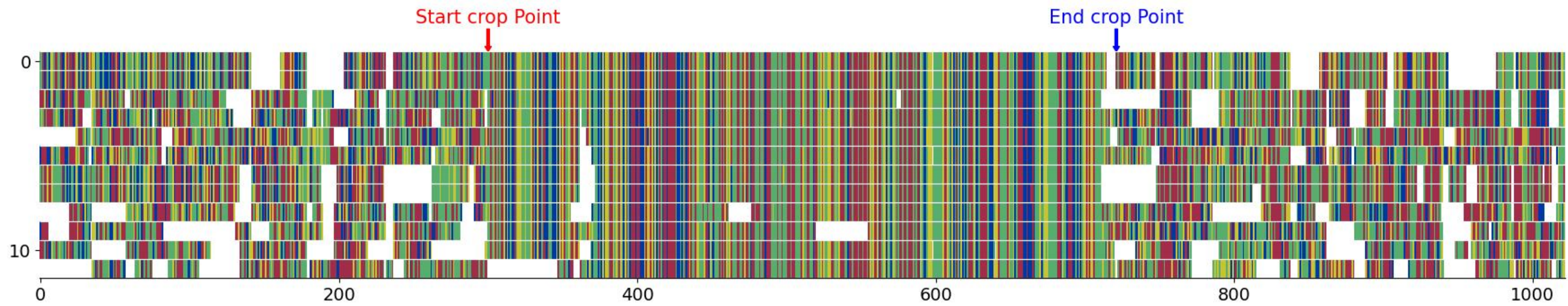
