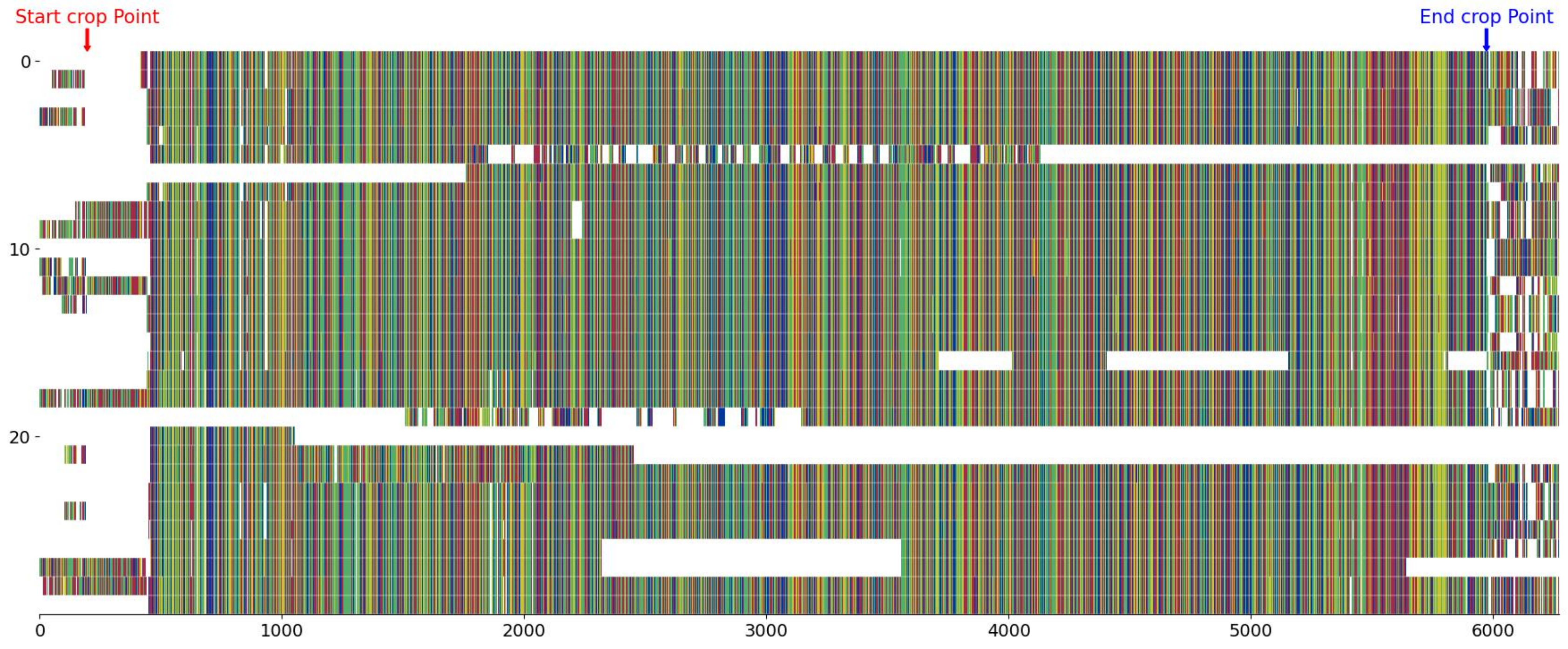


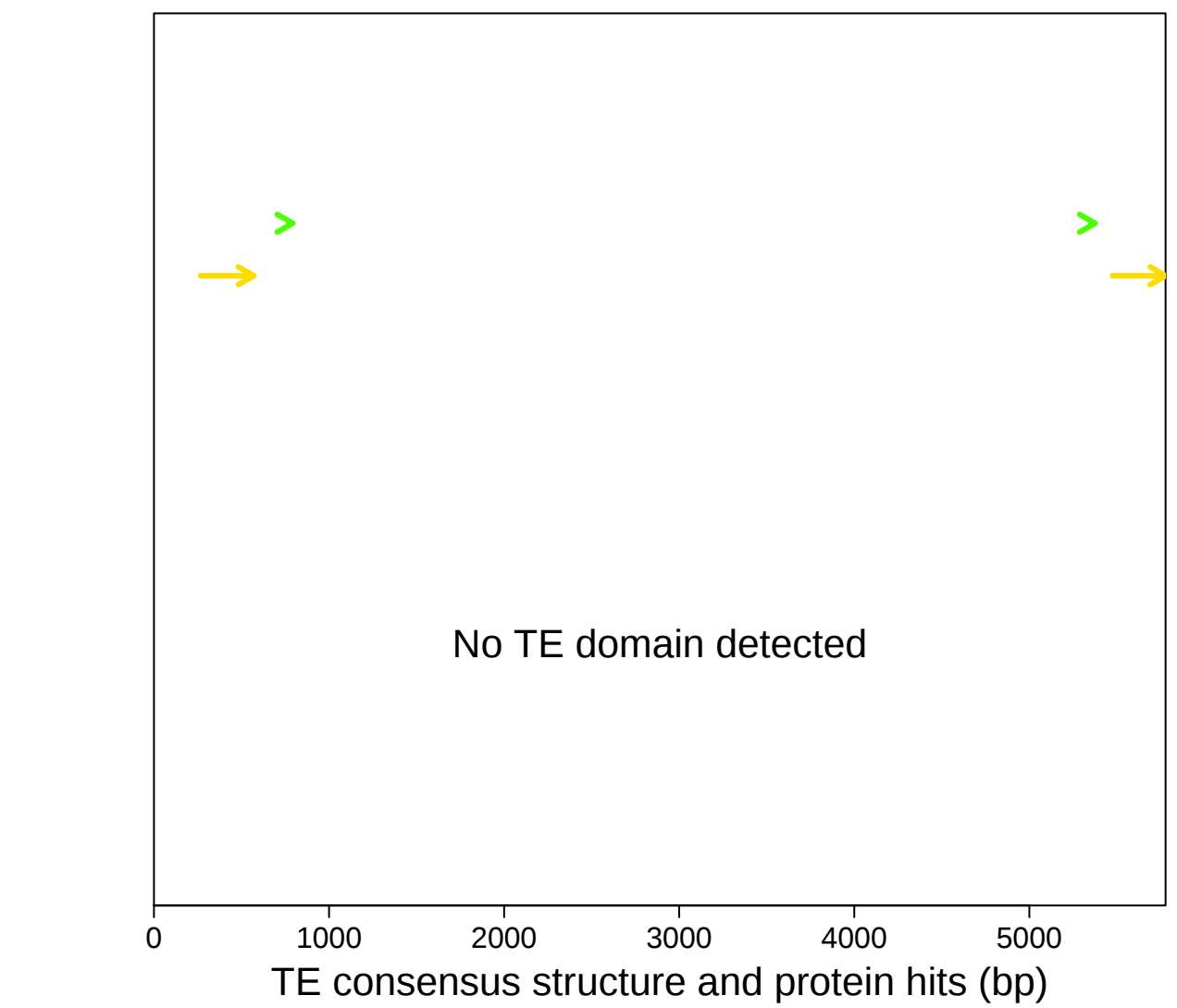
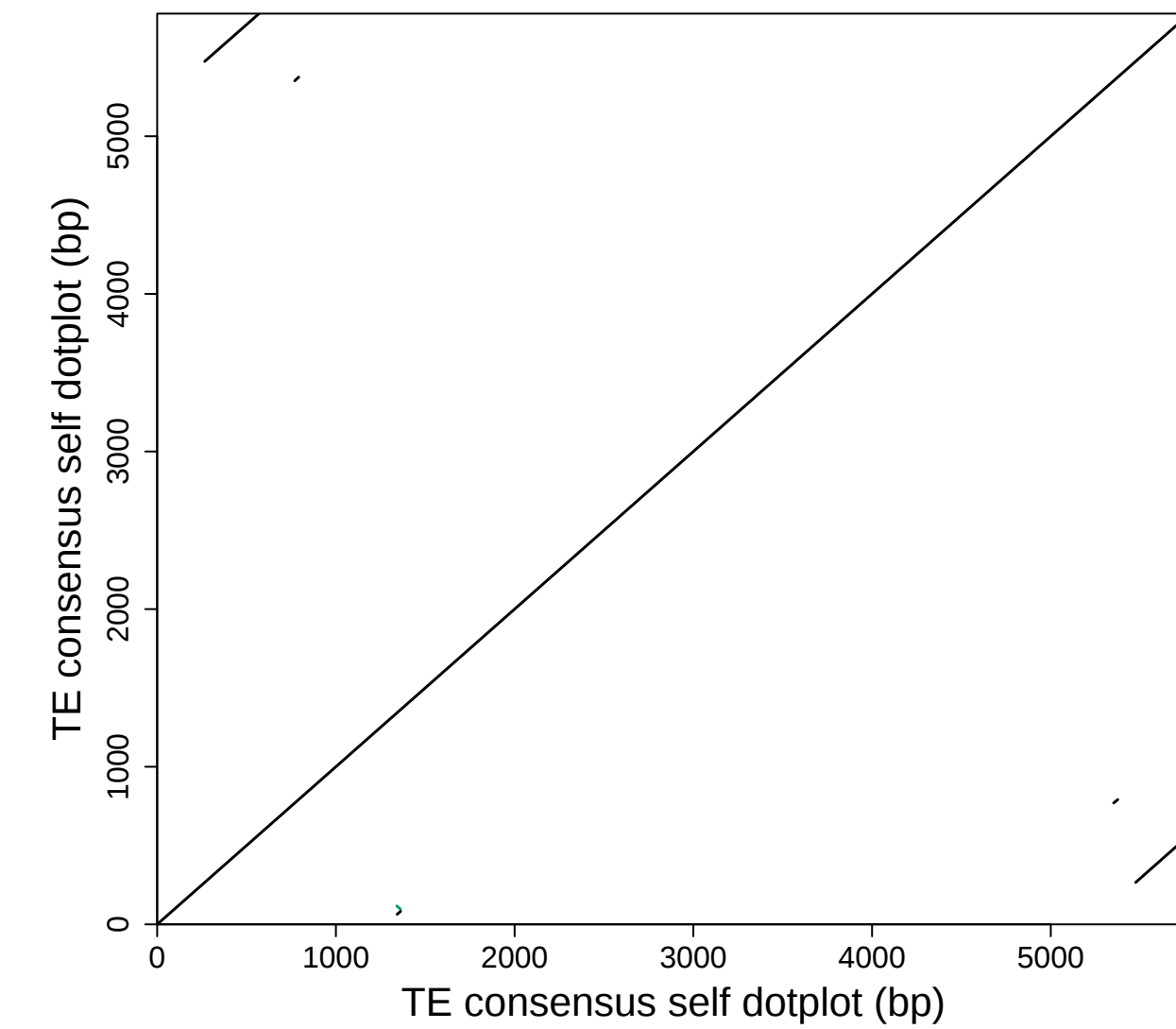
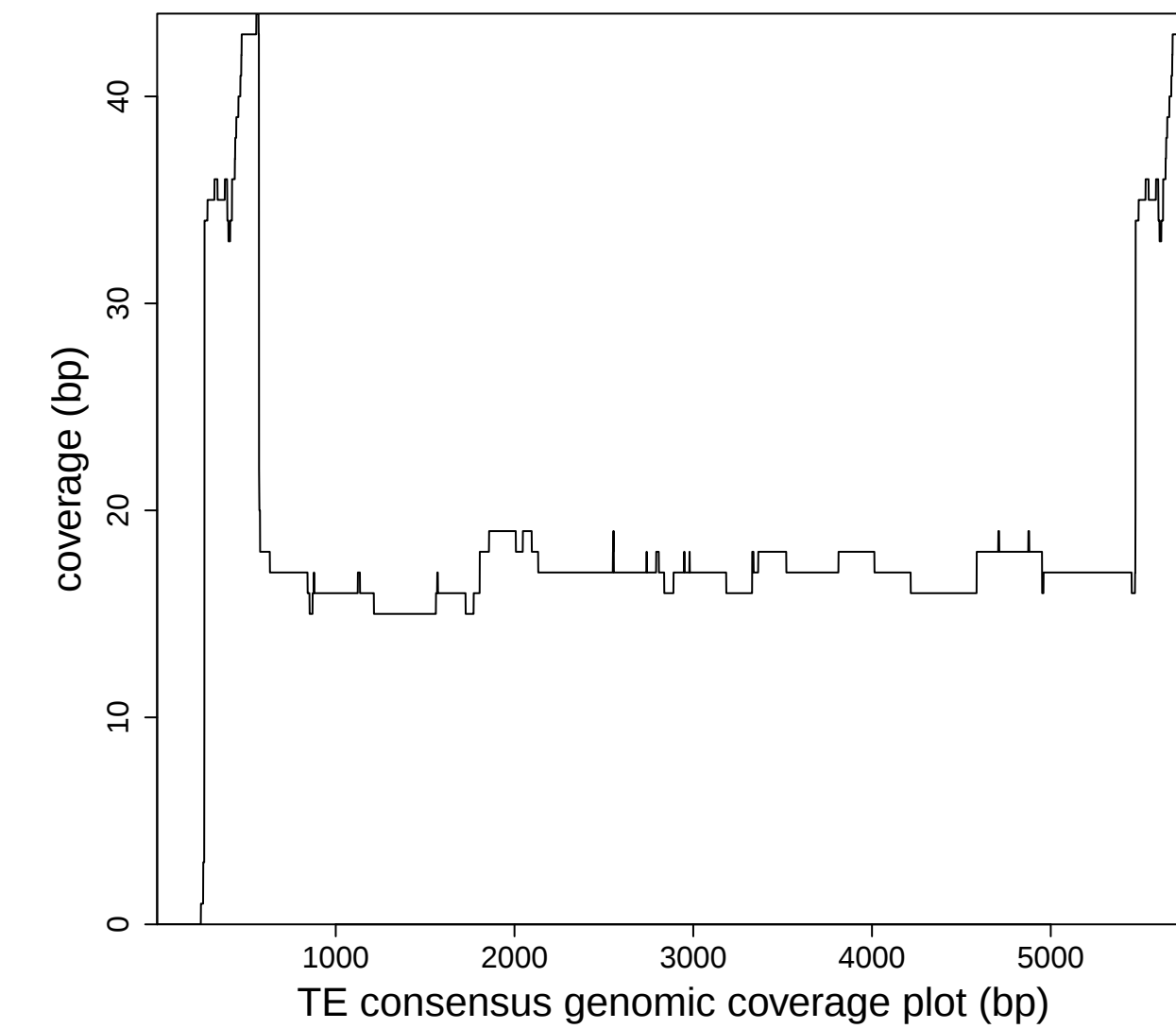
