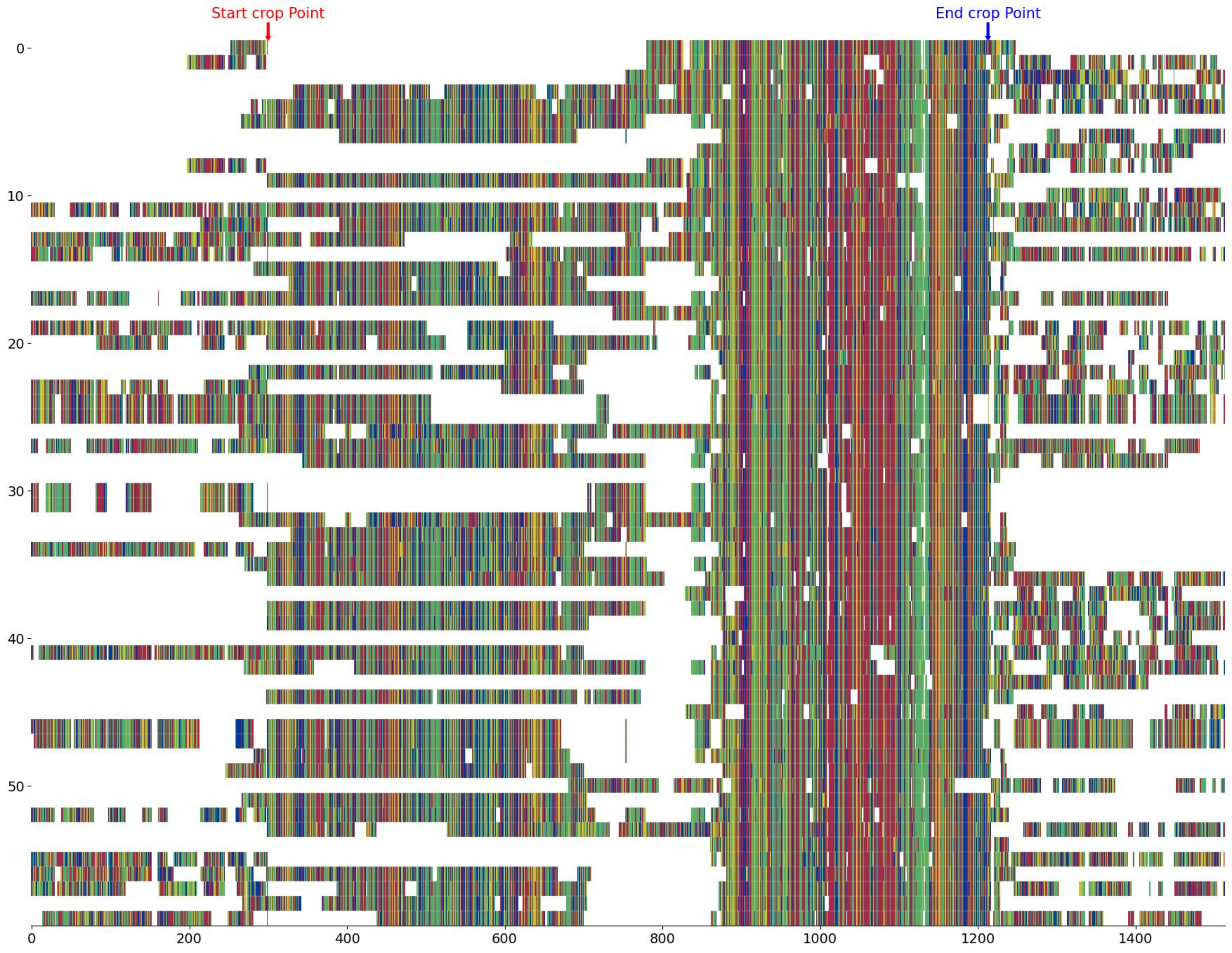


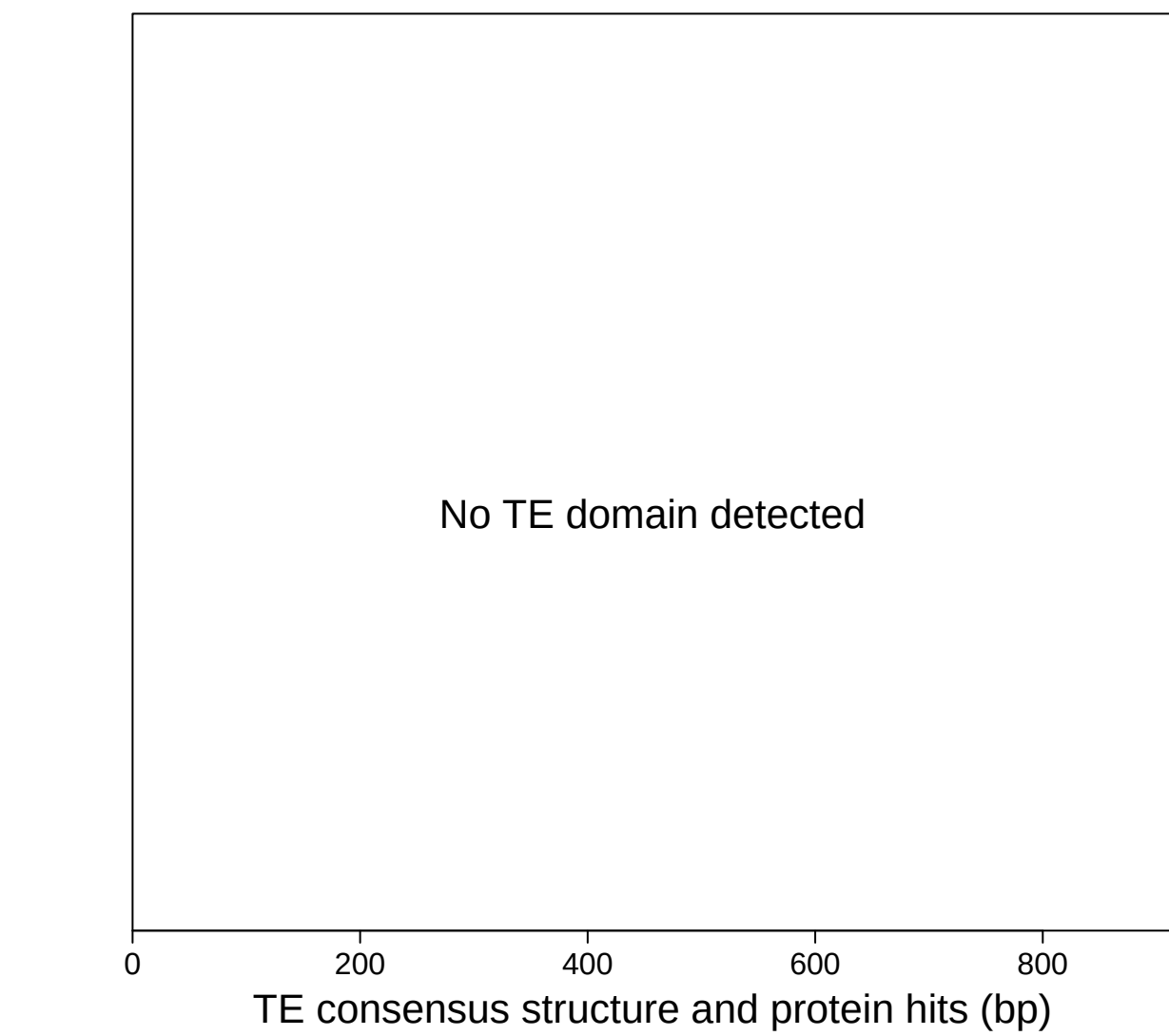
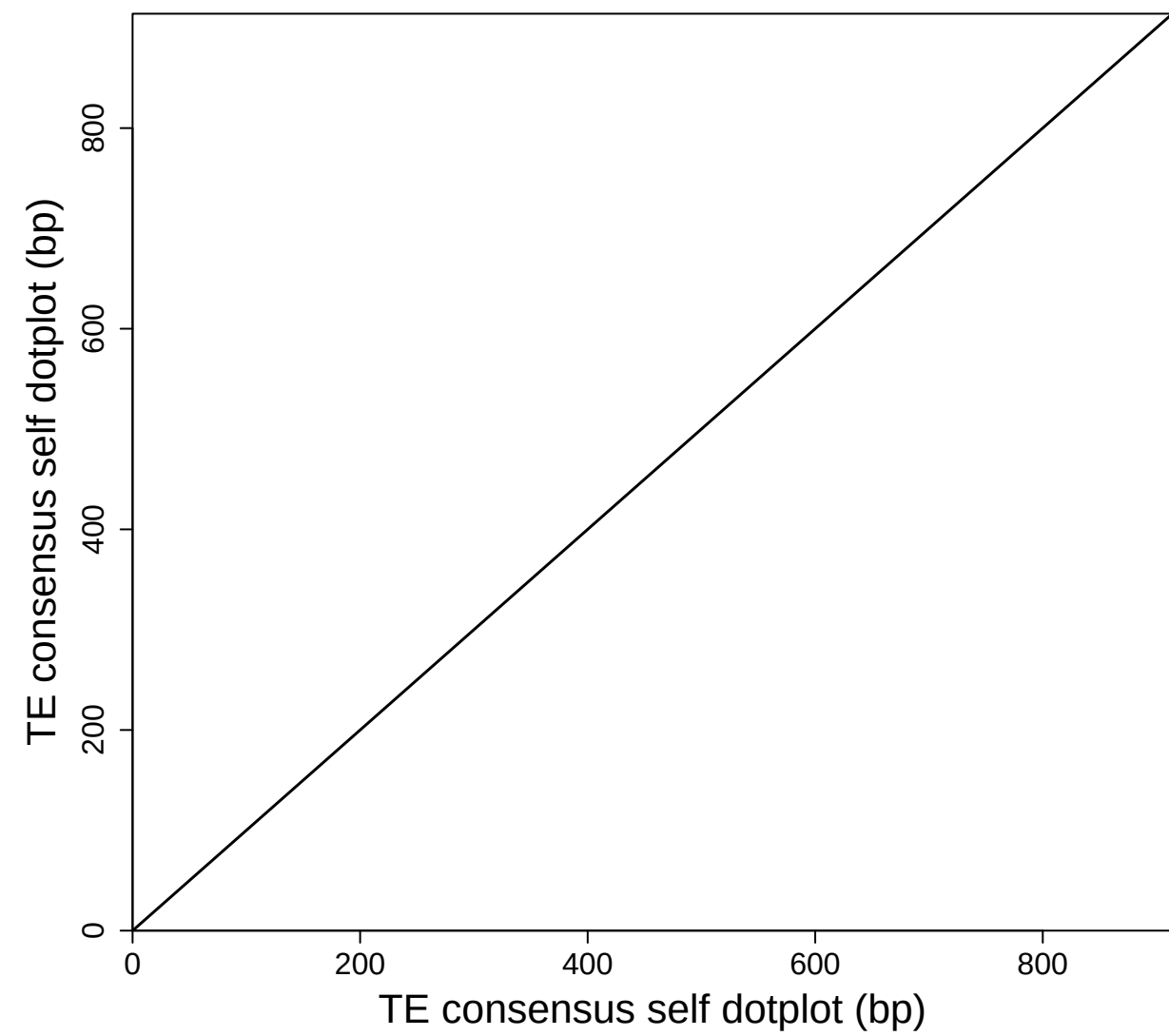
