

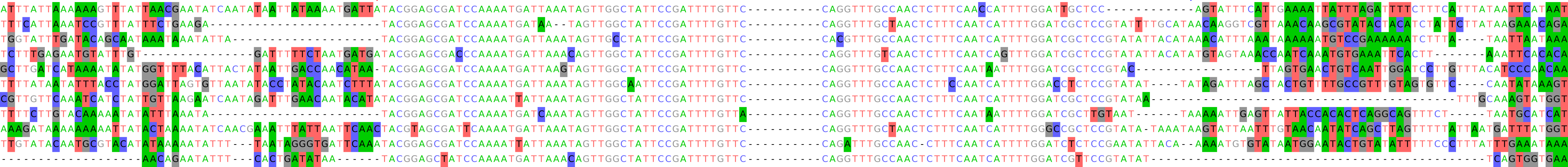
Start crop Point



End crop Point

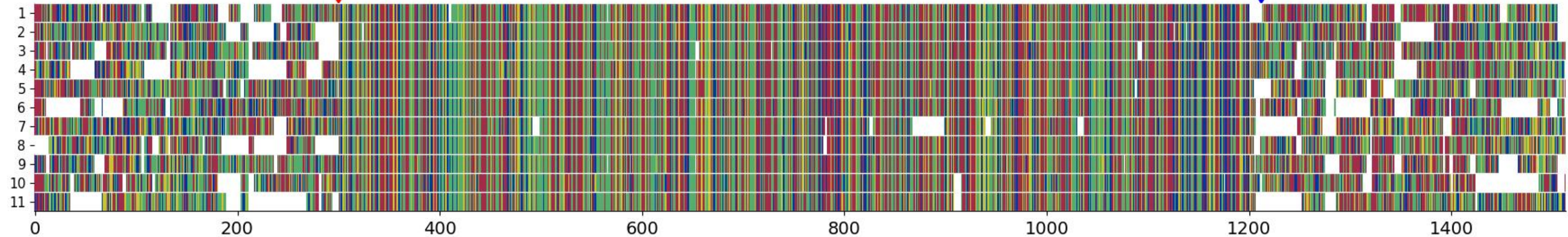


MSA length = 913