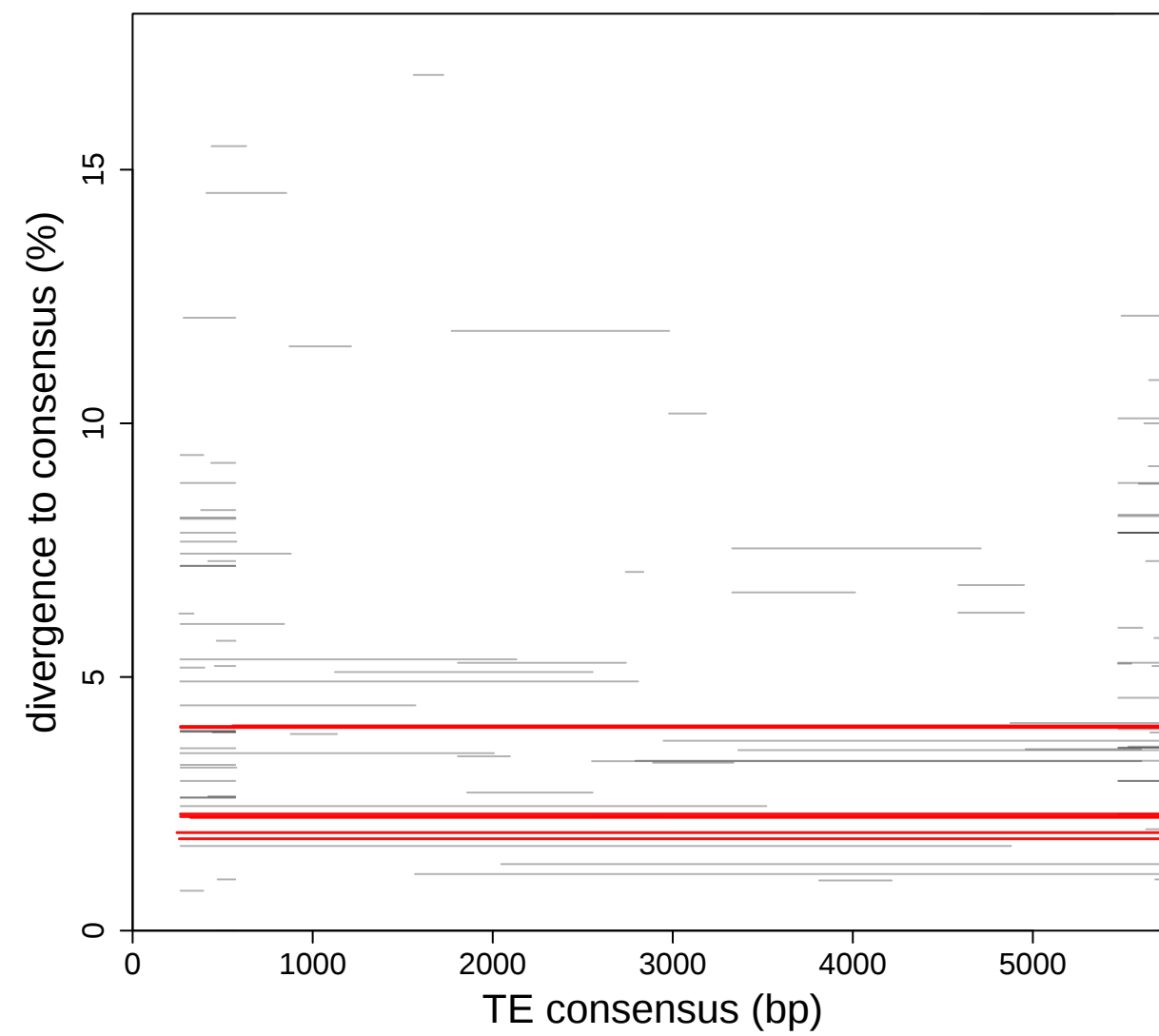
MSA length = 5778





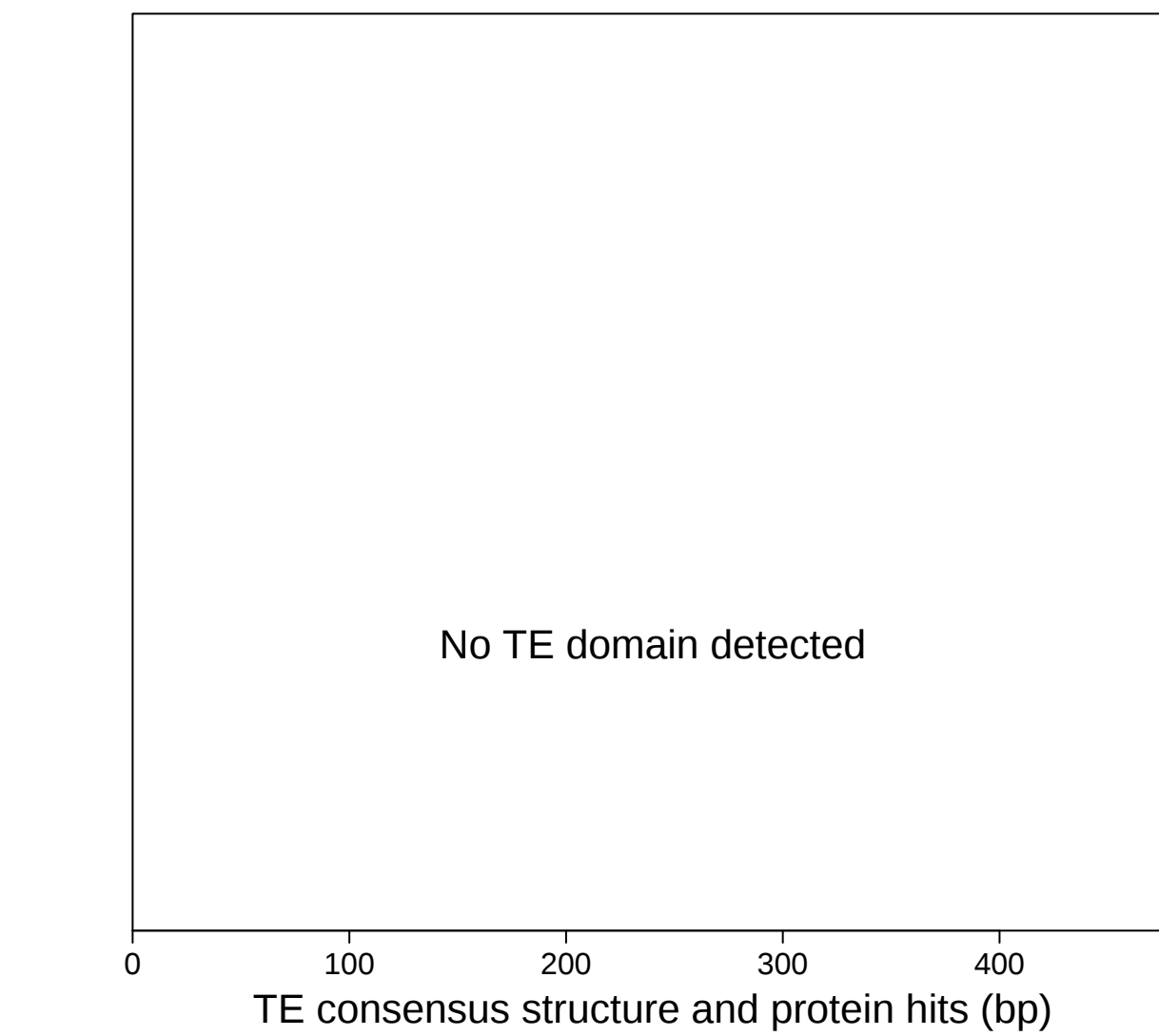
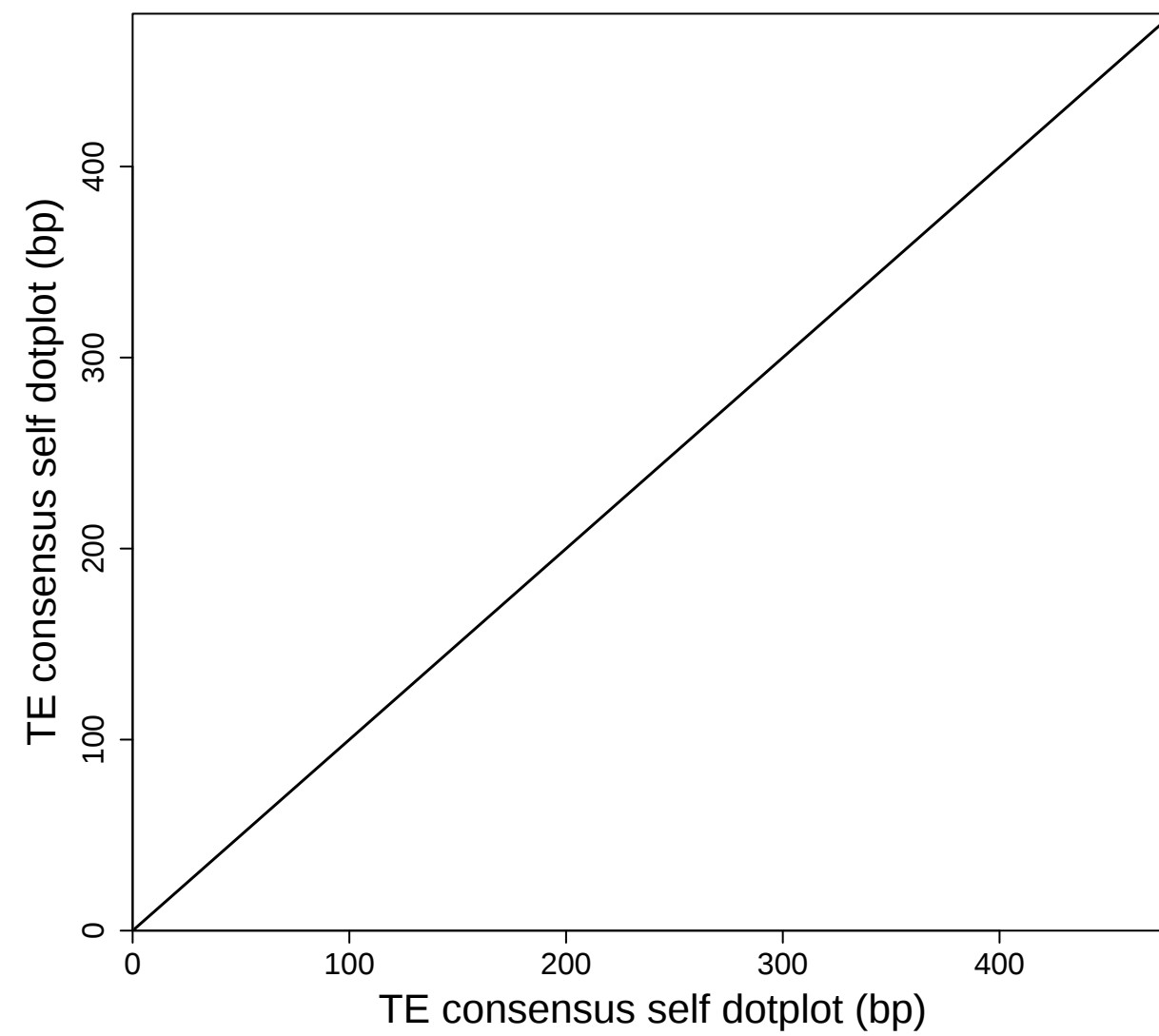