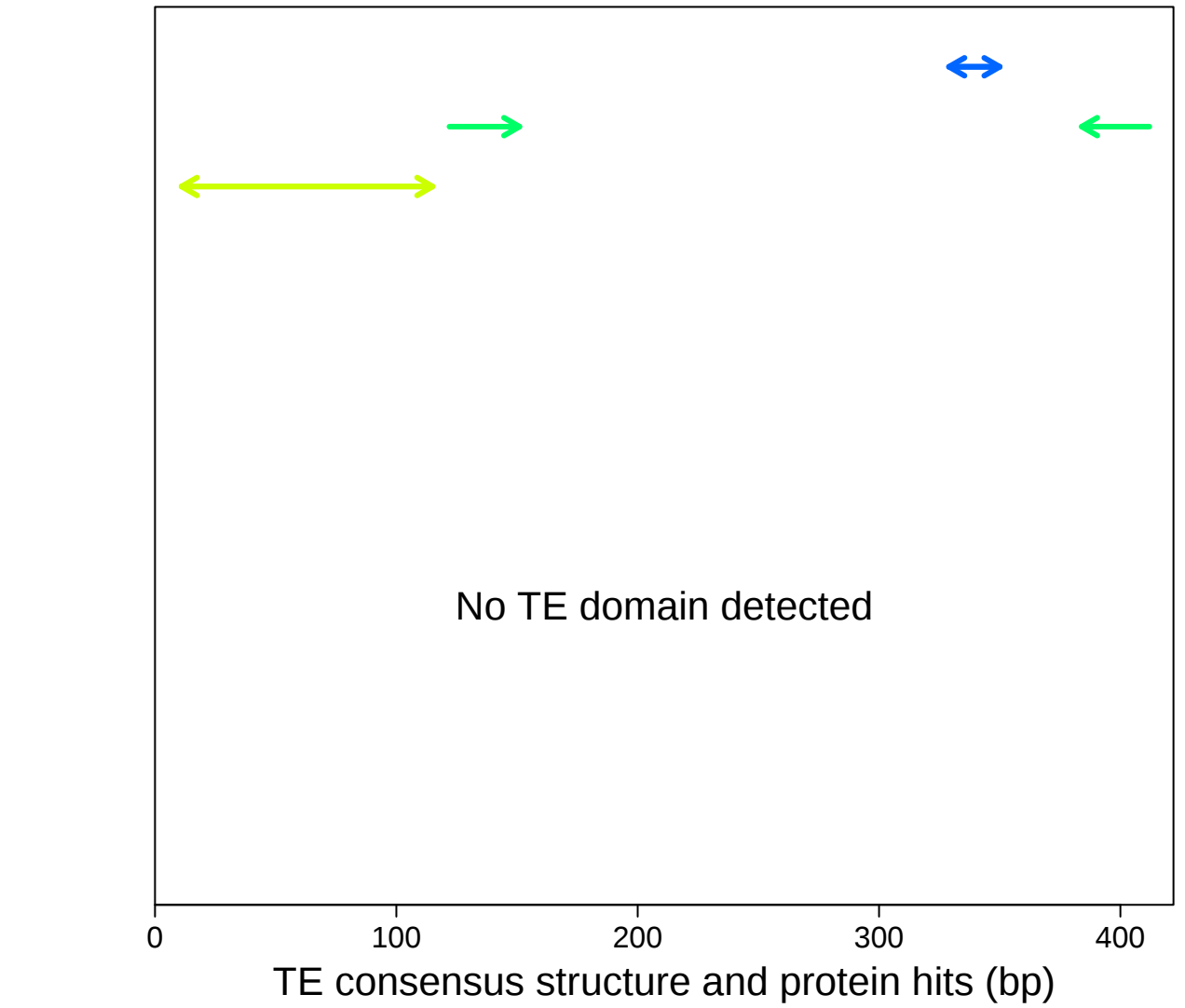
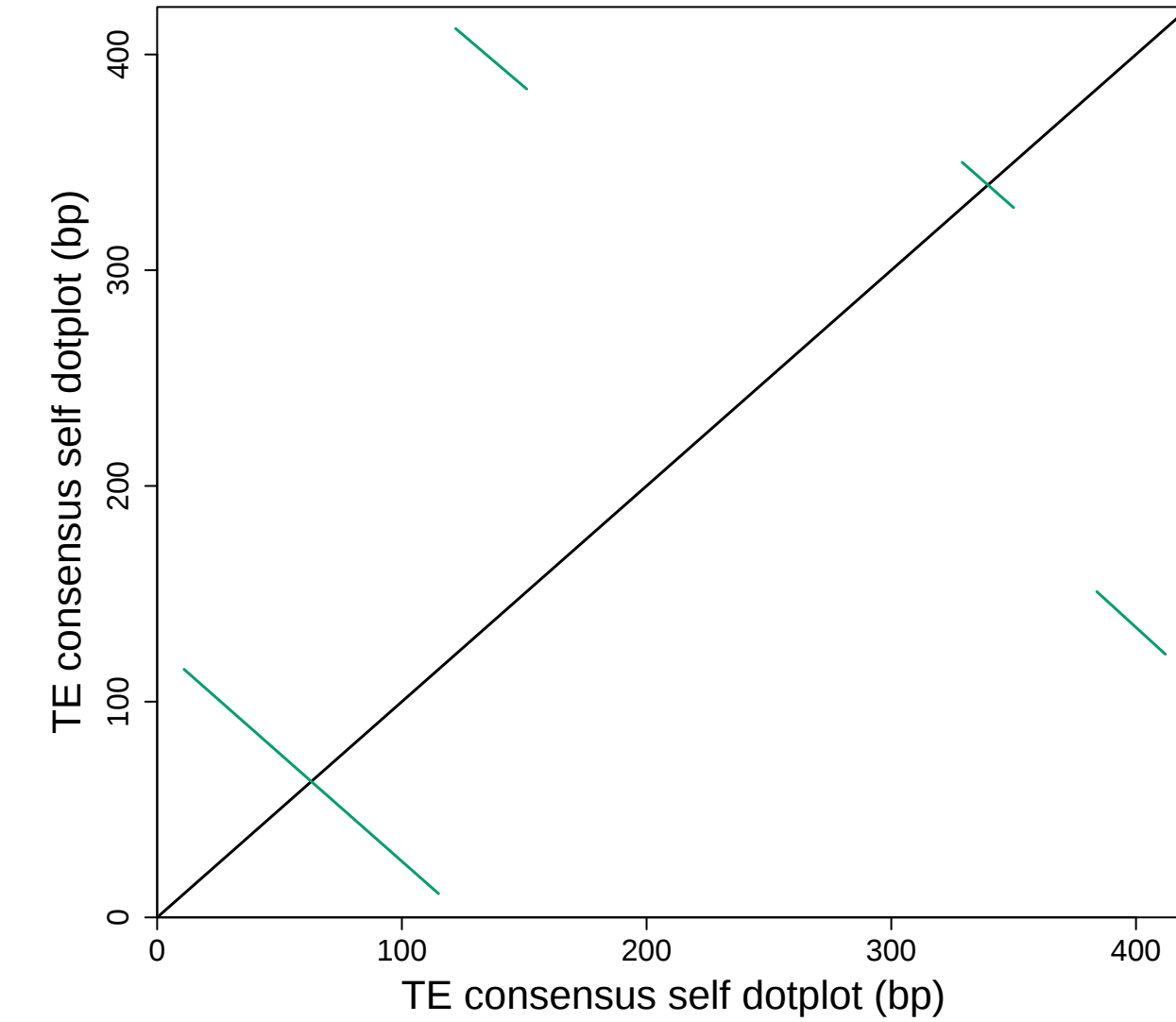