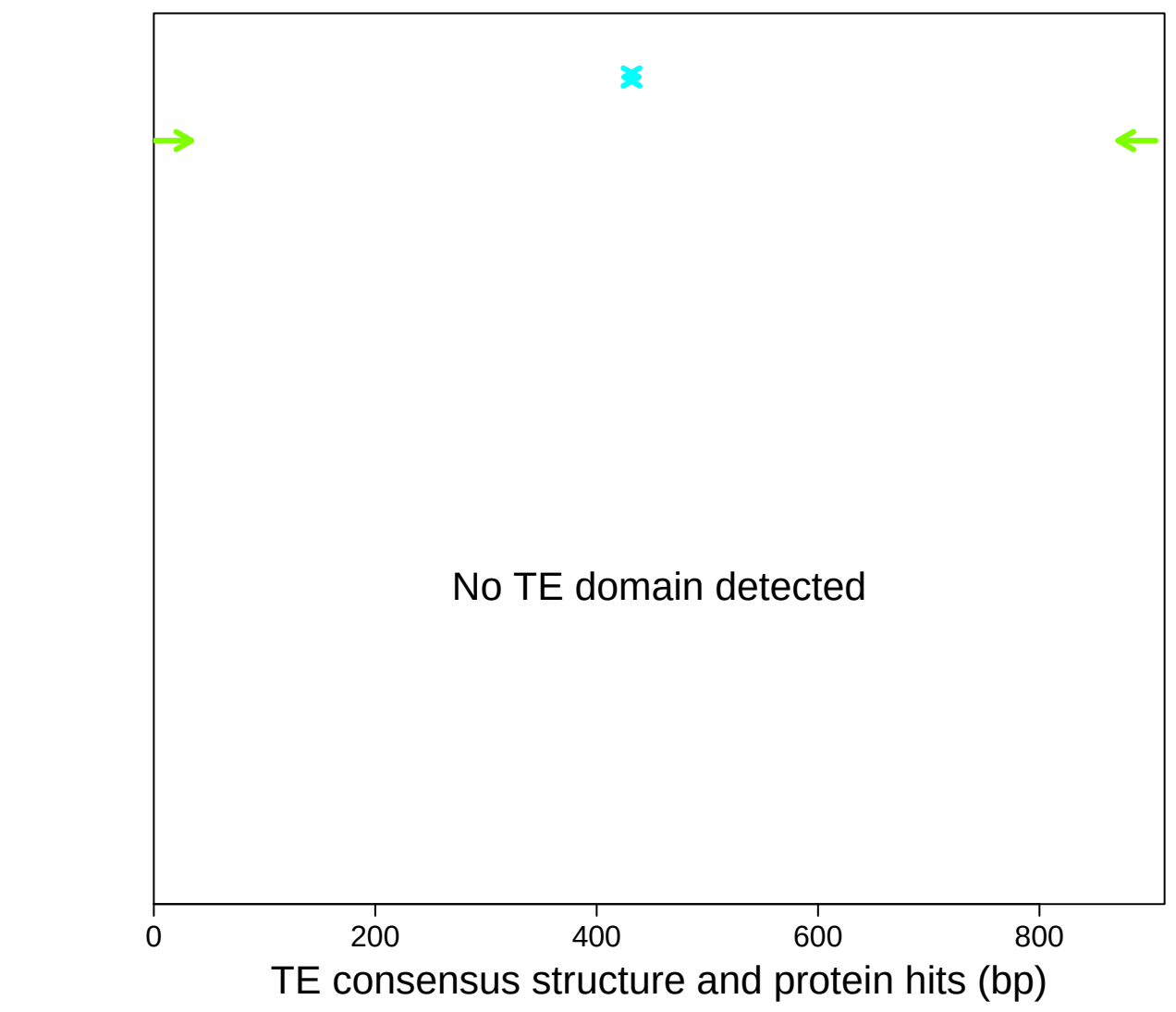
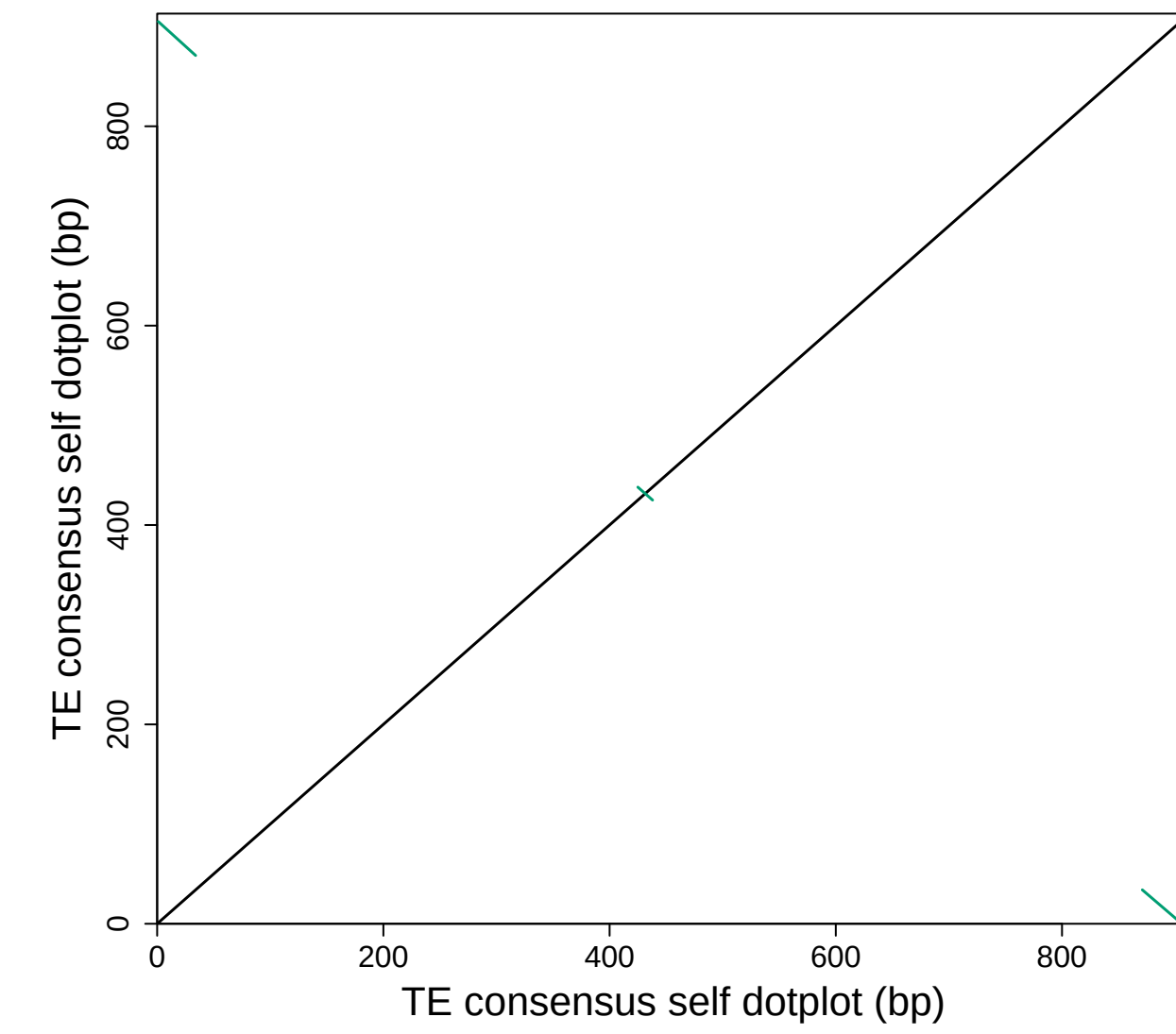
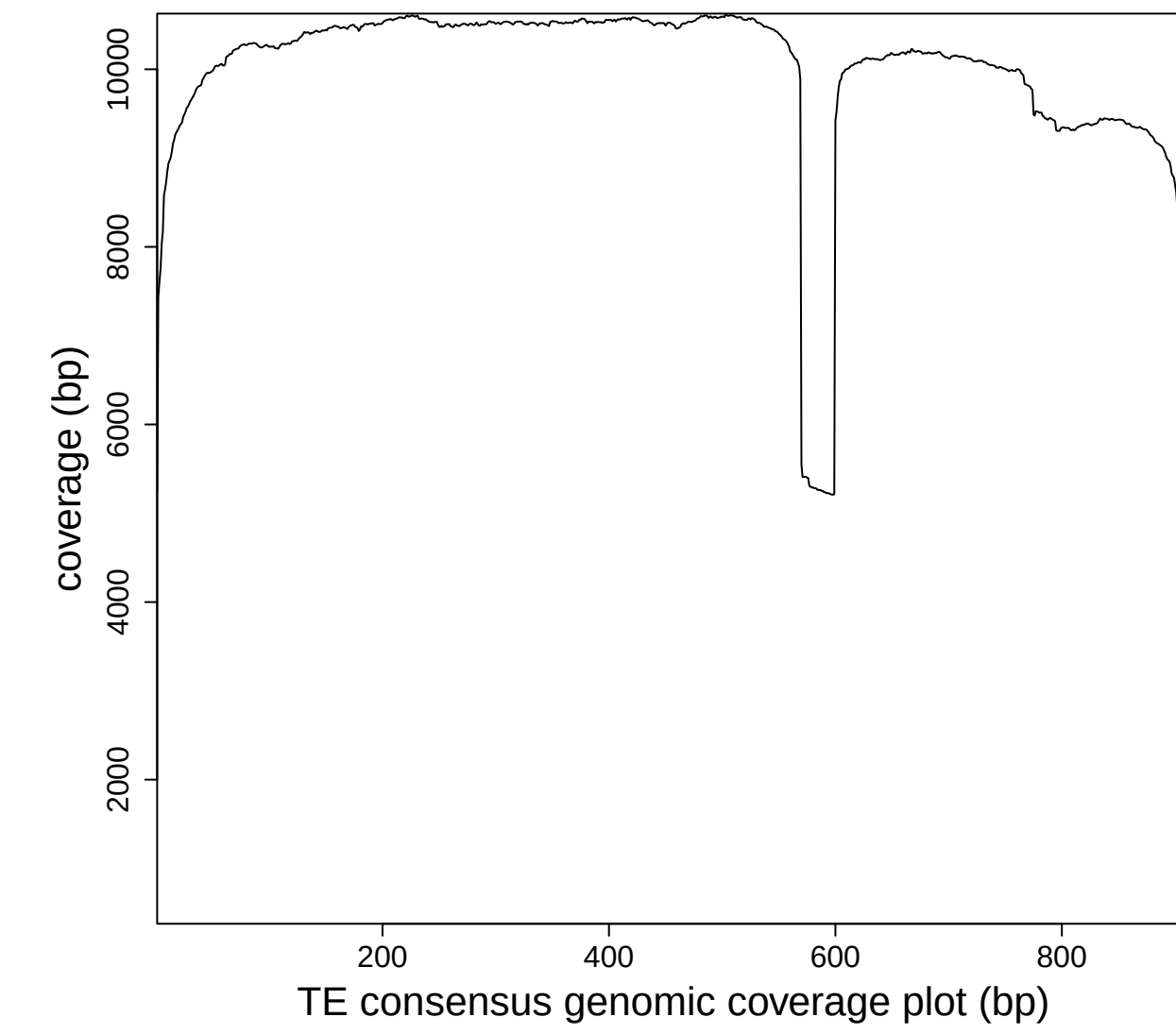
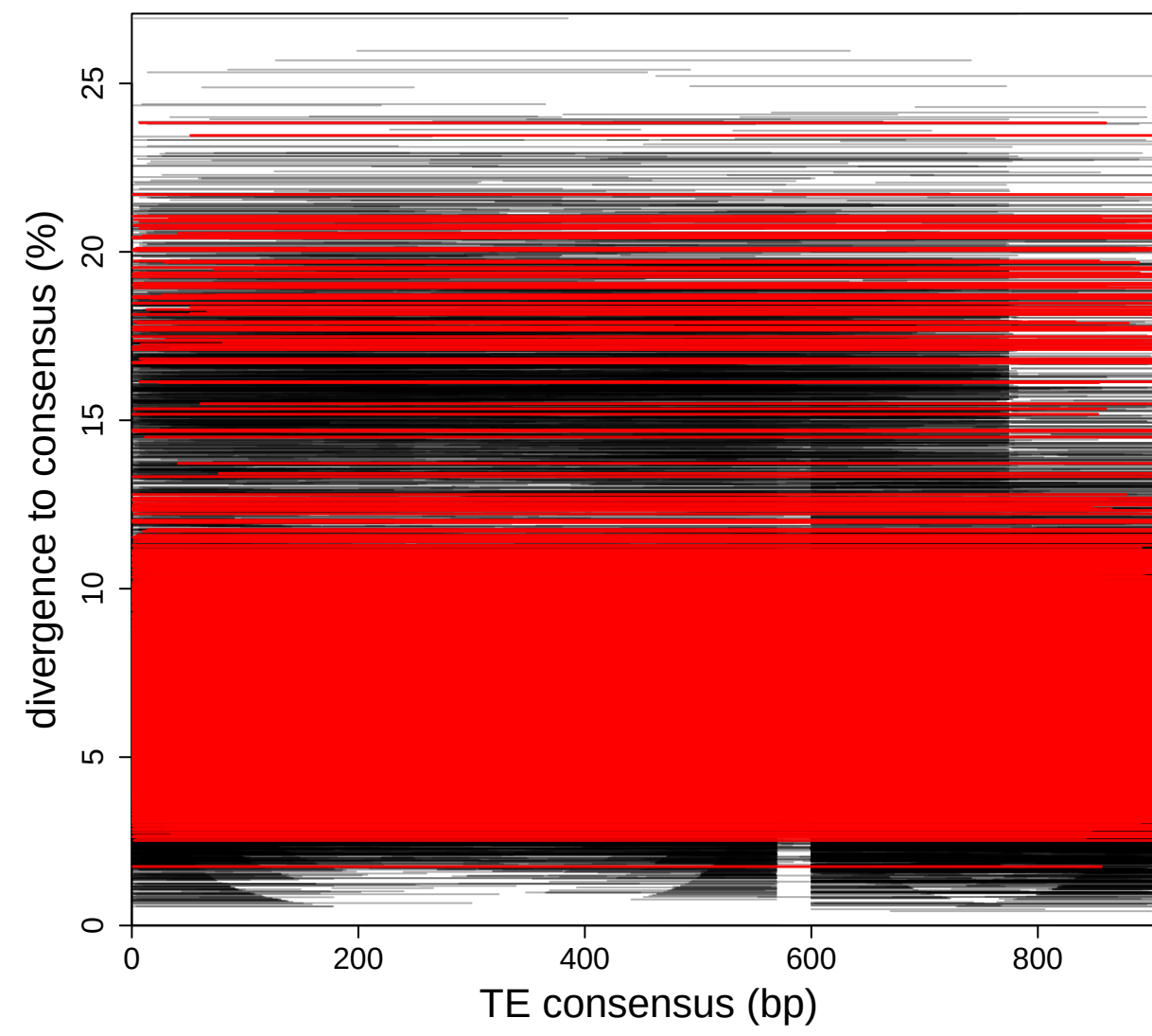