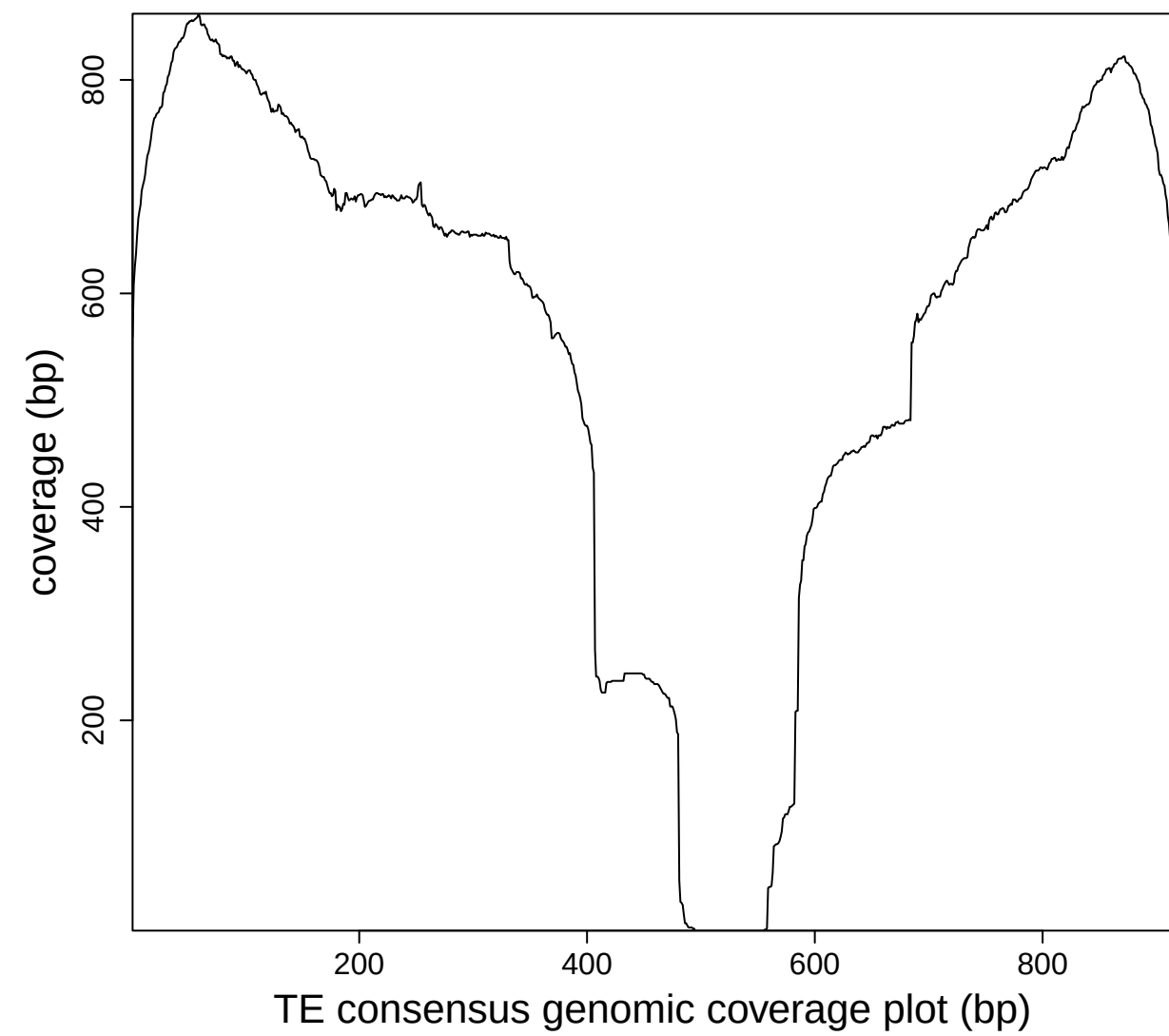
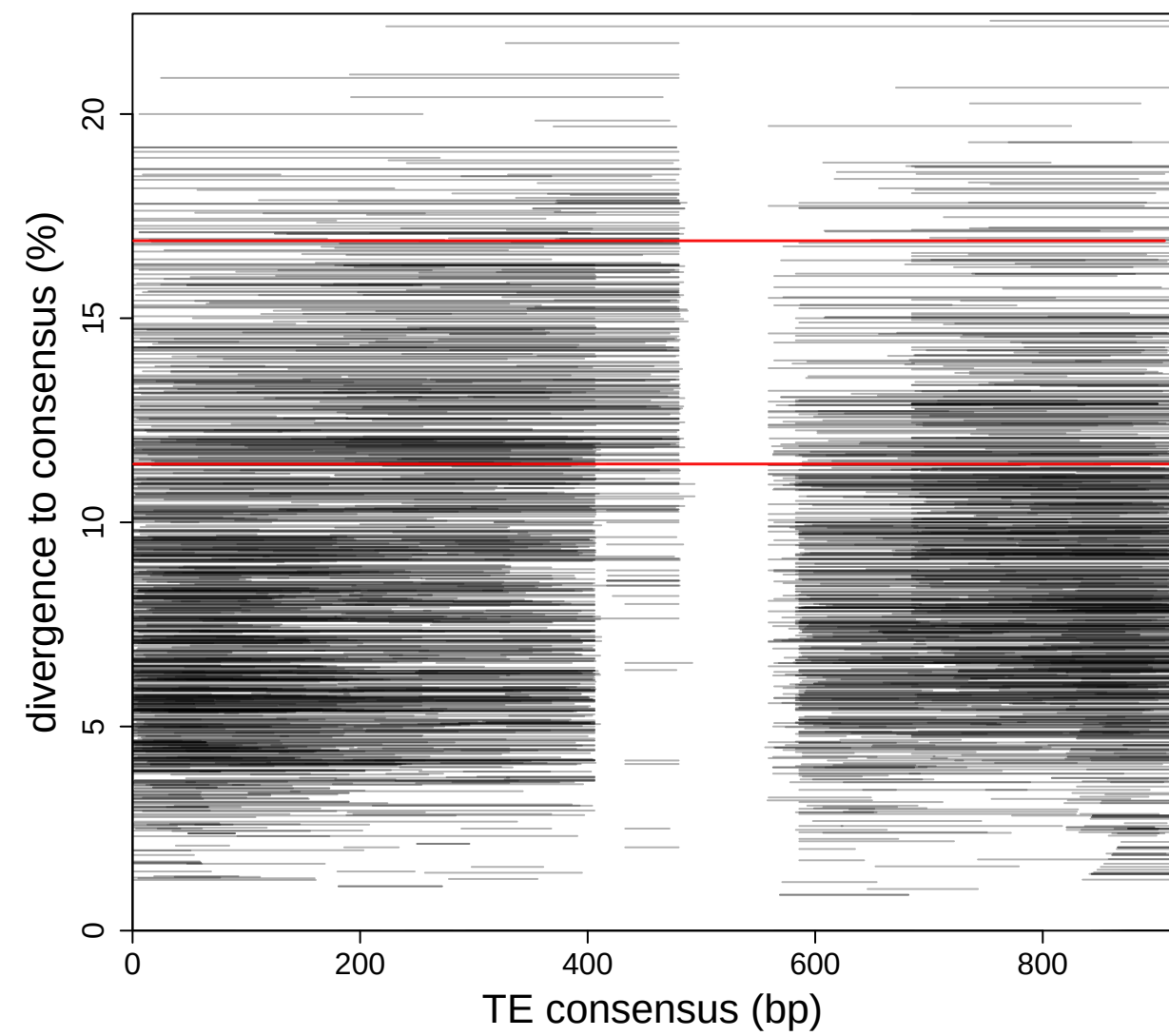




g_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcln.fa_a

