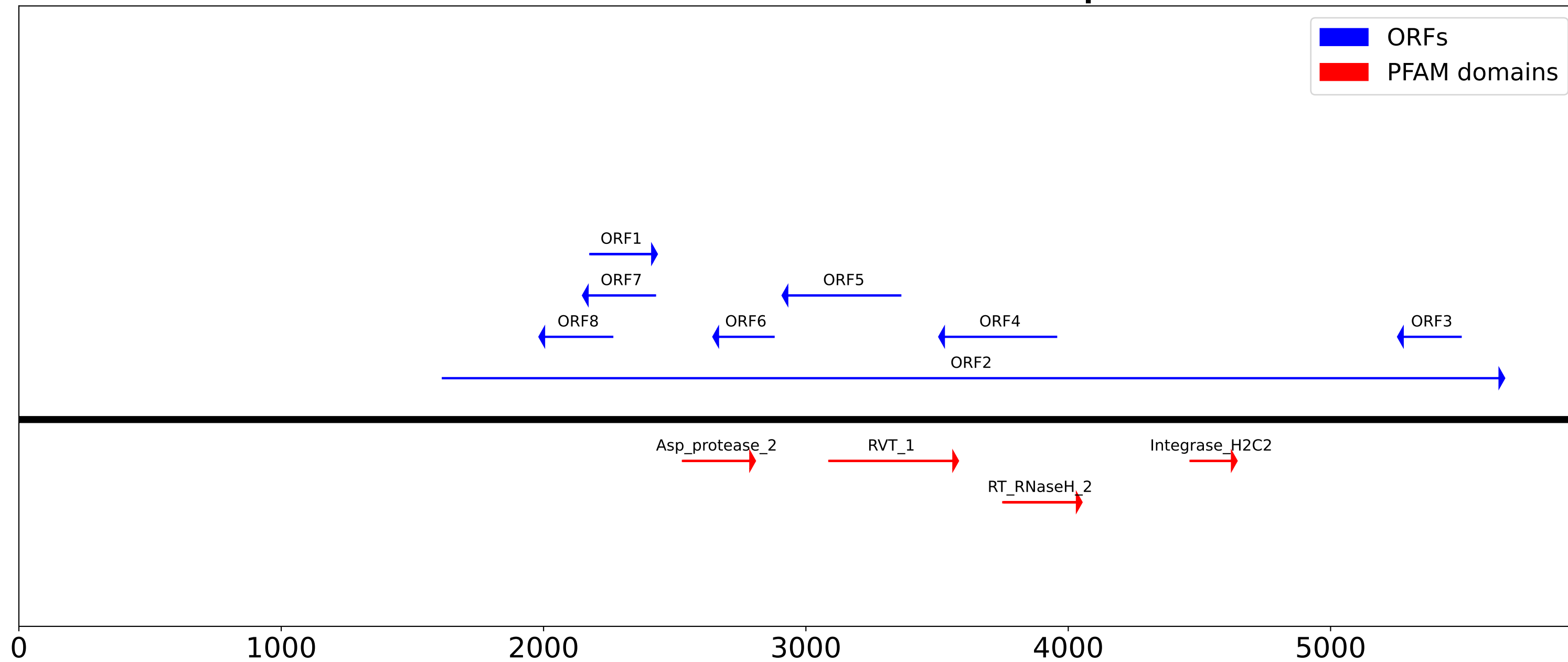
Before TEtrimmer 275 bp

TE: ltr_1_family_278

size: 275bp; fragments: 44; full length: 26 (≥ 247.5 bp)



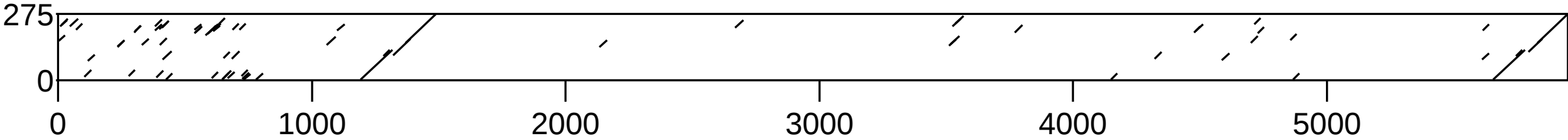
After TEtrimmer ORF and PFAM domain plot



Dotplot

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

nsus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)