Start crop Point

End crop Point



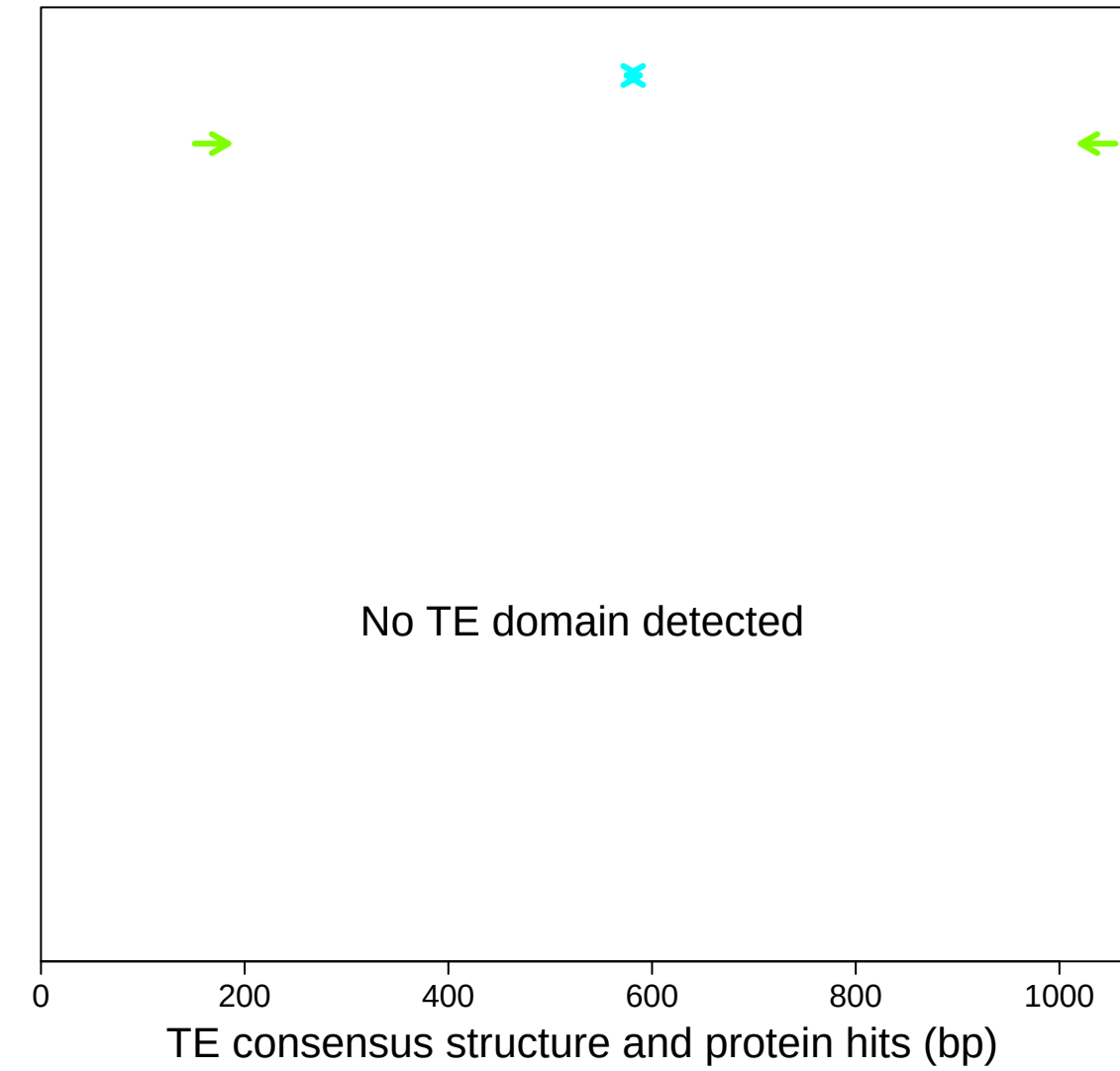
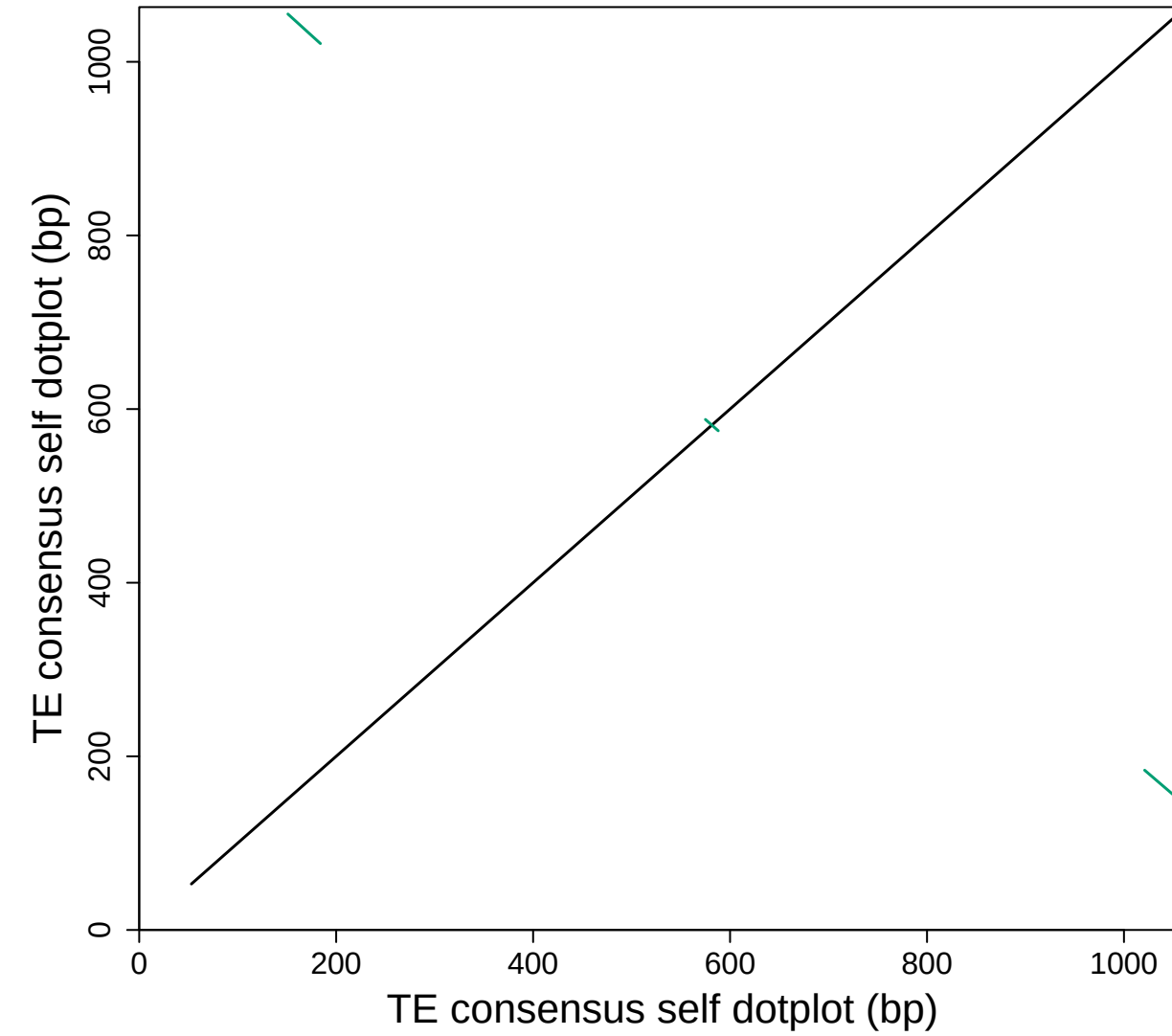