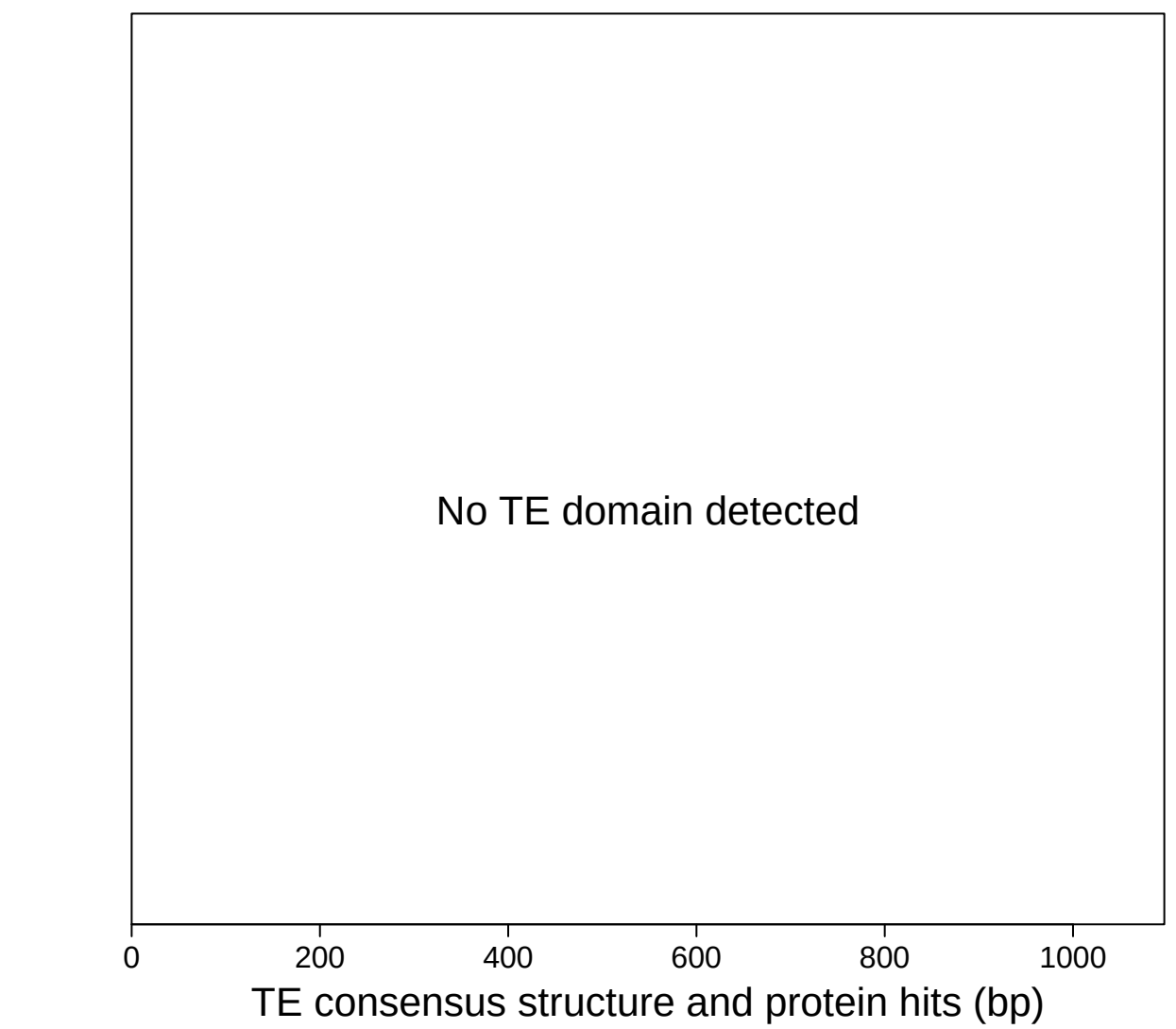
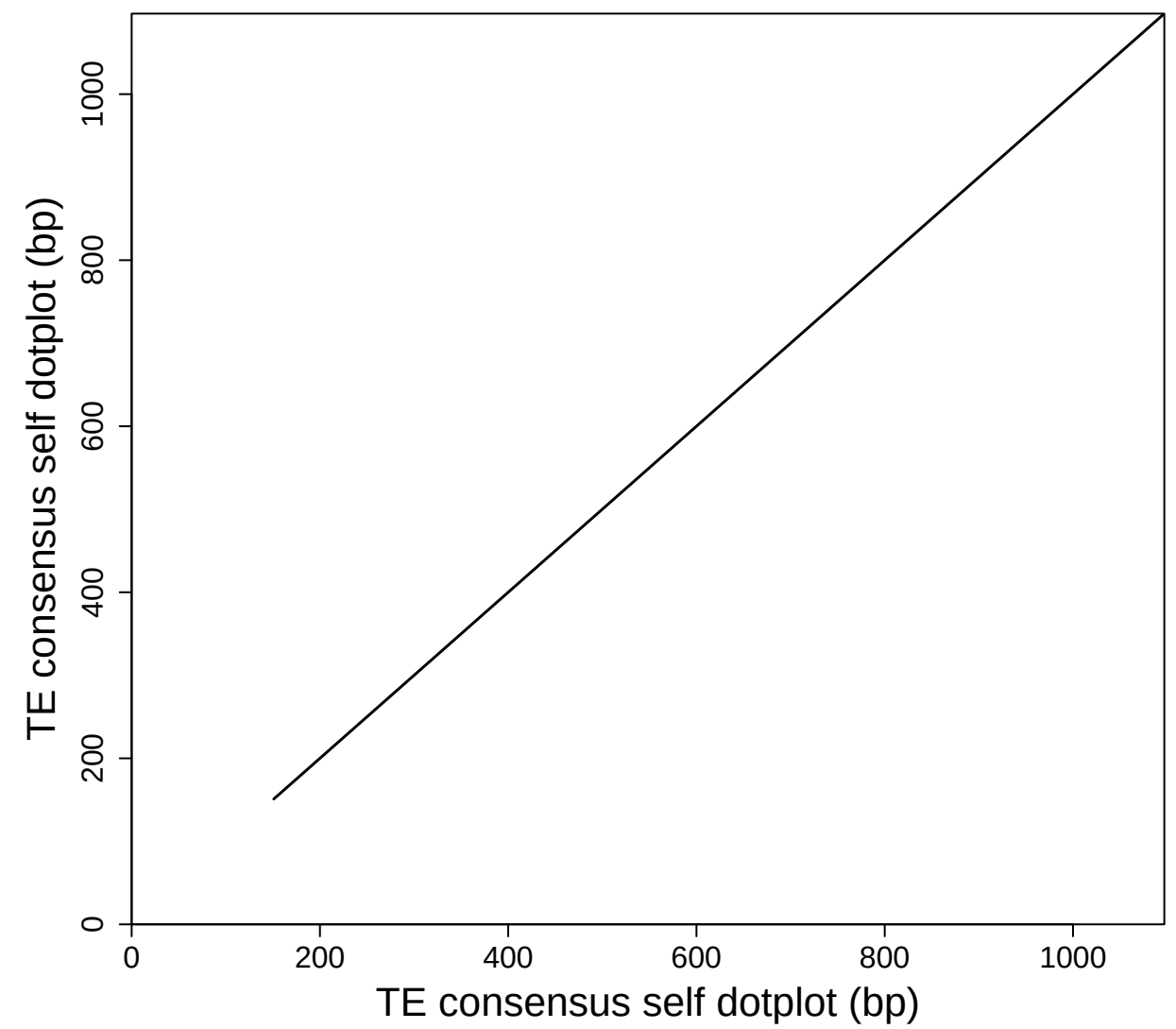
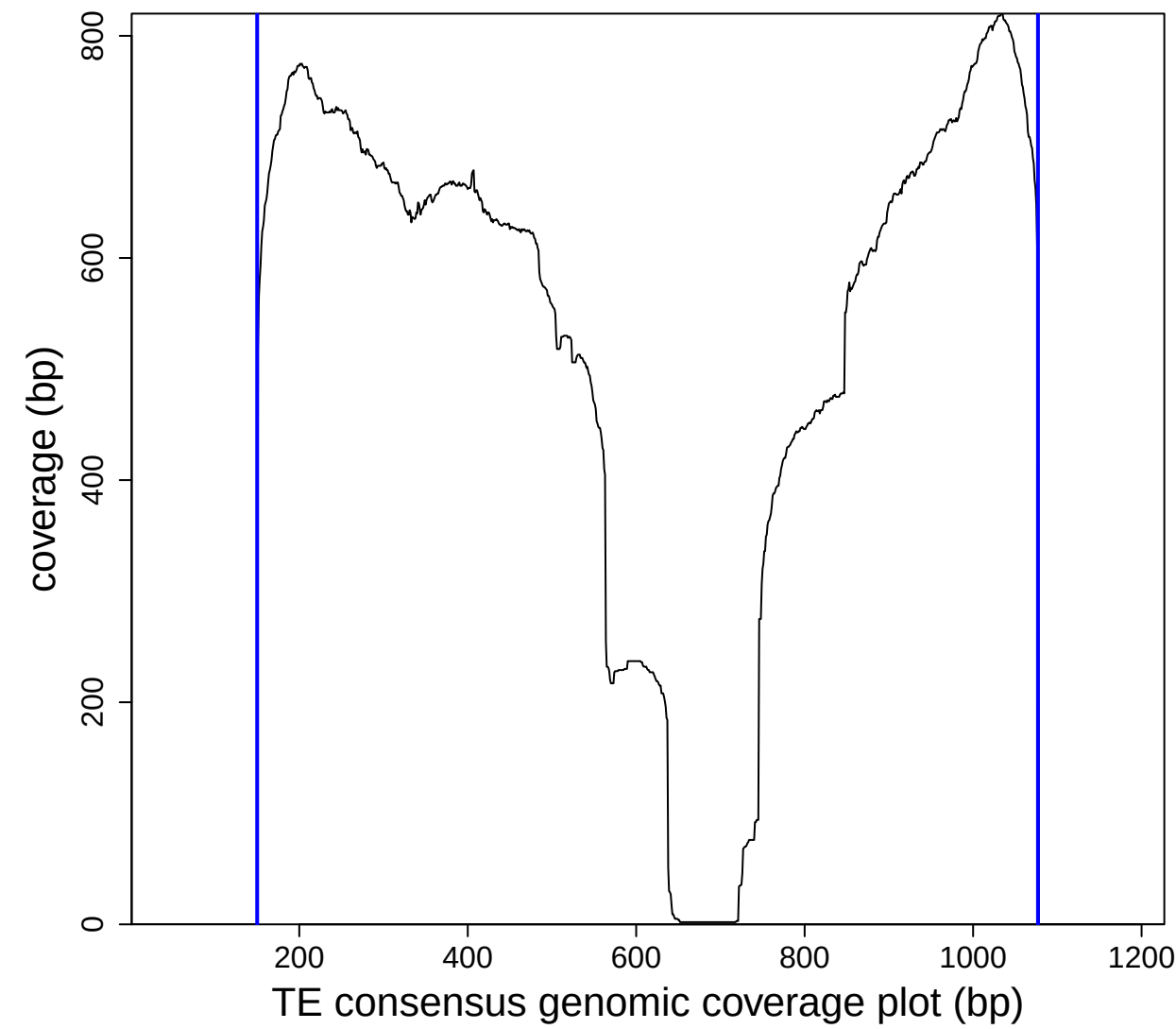