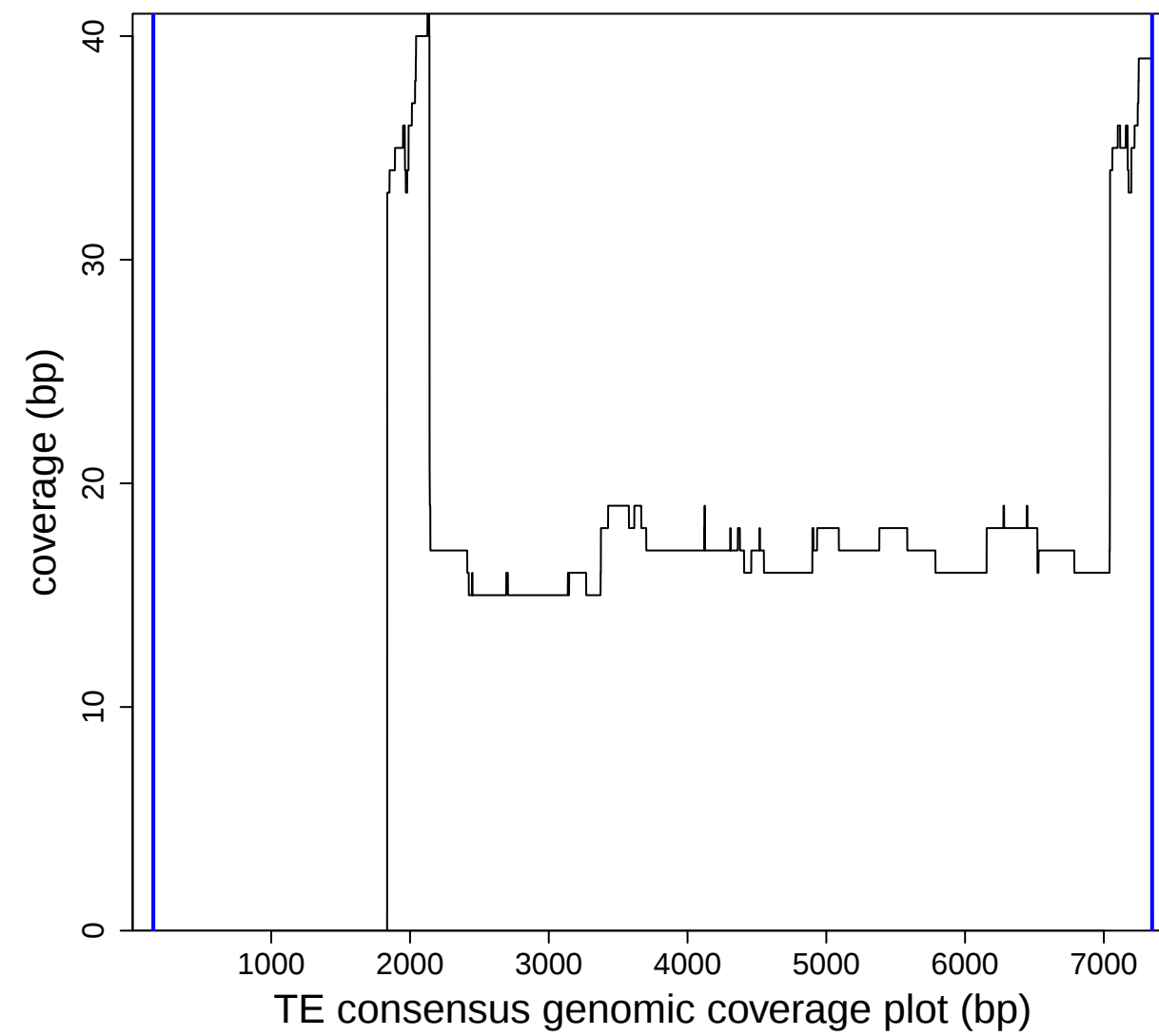
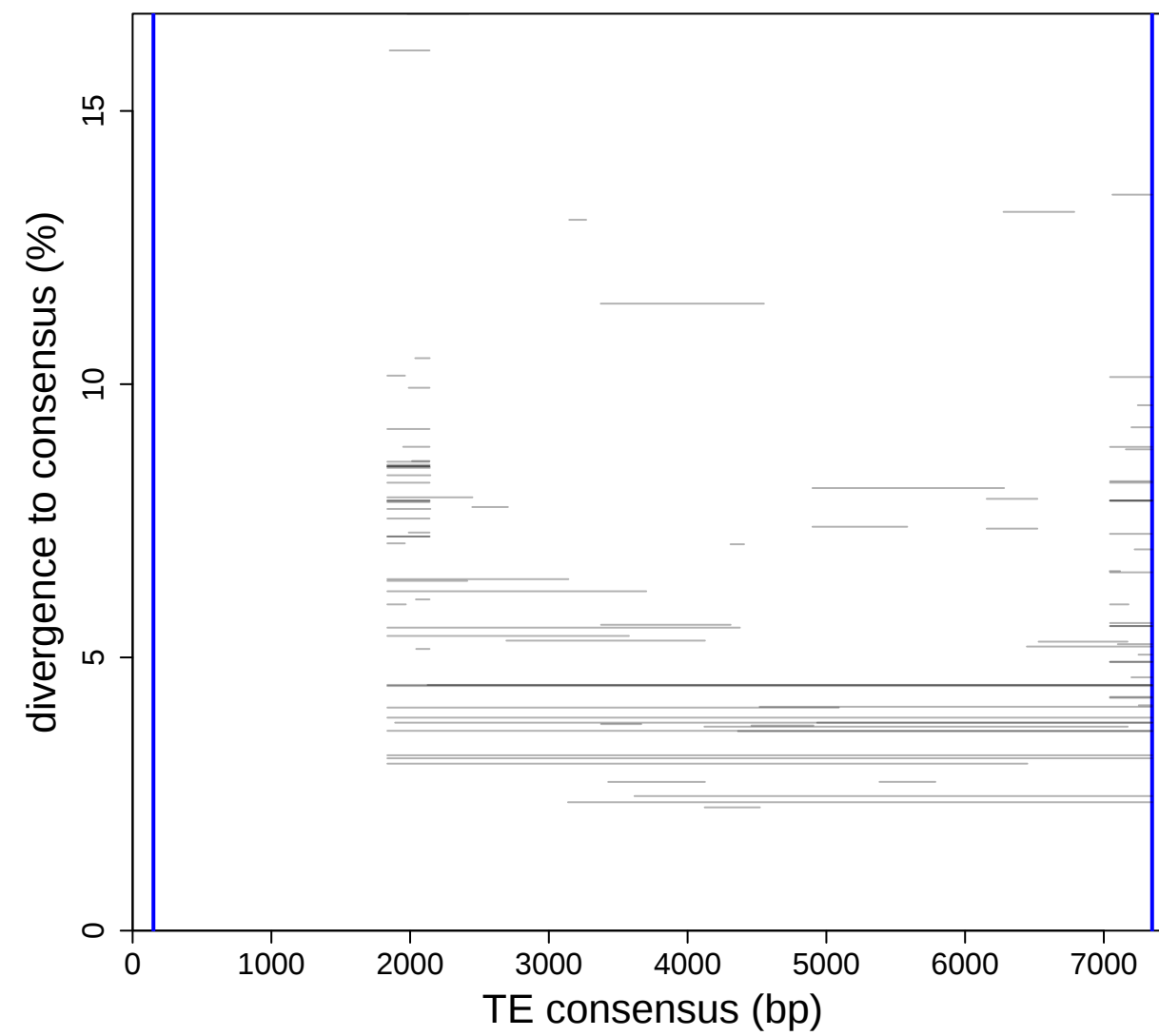
_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcIn.fa_al