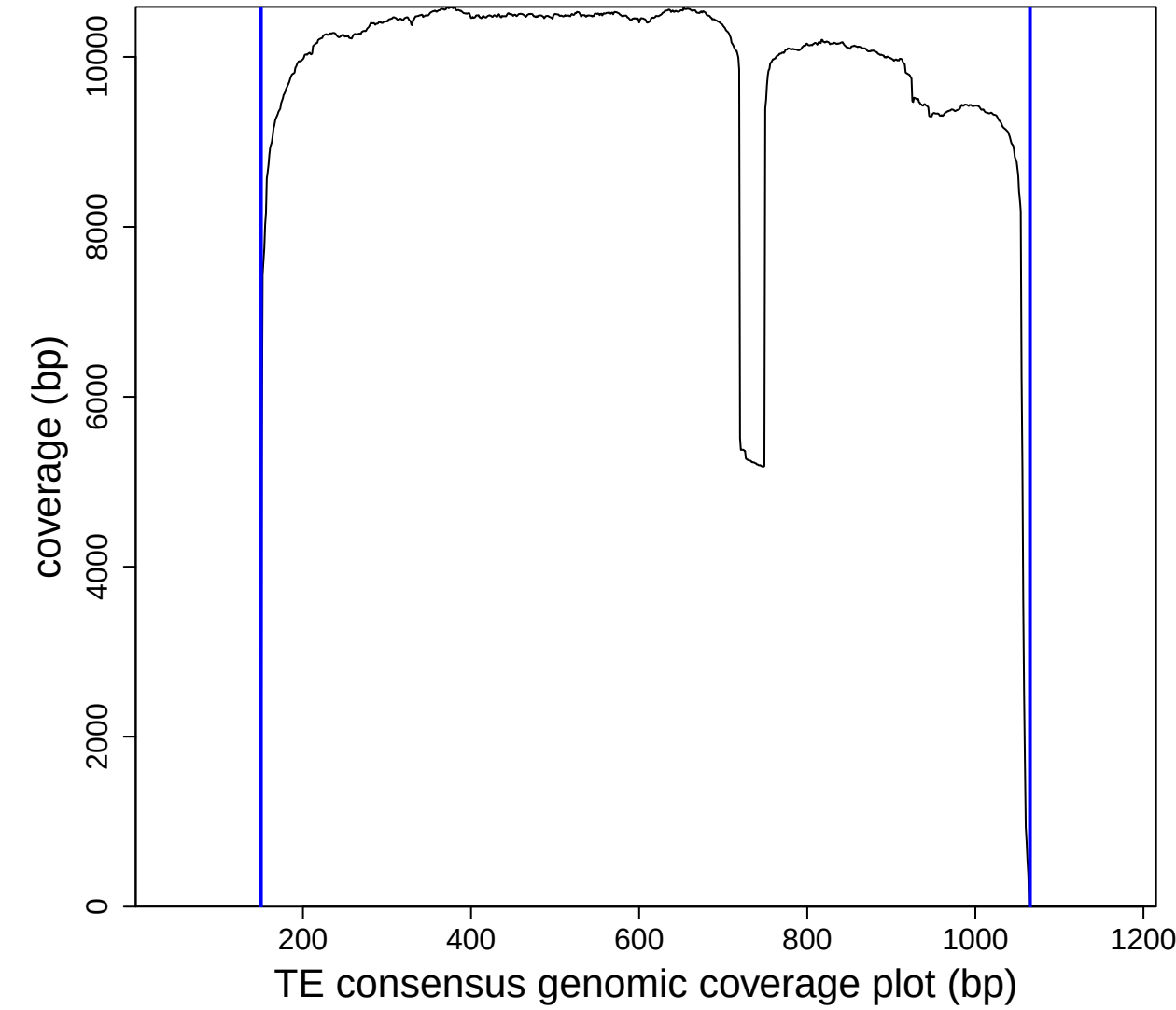
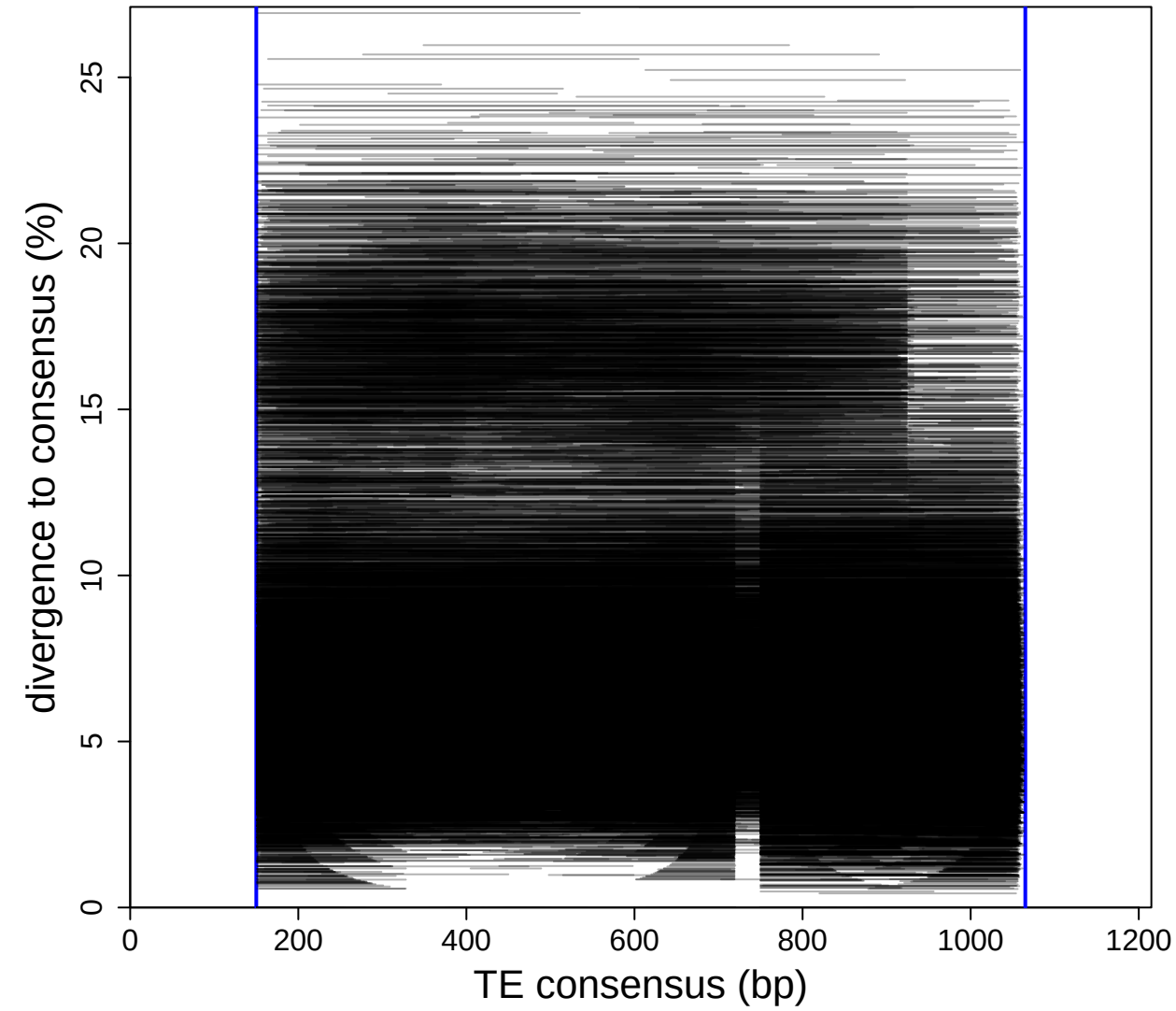
b.bed uf.bed g 1.bed fm 1.bed 0 0 bcln.fa aln.fa cl.fa gs.fa
size: 913bp; fragments: 