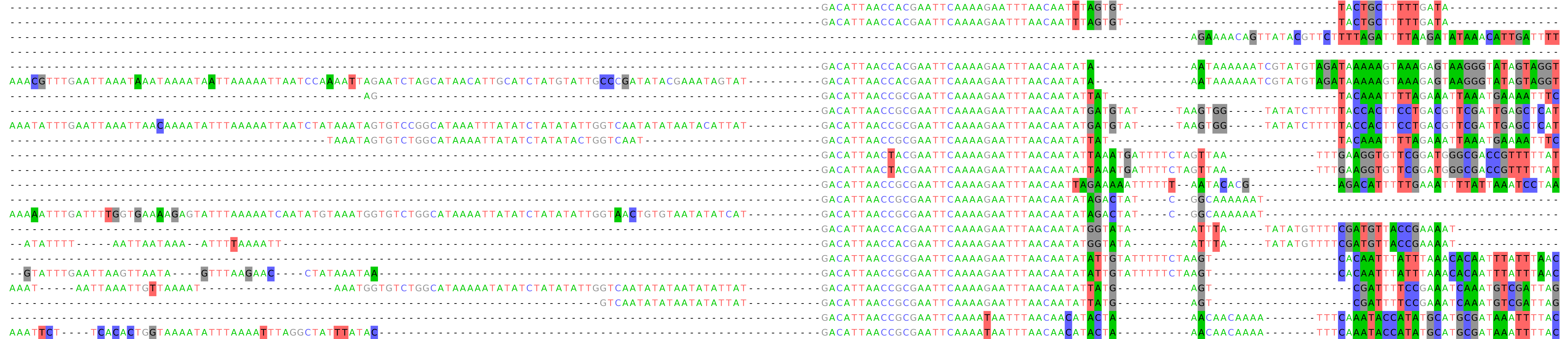
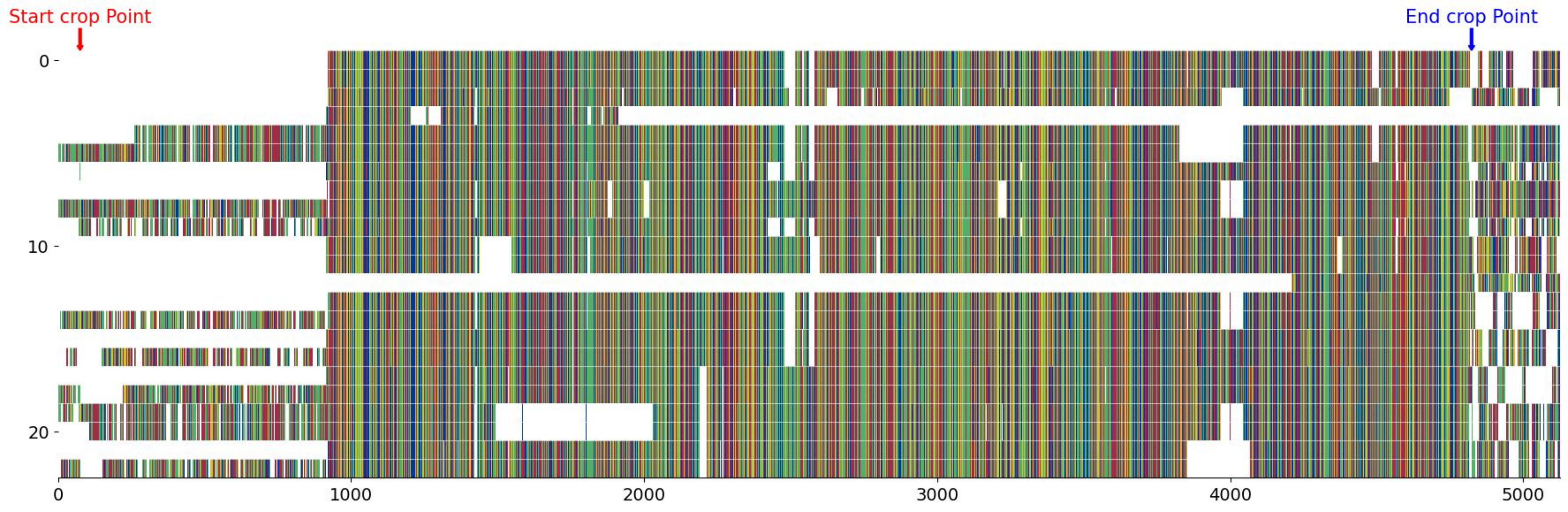


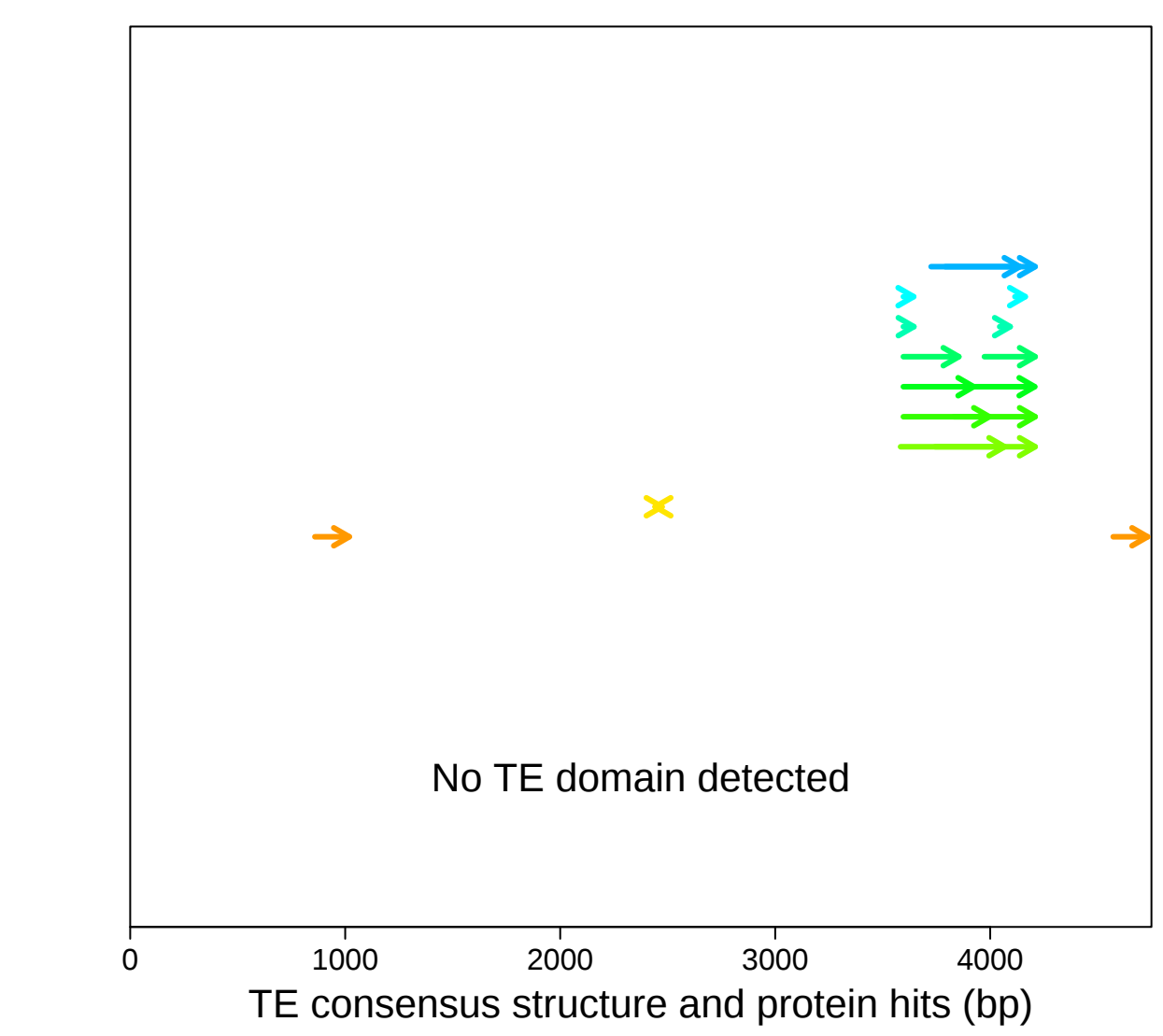
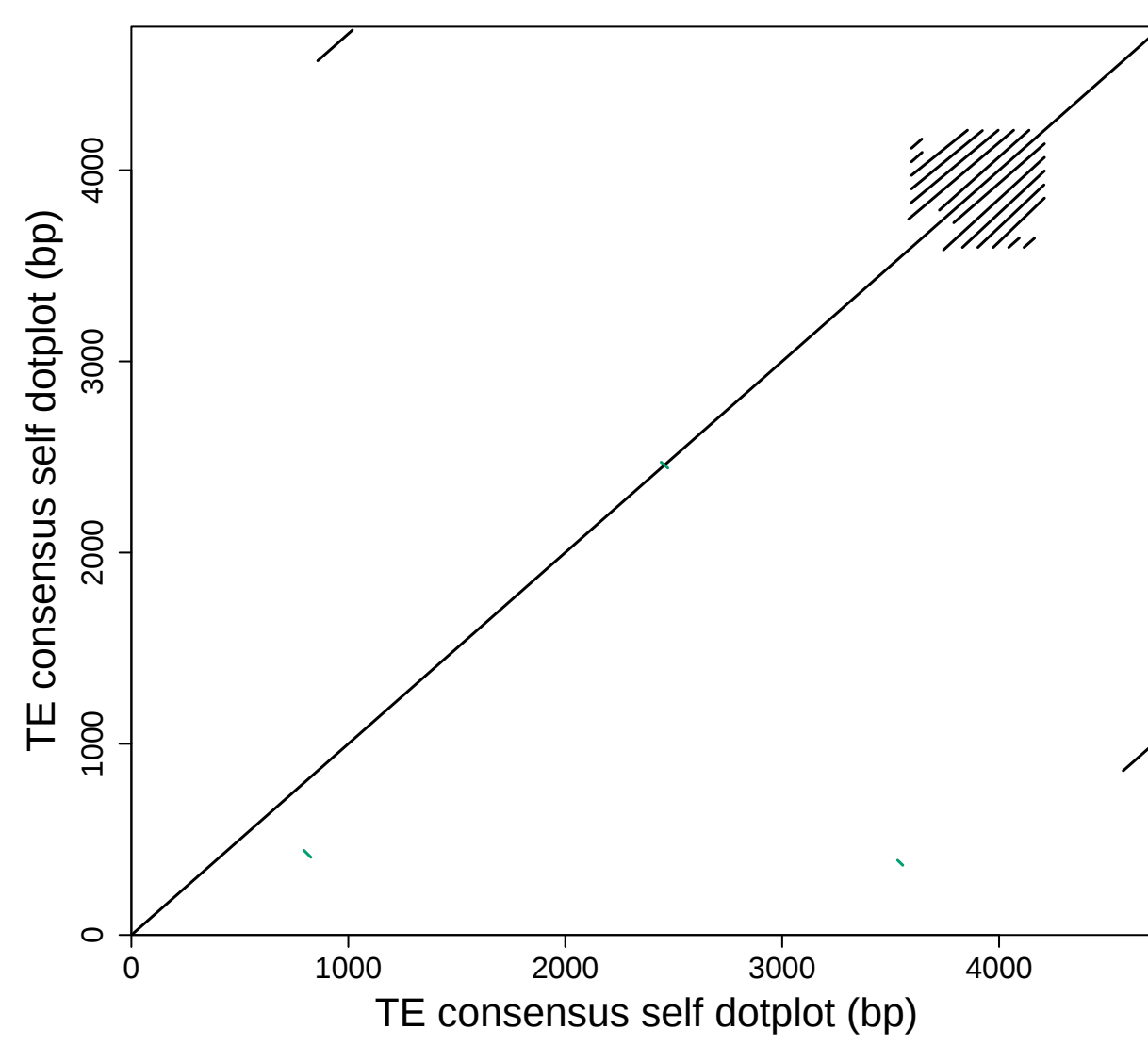