size: 5778bp; fragments: 105; full length: 8 (>=5200.2bp)

After TETrimmer 5778 bp



bed_g_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bclr
size: 7498bp; fragments: 95; full length: 0 (>=6748.2bp)

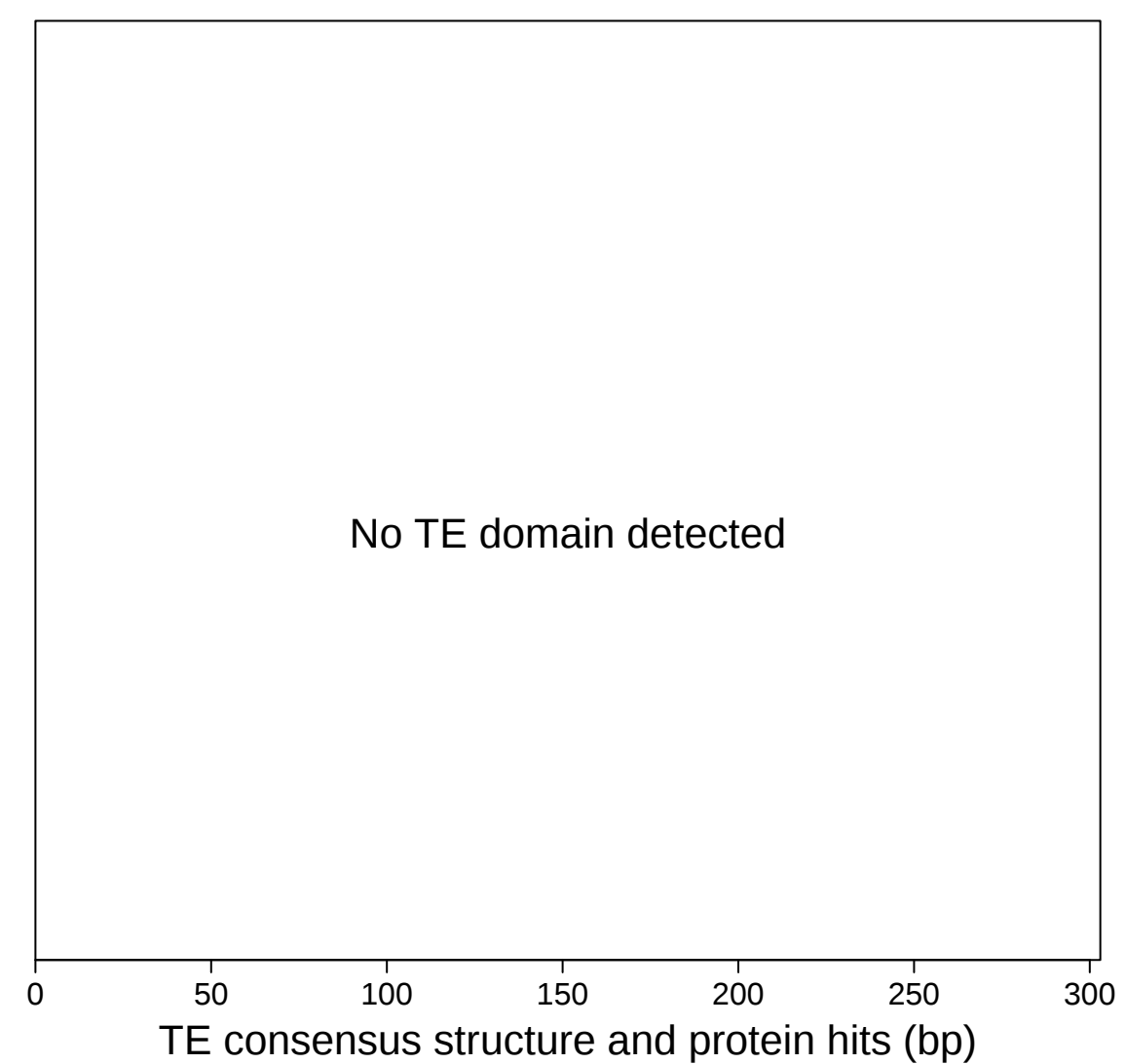
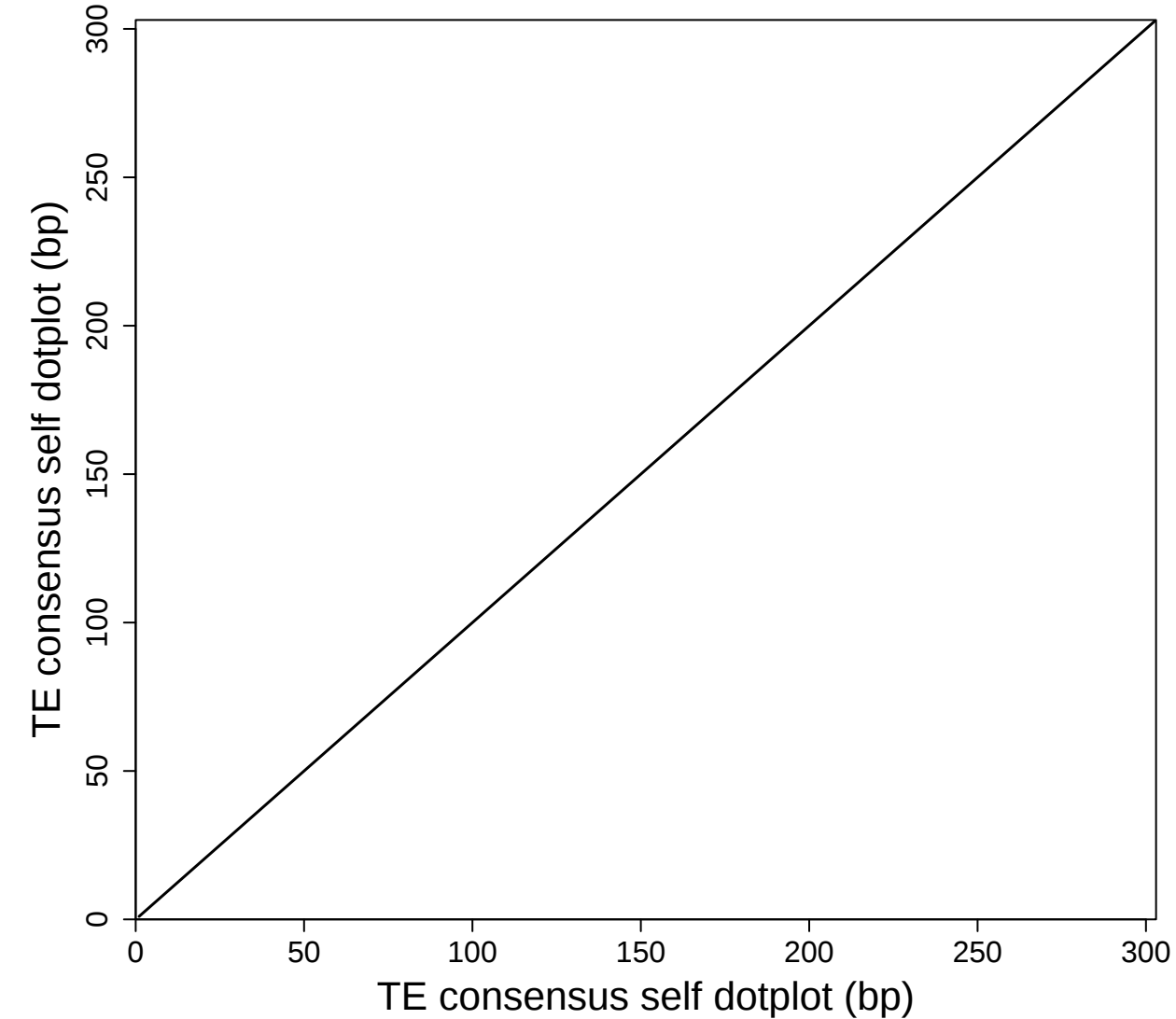
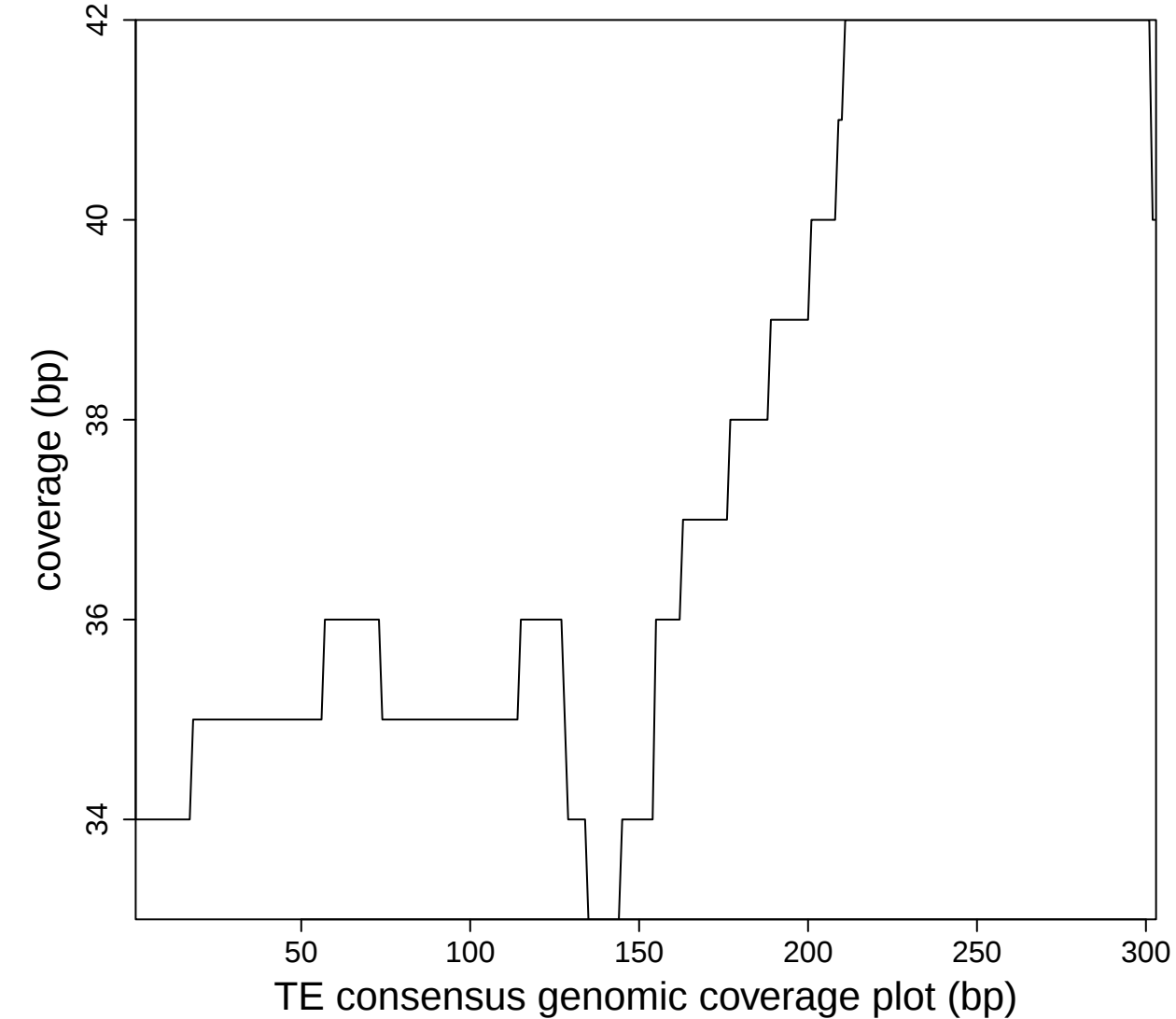