22153; full length: 2228 (≥ 821.7 bp)

After TEtrimmer 913 bp



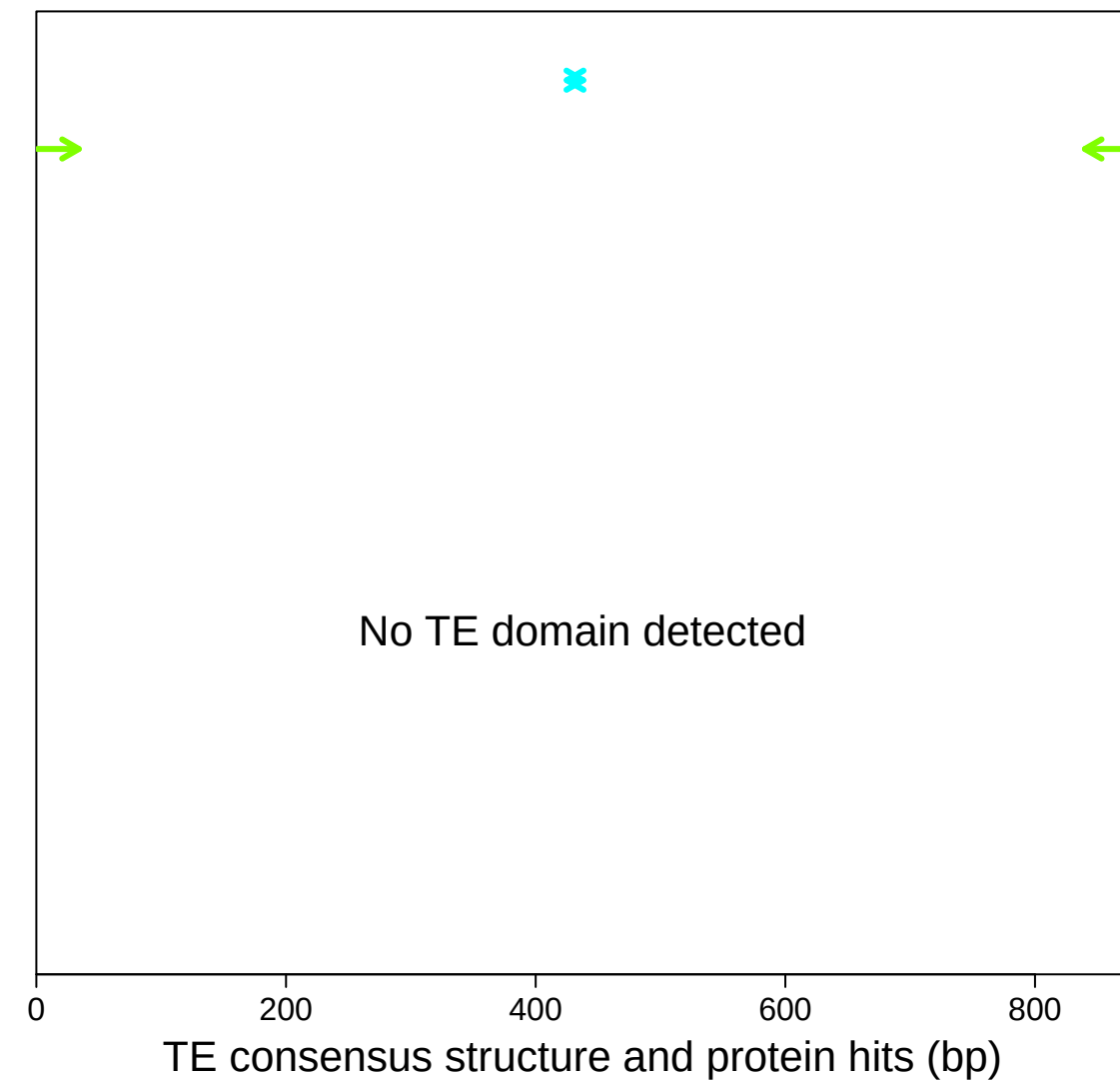
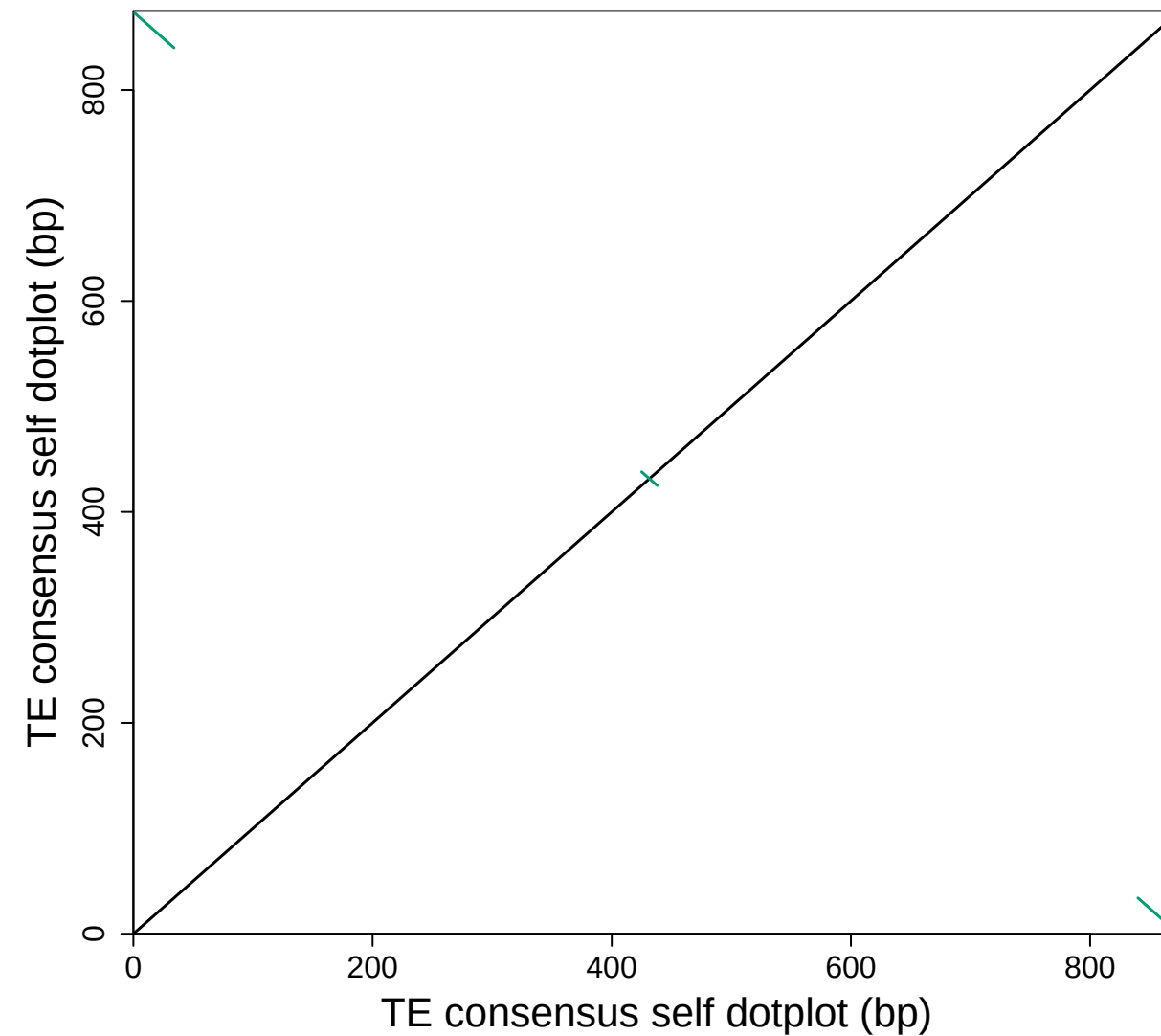