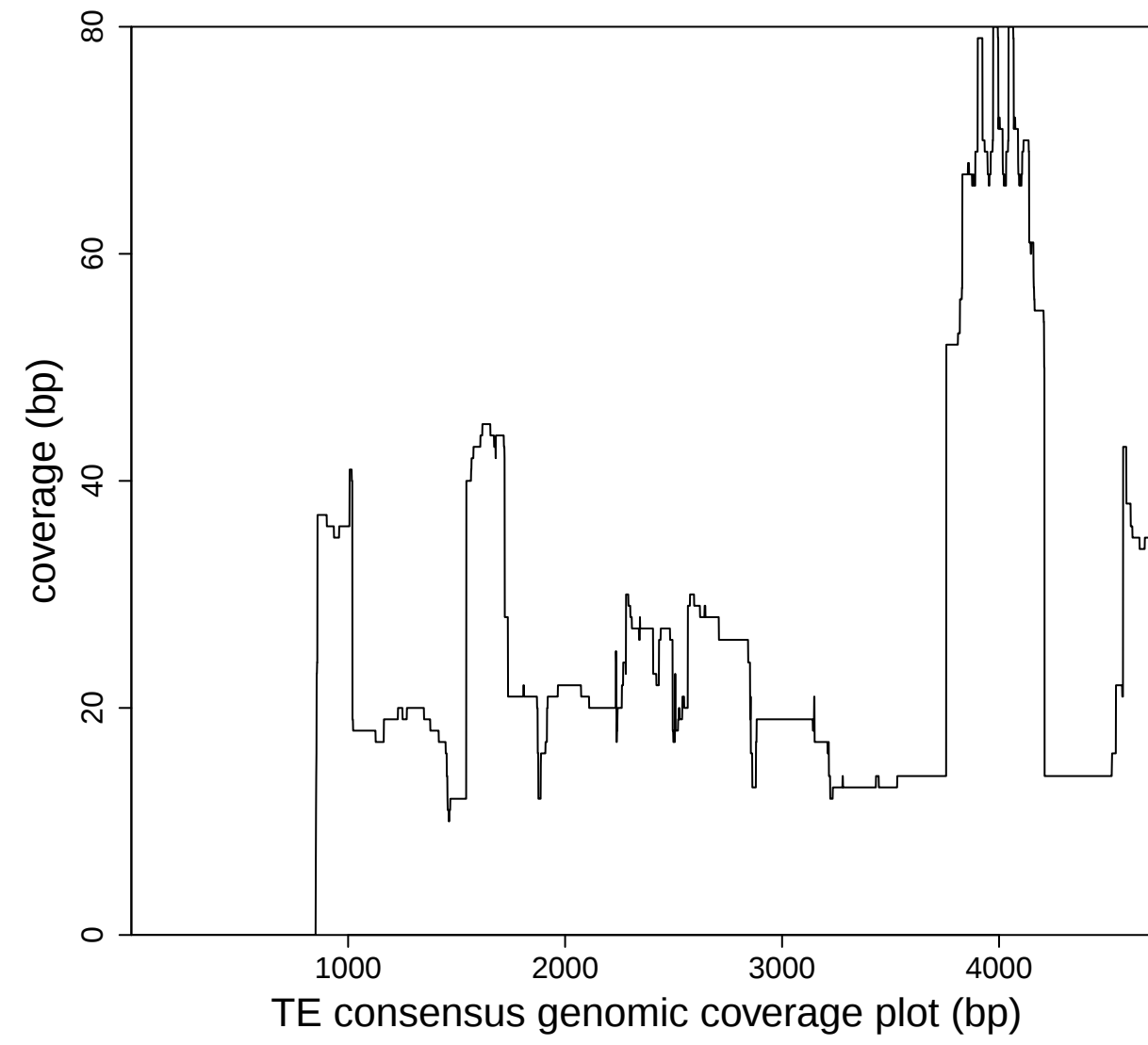
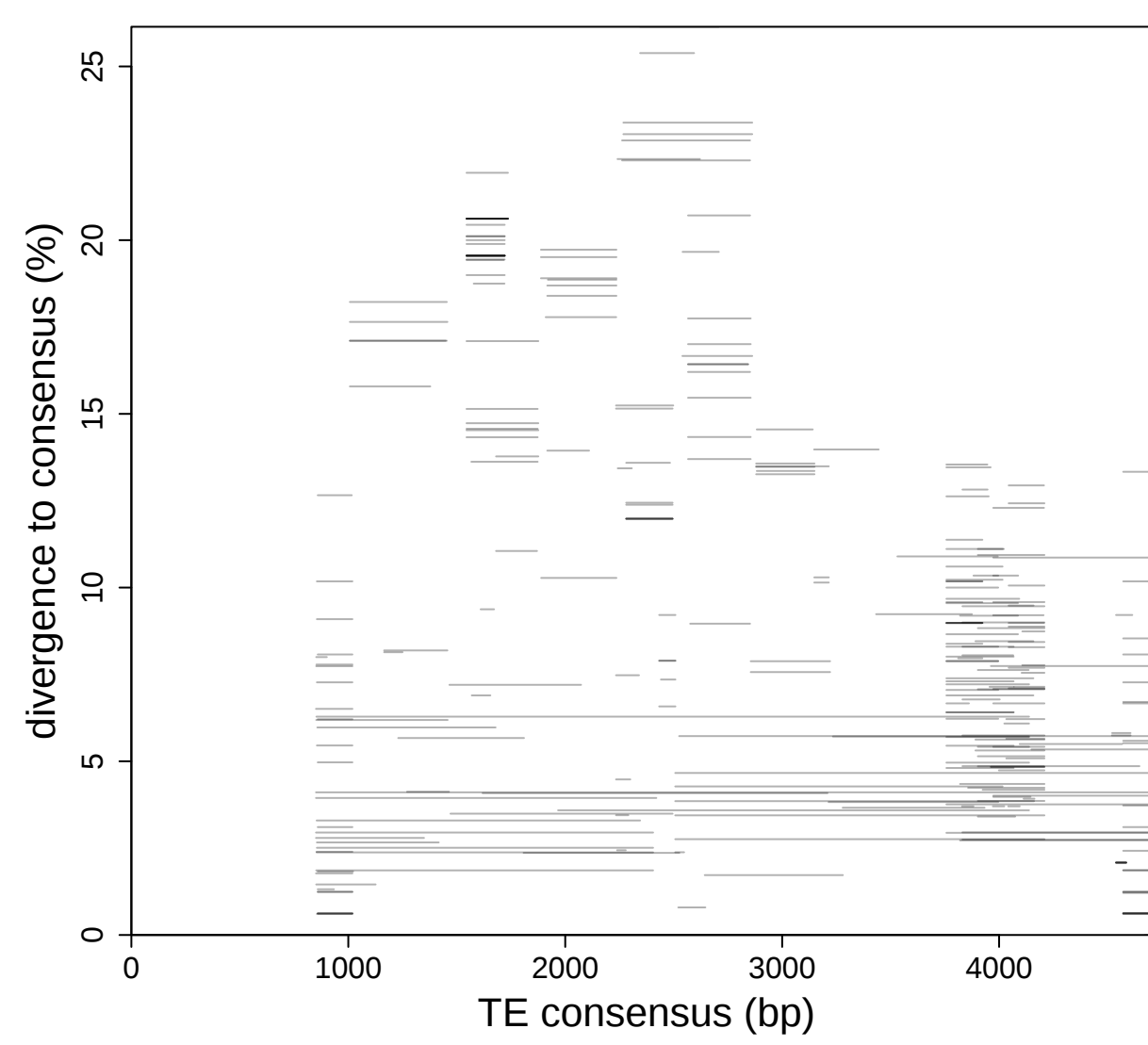
MSA length = 4750





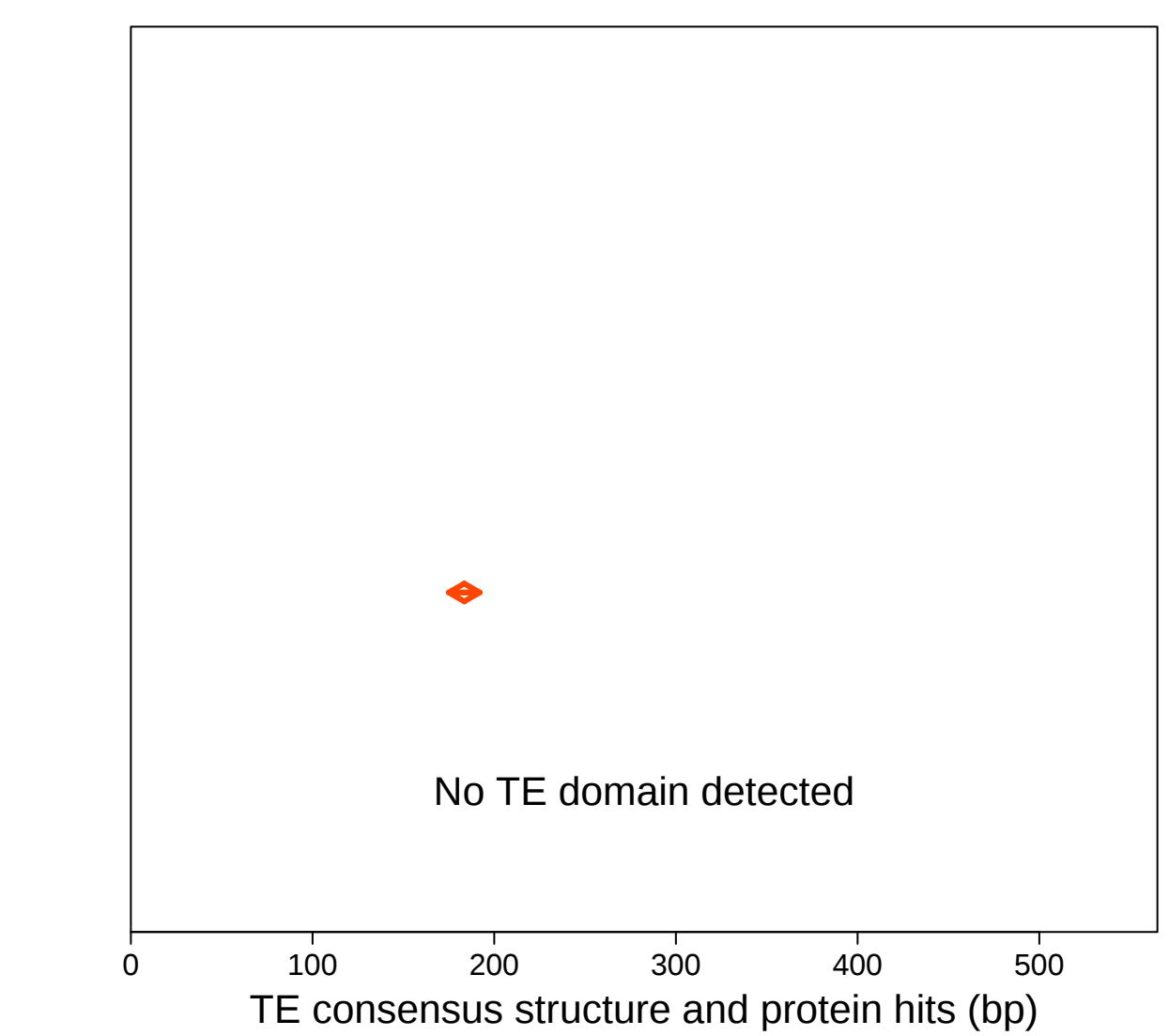