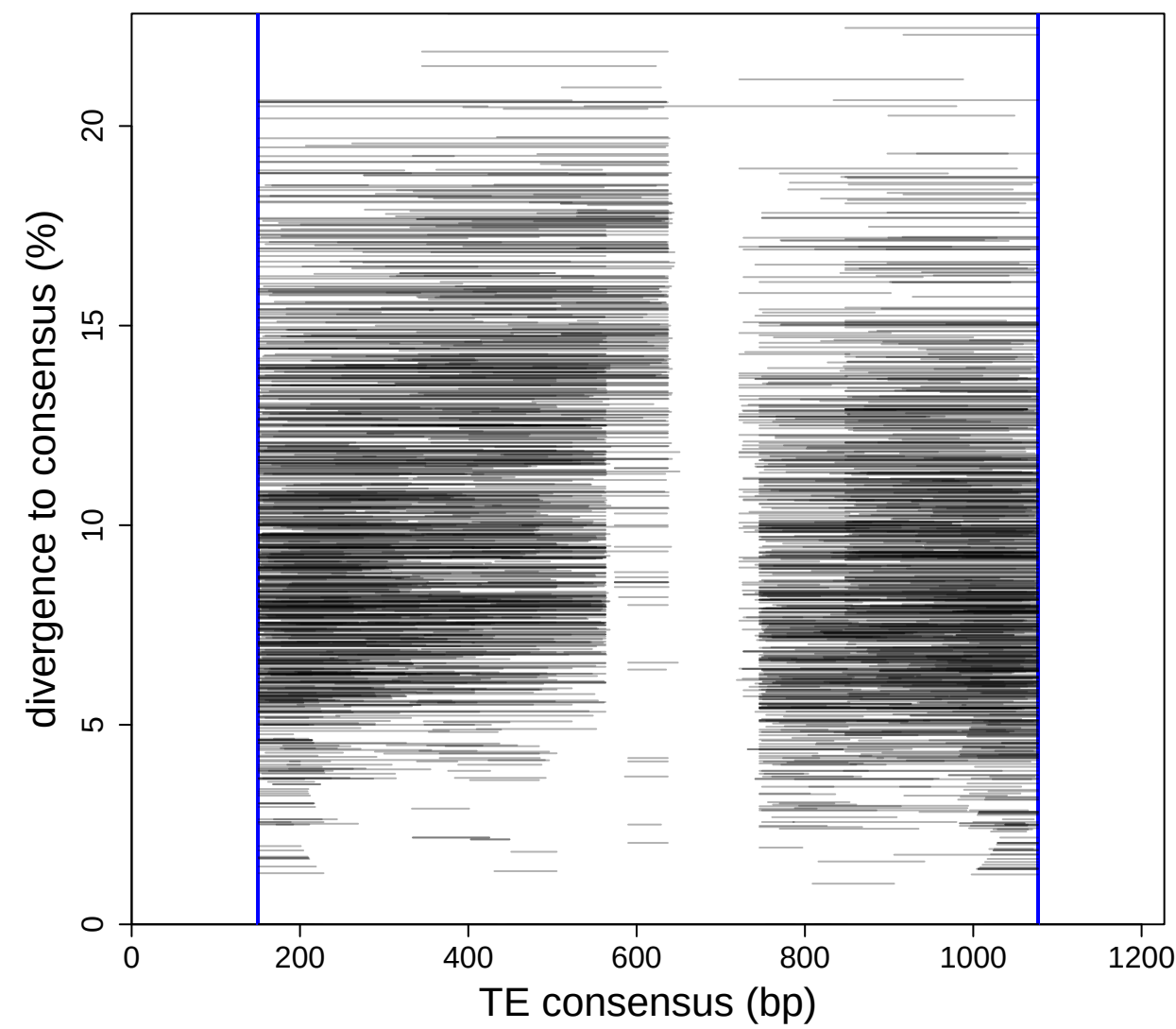
size: 914bp; fragments: 2343; full length: 2 (>=822.6bp)

After TEtrimmer 914 bp