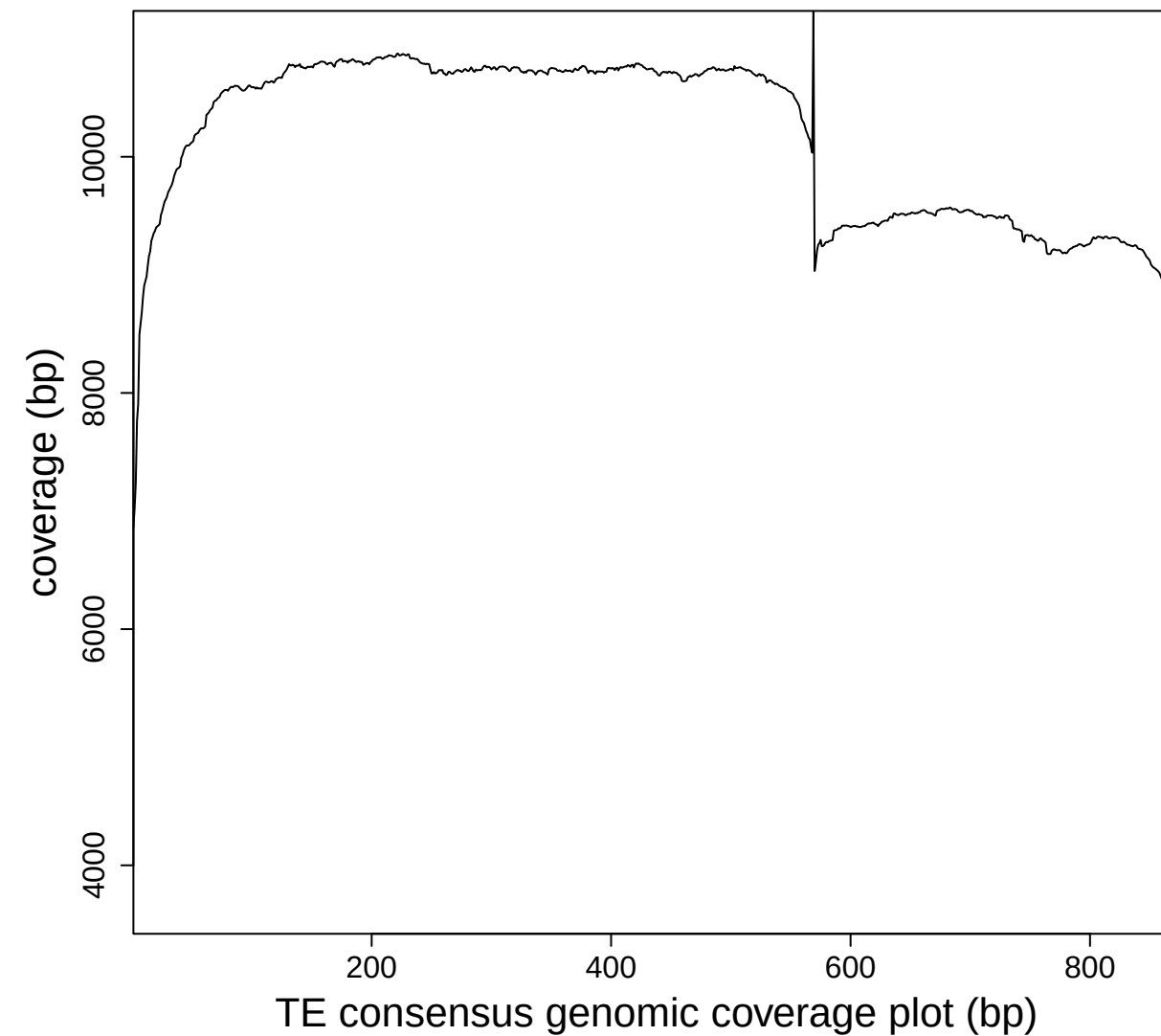
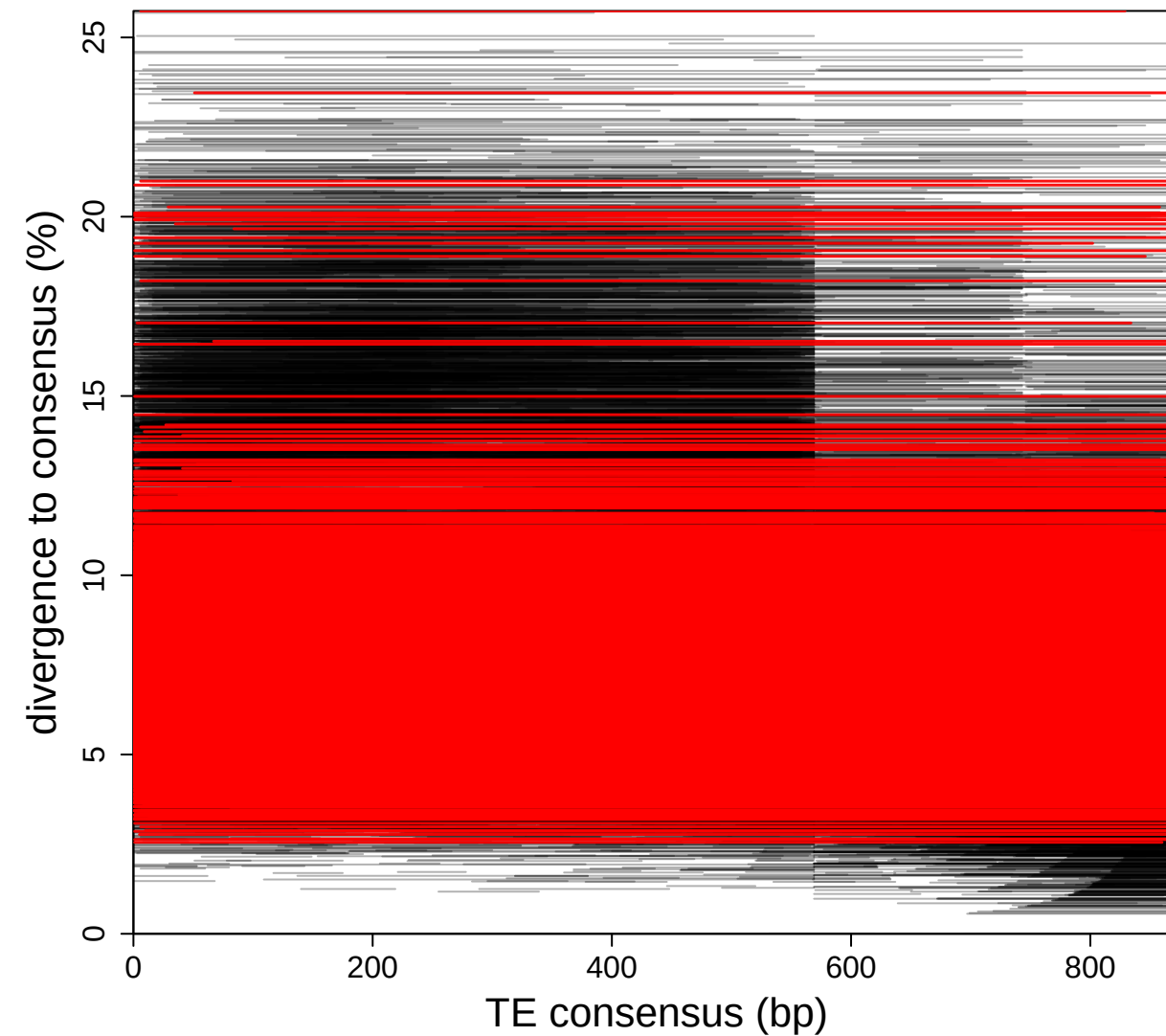
.fasta.b.bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa
size: 1215bp; fragments: 22101; full length: 0 (>=1093.5bp)

After TEtrimmer Extended plot Blue lines are boundaries



TE: rnd_2_family_49
size: 875bp; fragments: 19570; full length: 4008 (≥ 787.5 bp)

Before TEtrimmer 875 bp



Dotplot

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