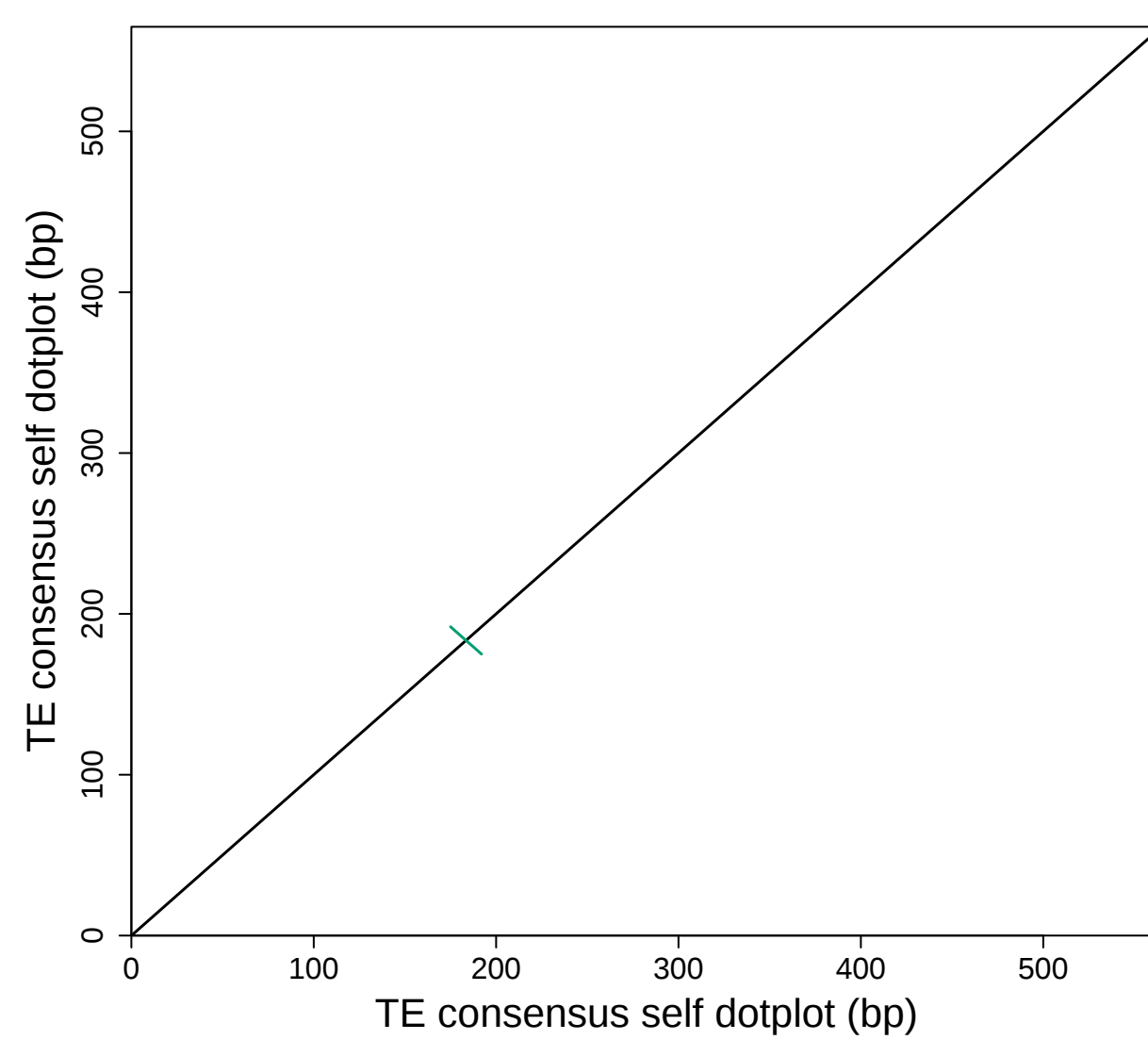
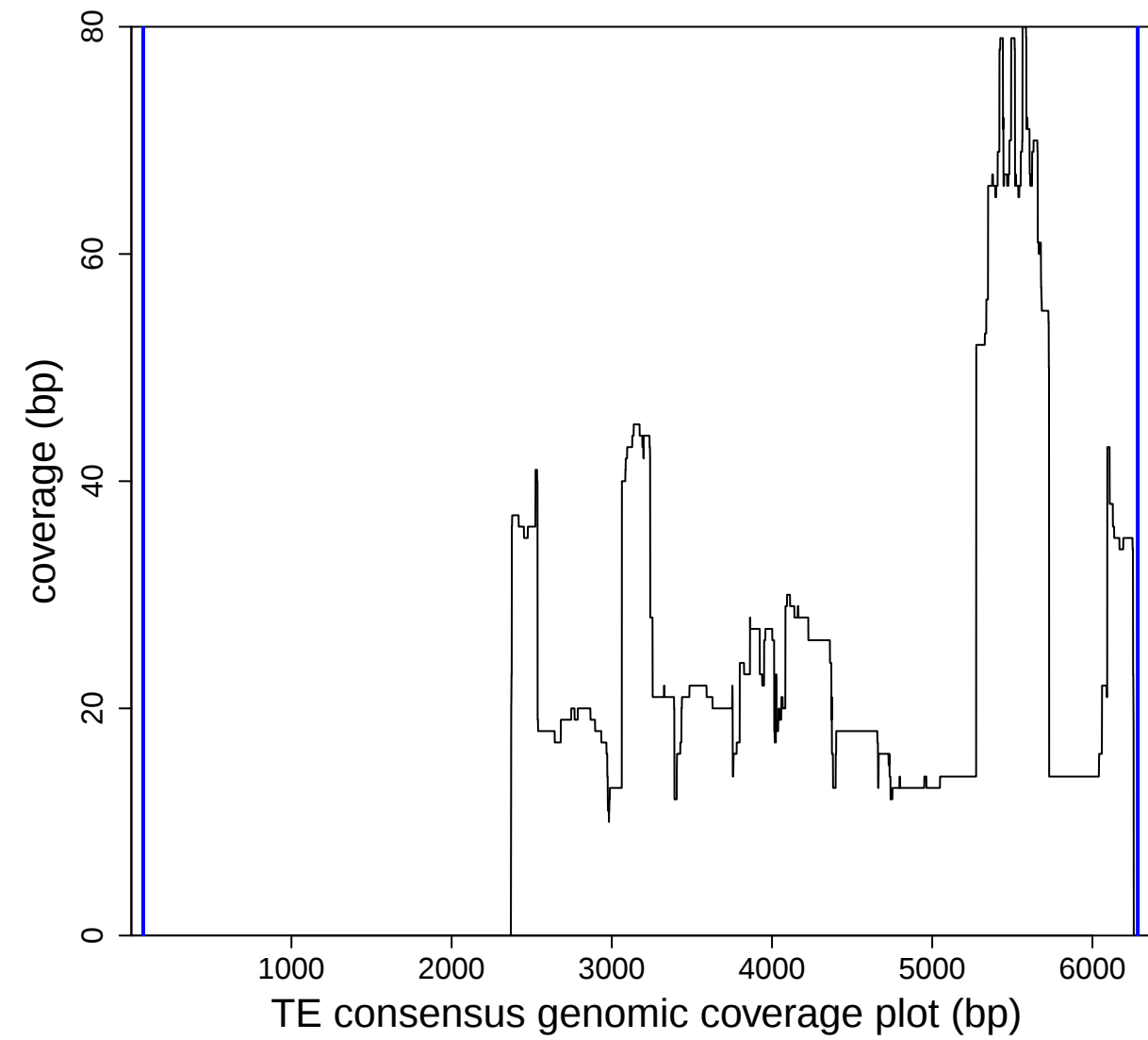
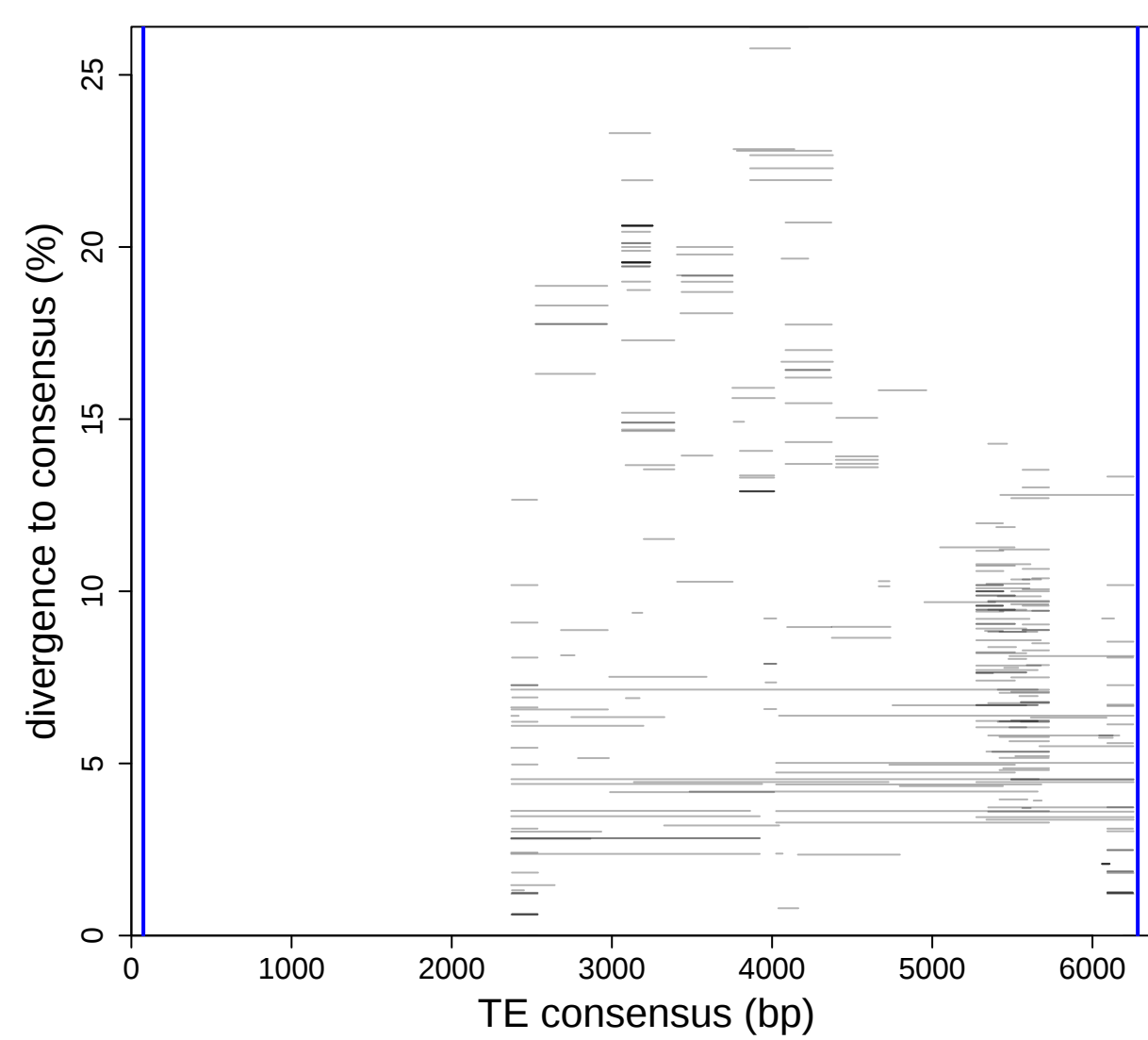
_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcIn.fa_al