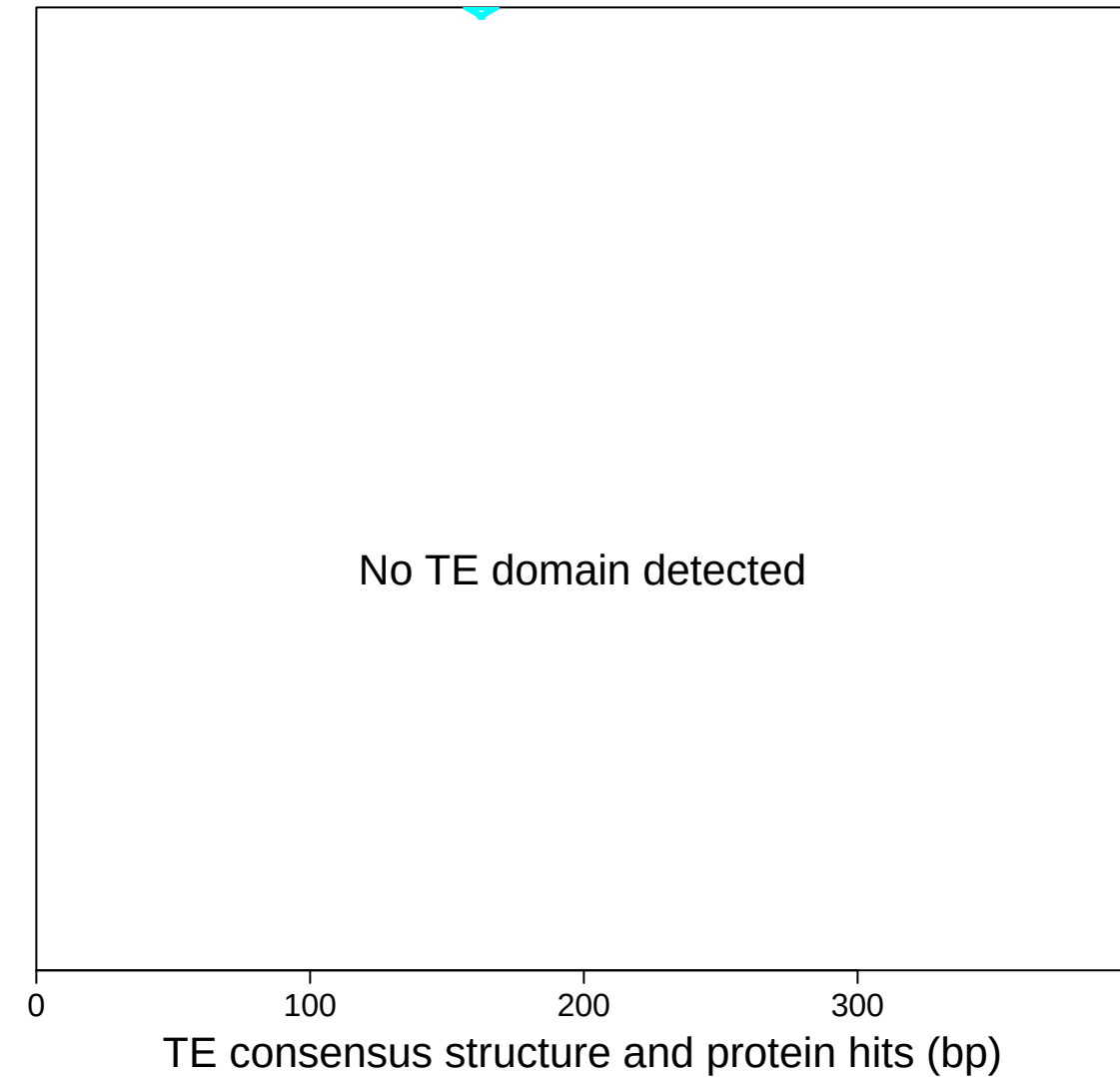
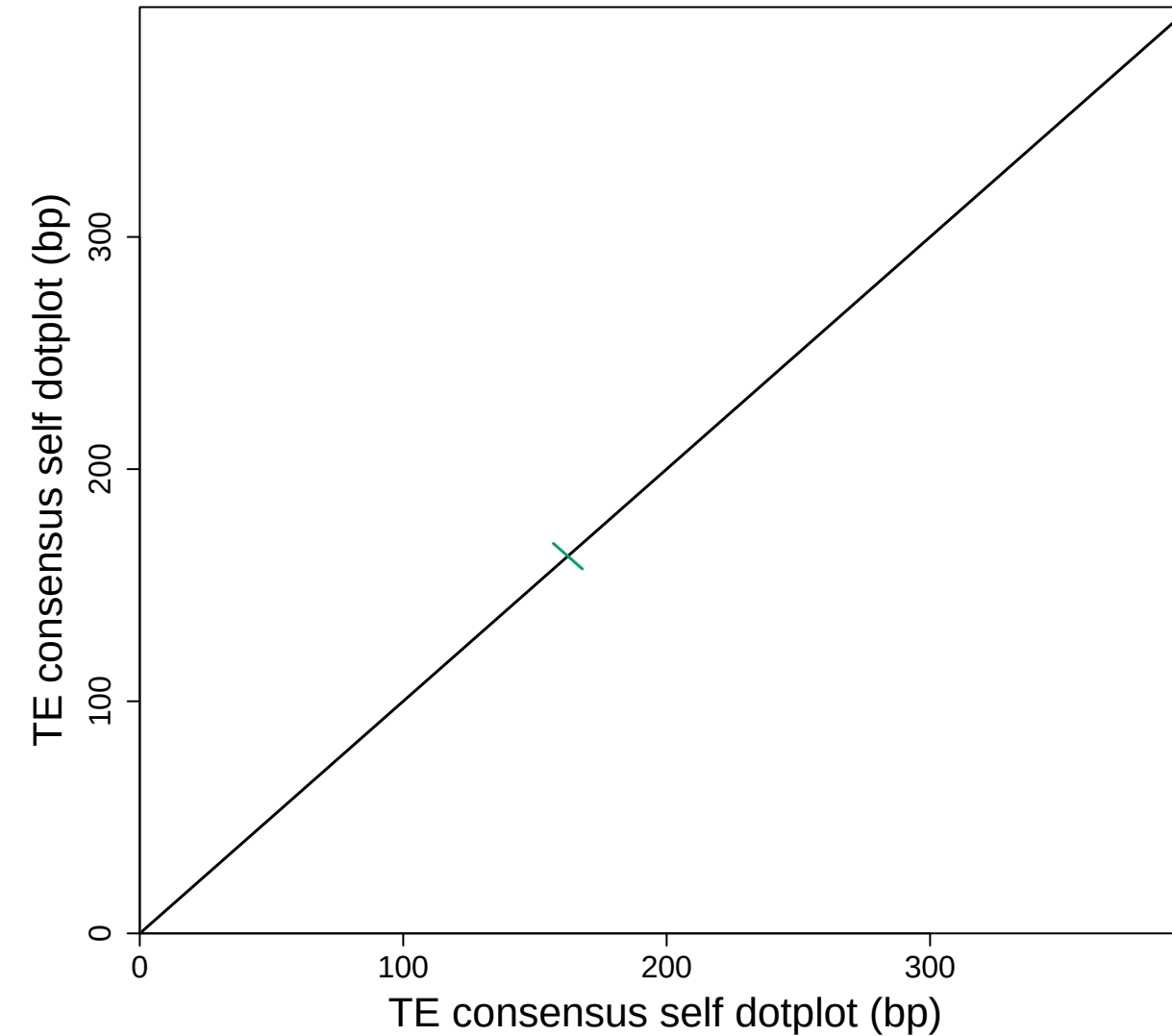
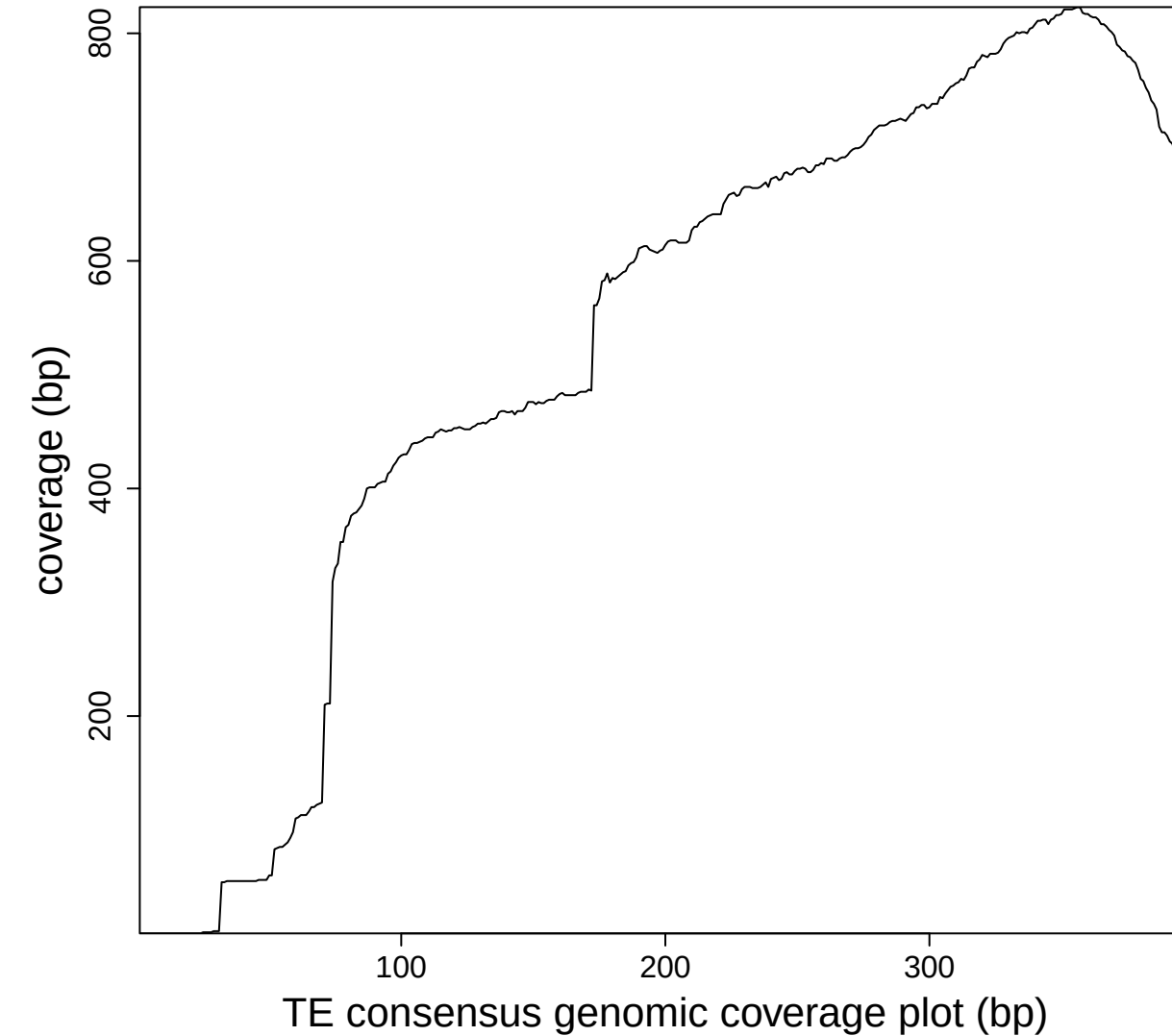
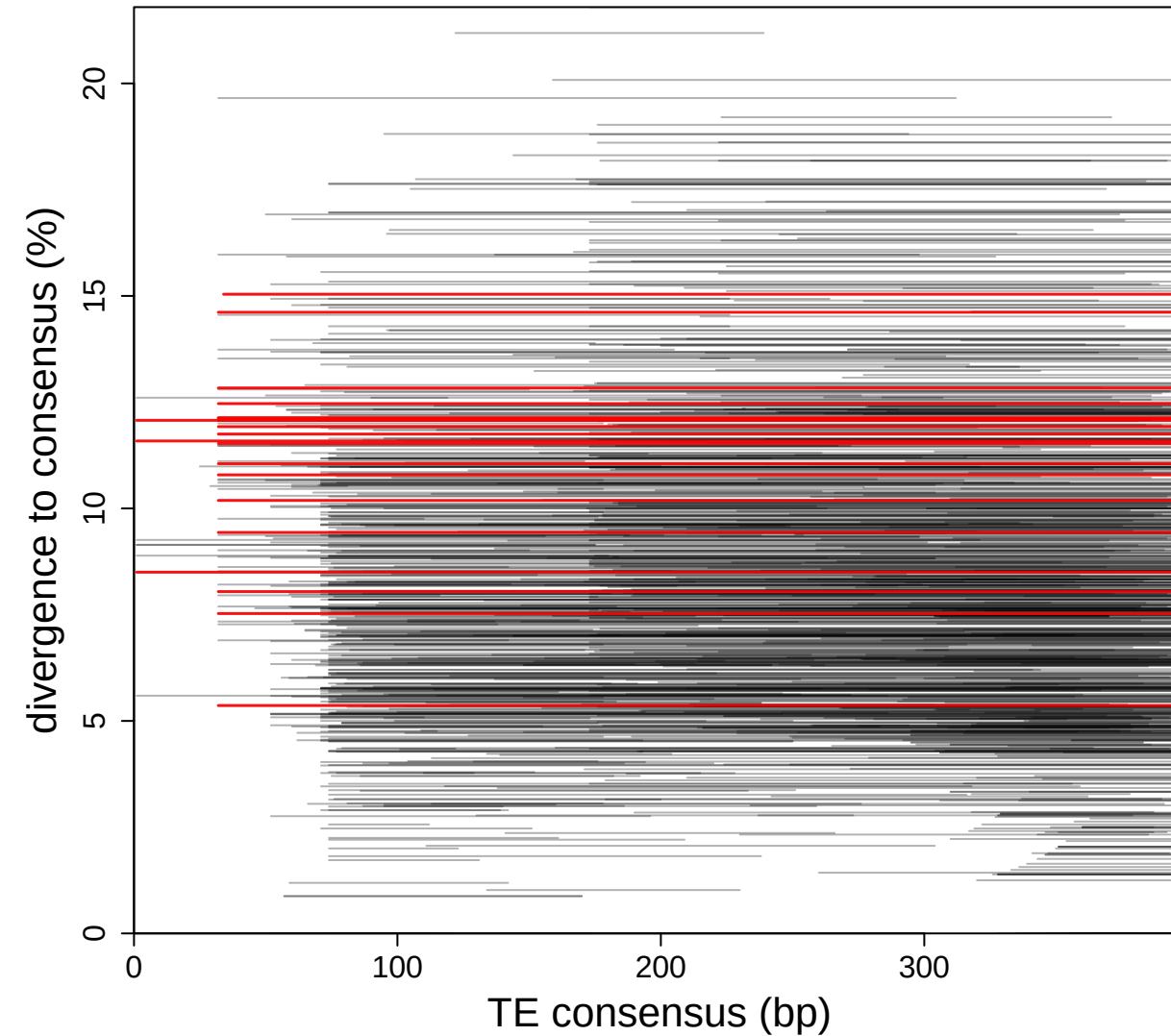