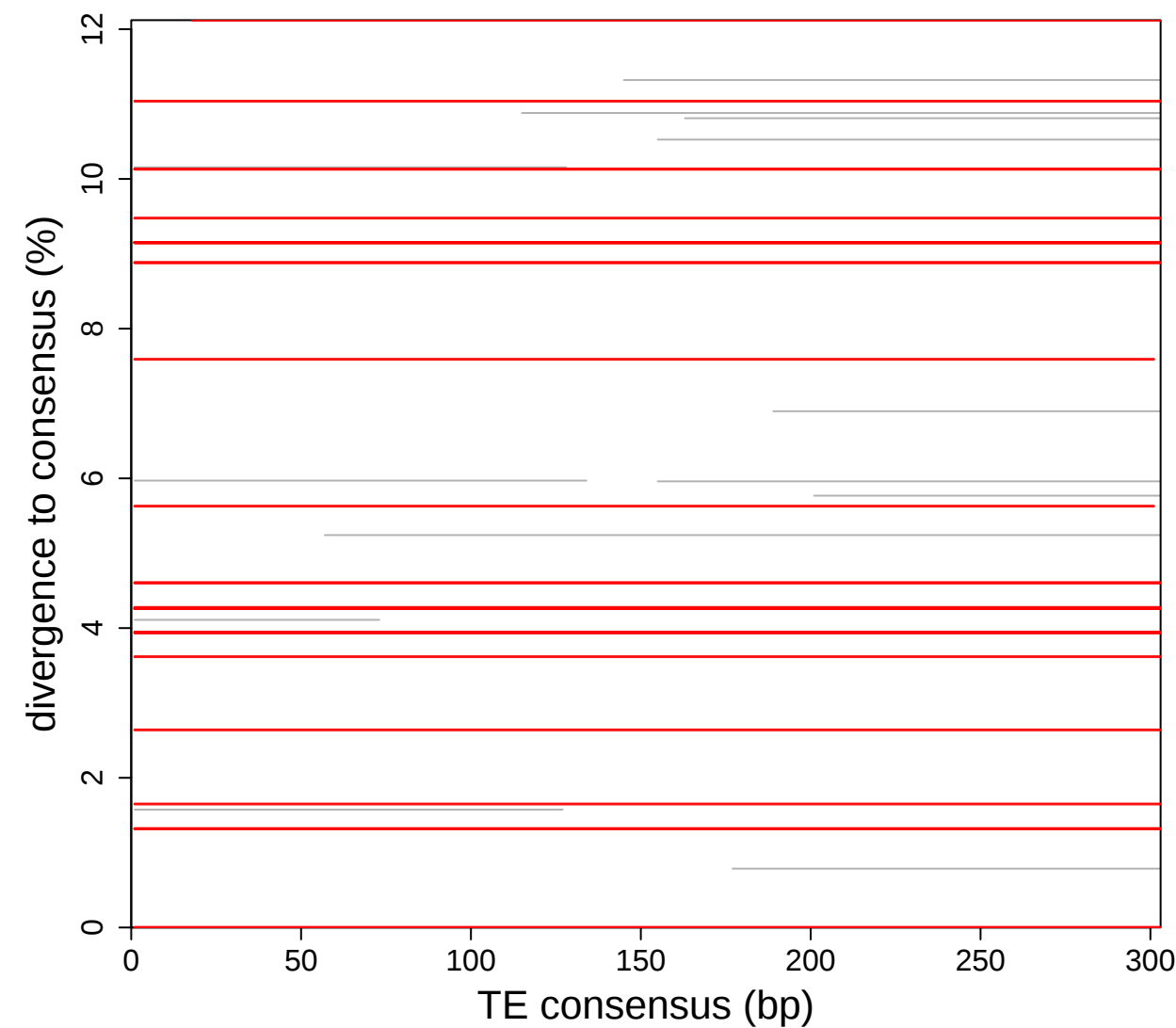
After TEtrimmer Extended plot Blue lines are boundaries



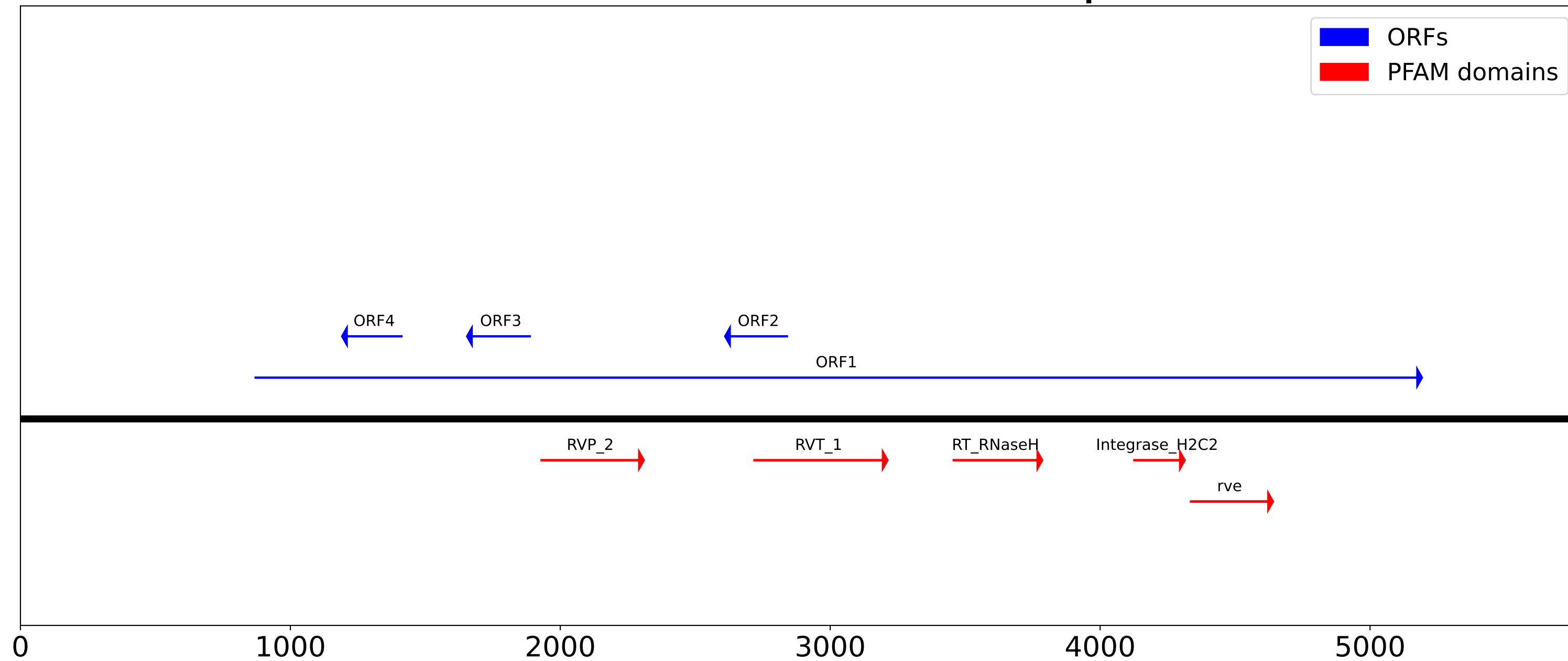
Before TEtrimmer 303 bp

TE: ltr_1_family_316

size: 303bp; fragments: 46; full length: 31 (≥ 272.7 bp)



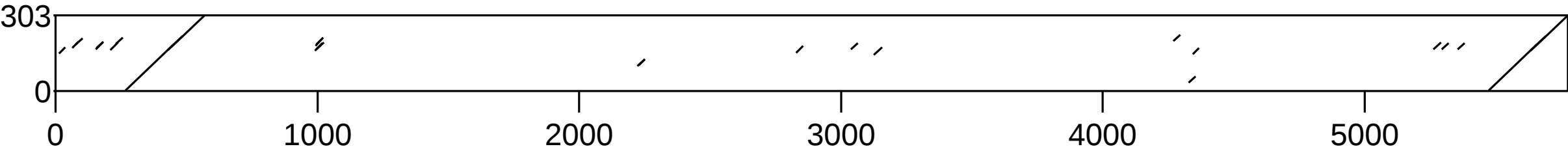
After TEtrimmer ORF and PFAM domain plot



Dotplot

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

Consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)