size: 4750bp; fragments: 296; full length: 0 (>=4275bp)

After TEtrimmer 4750 bp



bed_g_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bclr
size: 6433bp; fragments: 289; full length: 0 (>=5789.7bp)

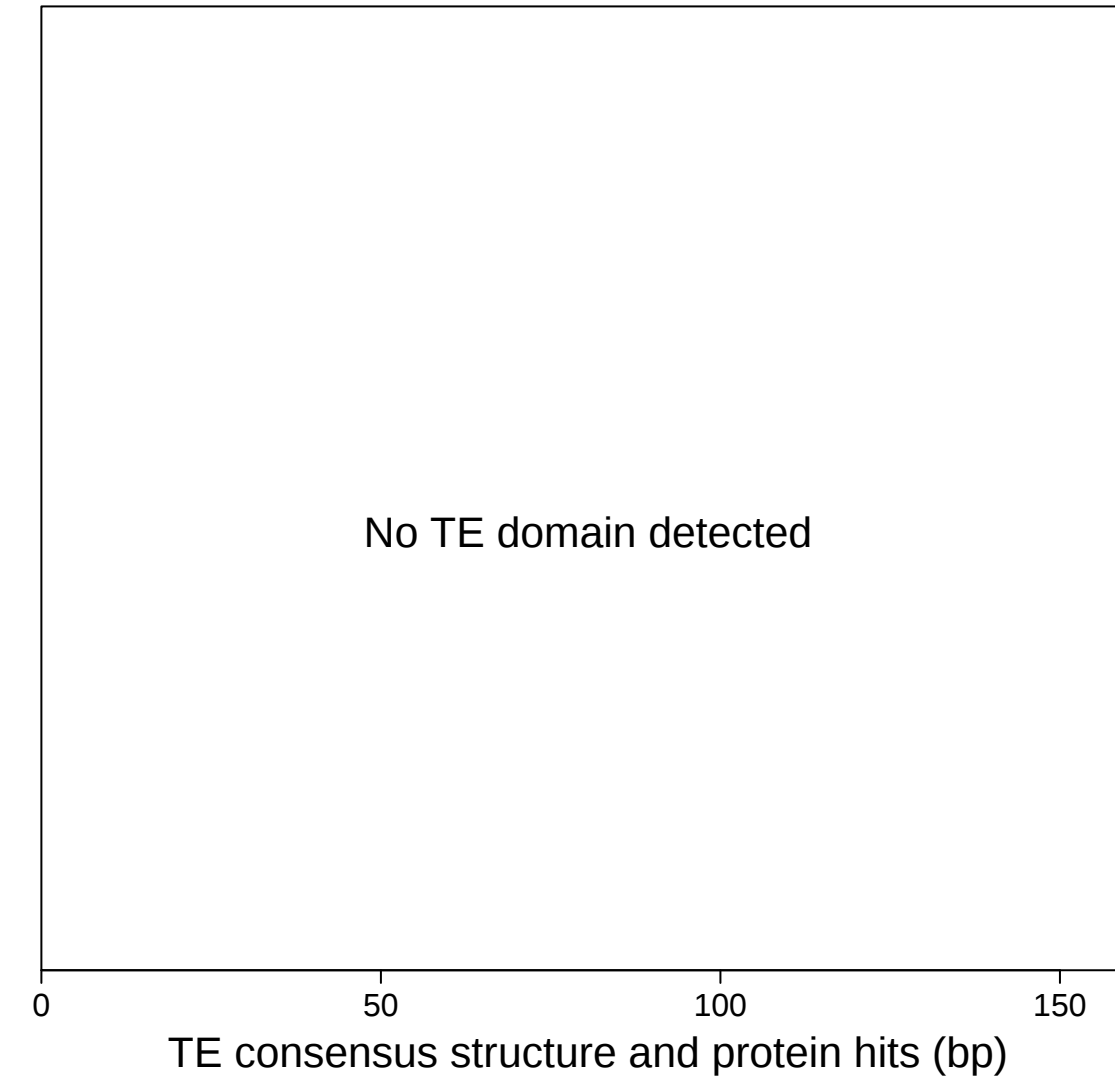
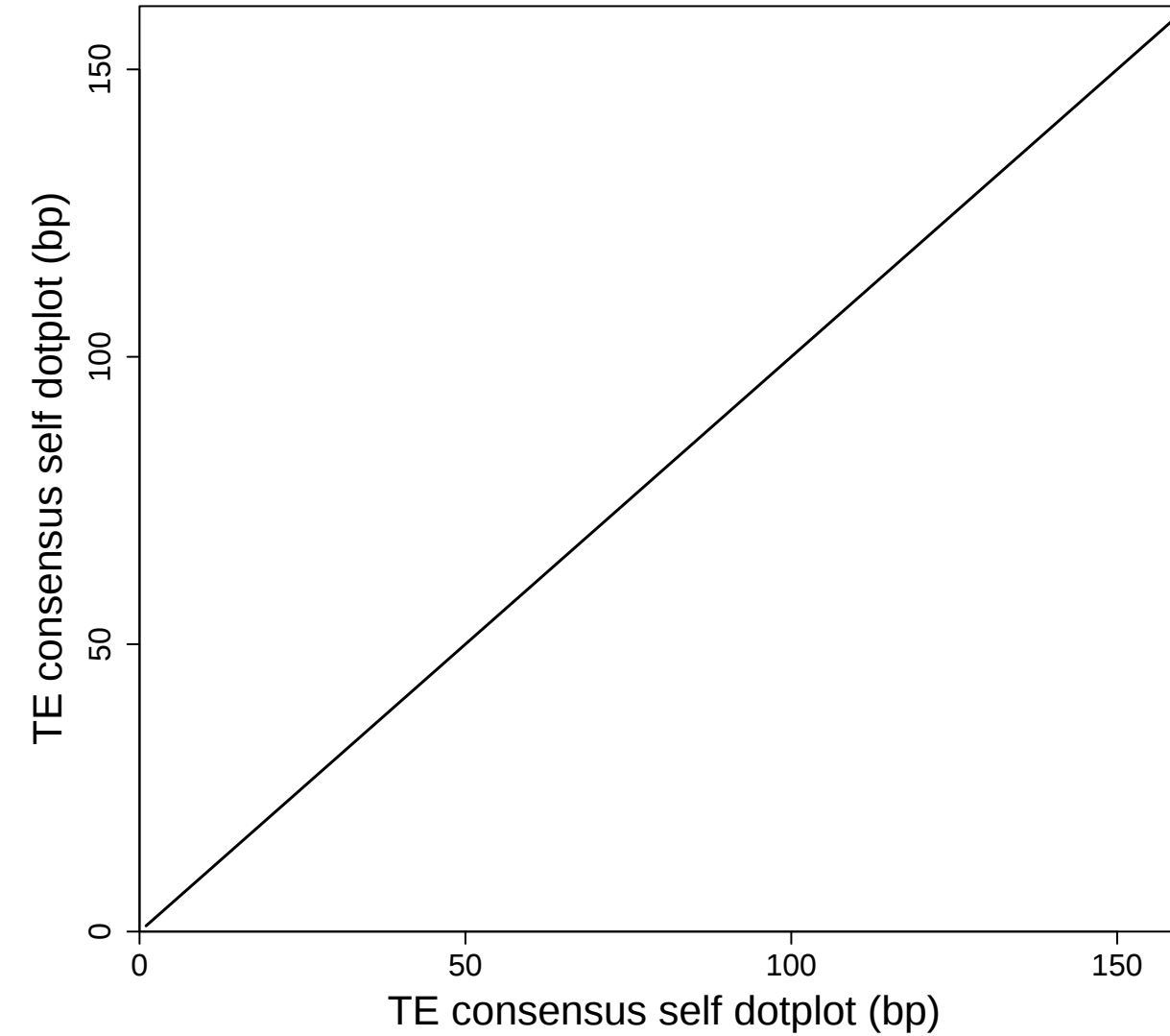