f.bed_g_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcl
size: 1227bp; fragments: 2215; full length: 0 (>=1104.3bp)

After TETrimmer Extended plot Blue lines are boundaries

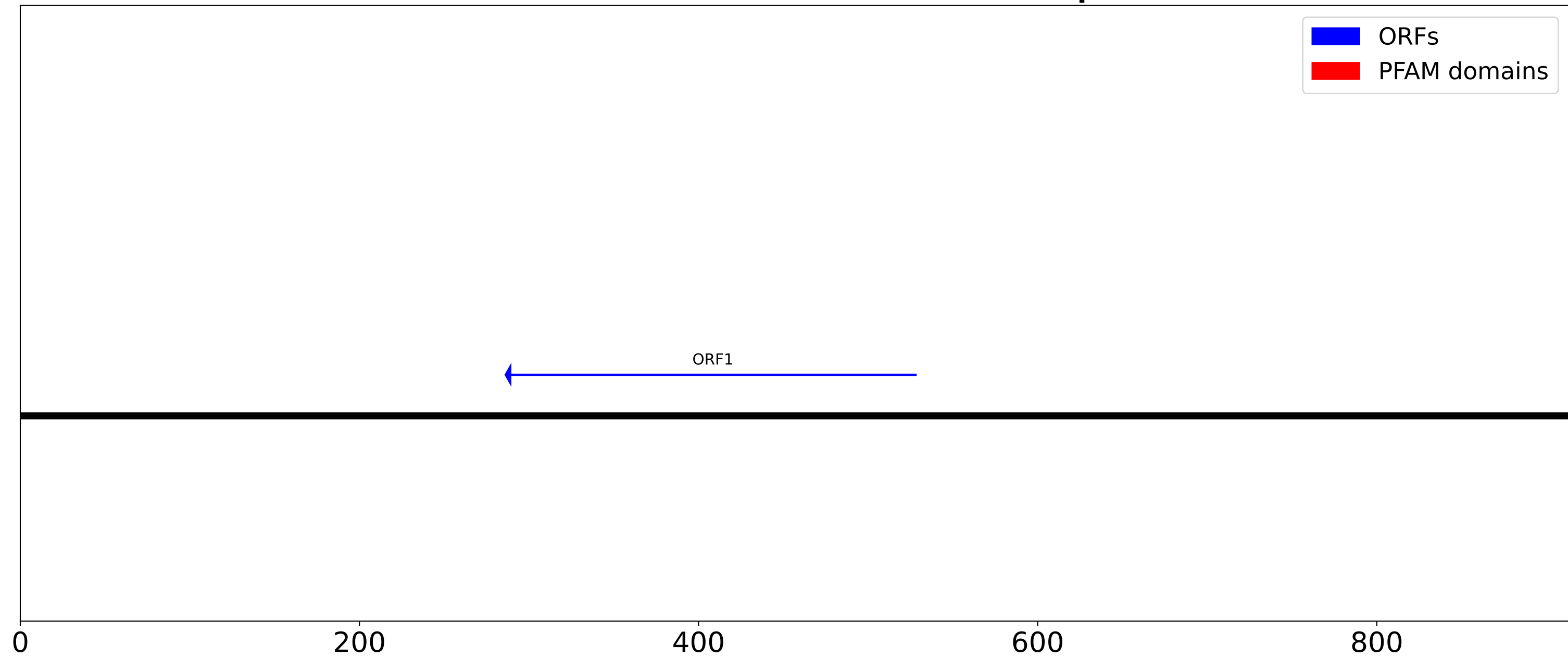


TE: rnd_5_family_808
size: 399bp; fragments: 1053; full length: 20 (≥ 359.1 bp)



Before TEtrimmer 399 bp

After TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)