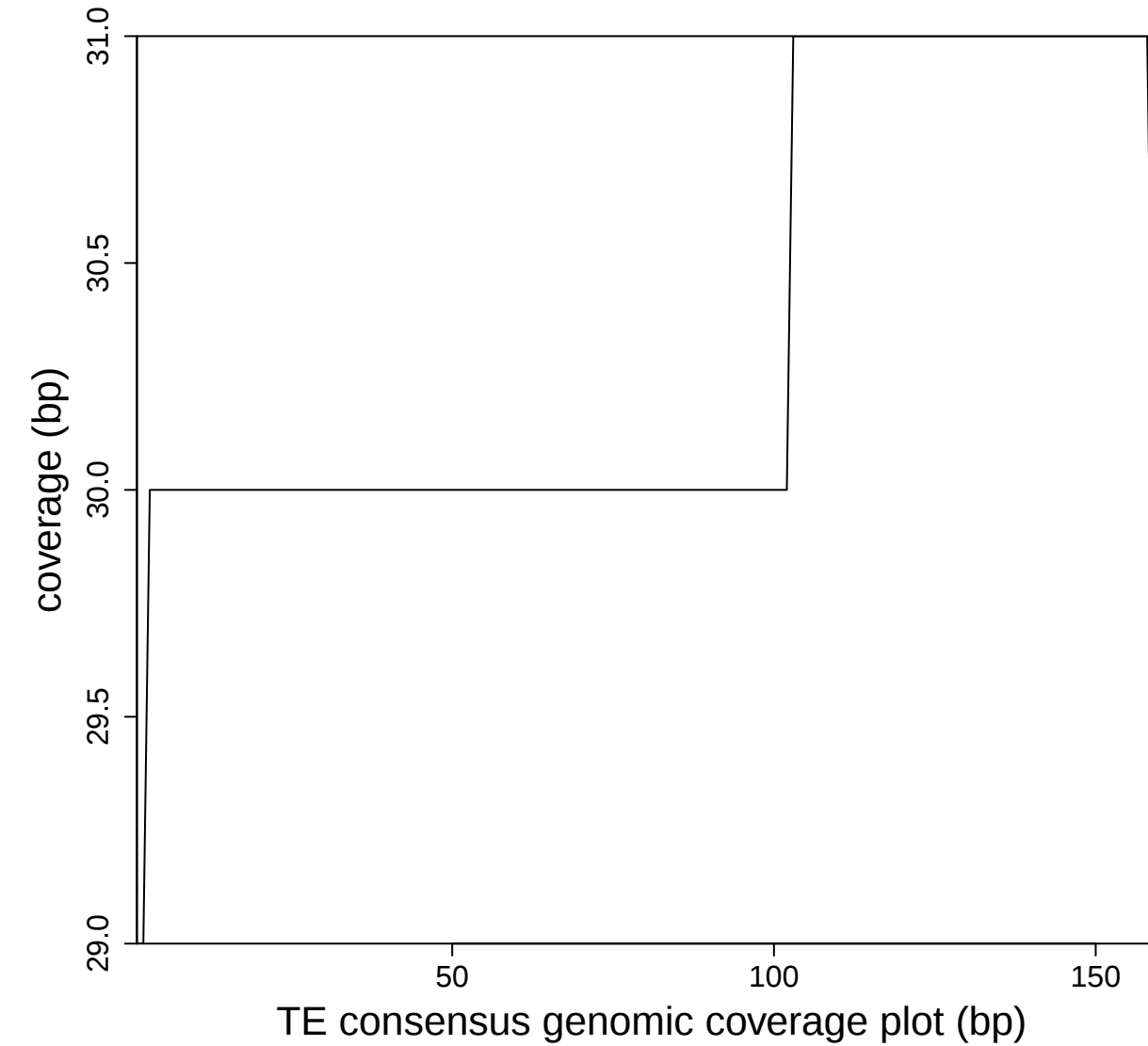
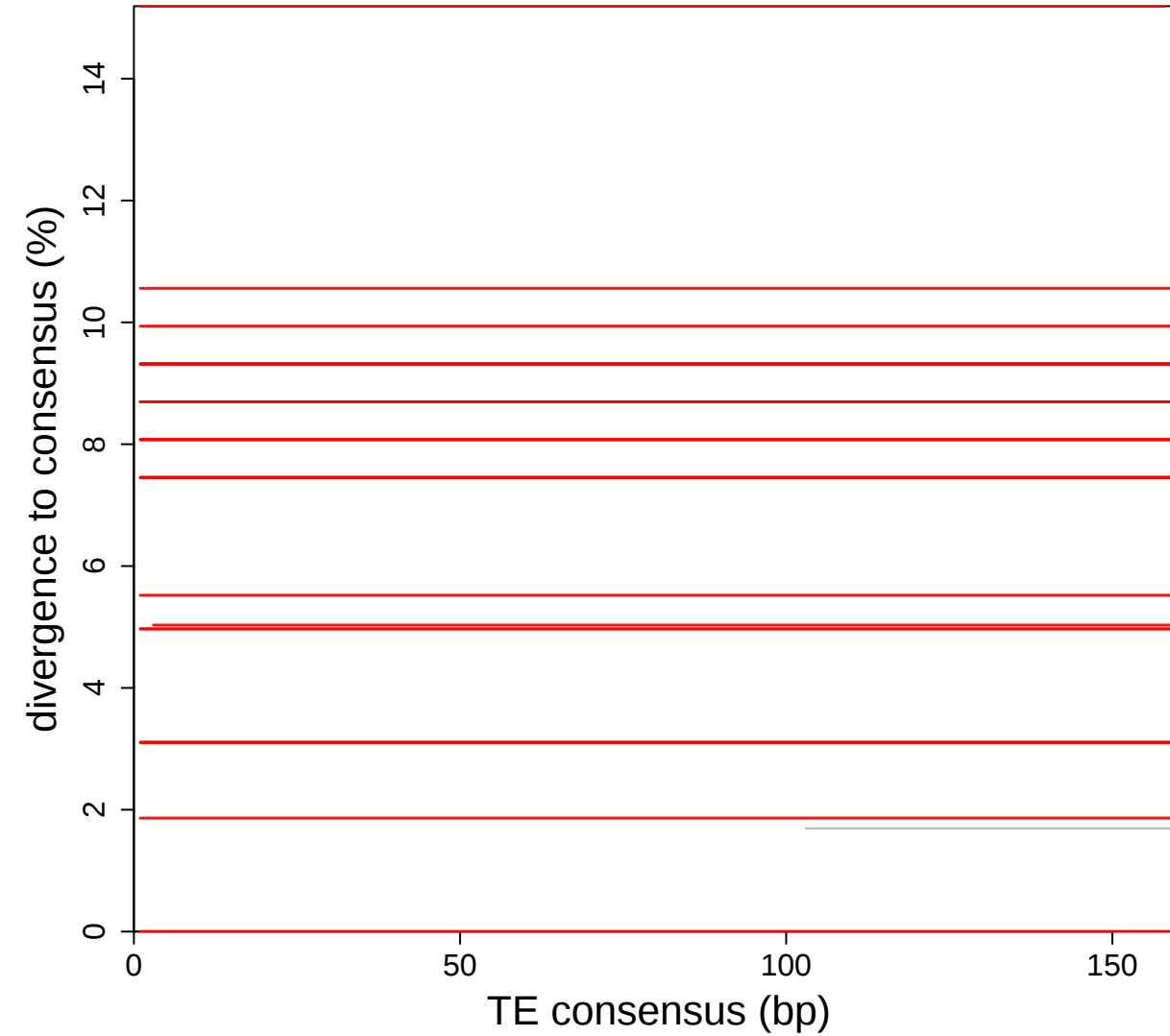
After TETrimmer Extended plot Blue lines are boundaries



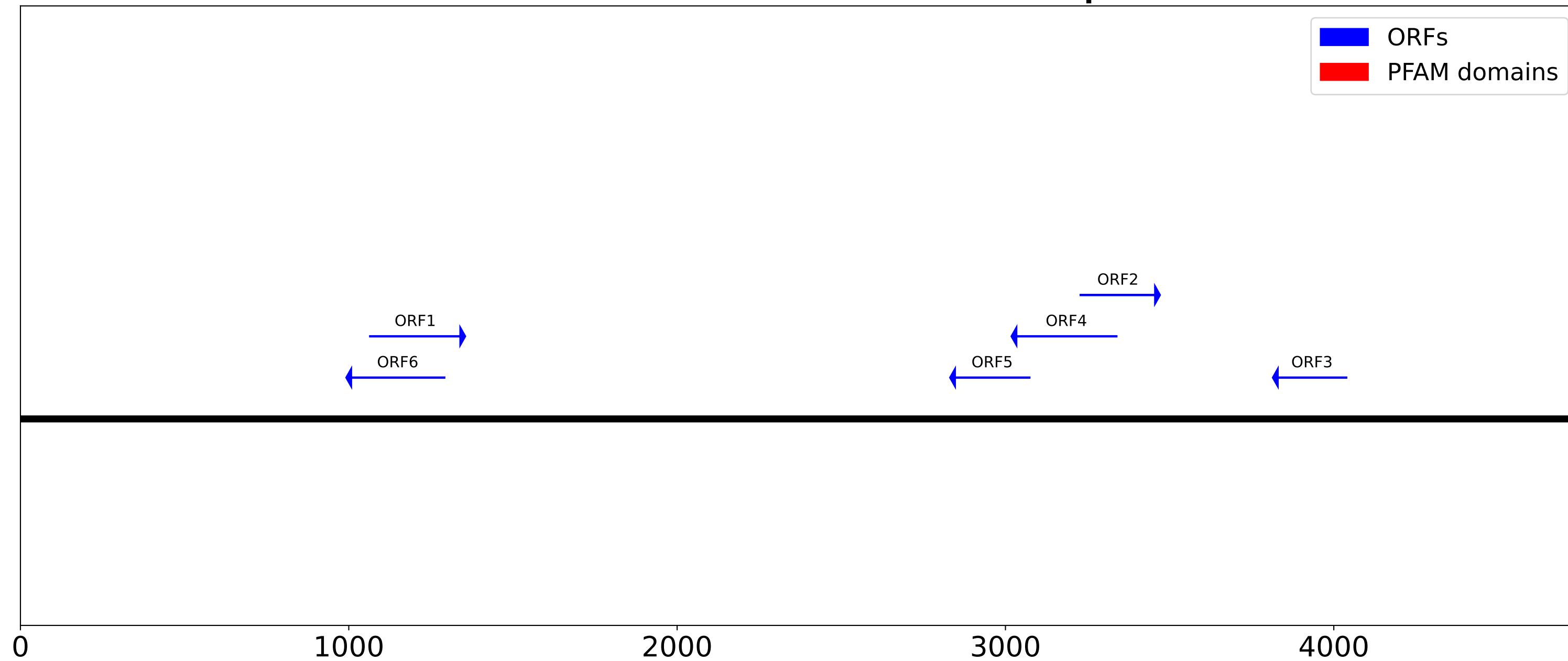
Before TEtrimmer 161 bp

TE: ltr_1_family_128

size: 161bp; fragments: 31; full length: 30 (≥ 144.9 bp)



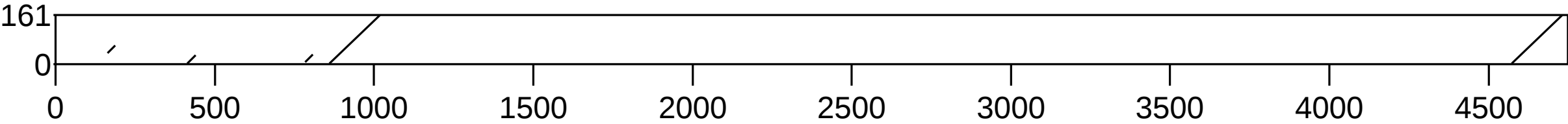
After TEtrimmer ORF and PFAM domain plot



Dotplot

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

nsus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)