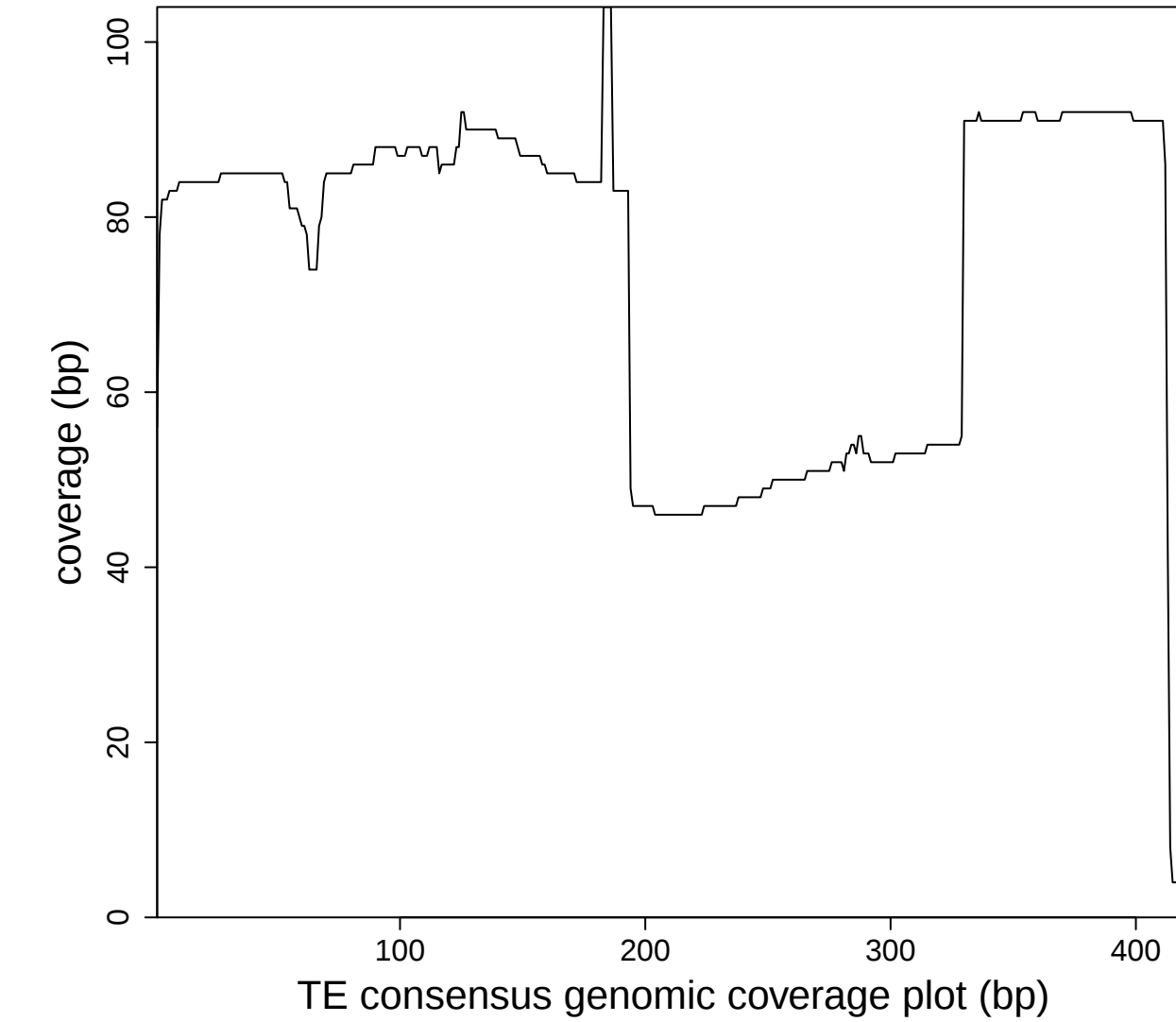
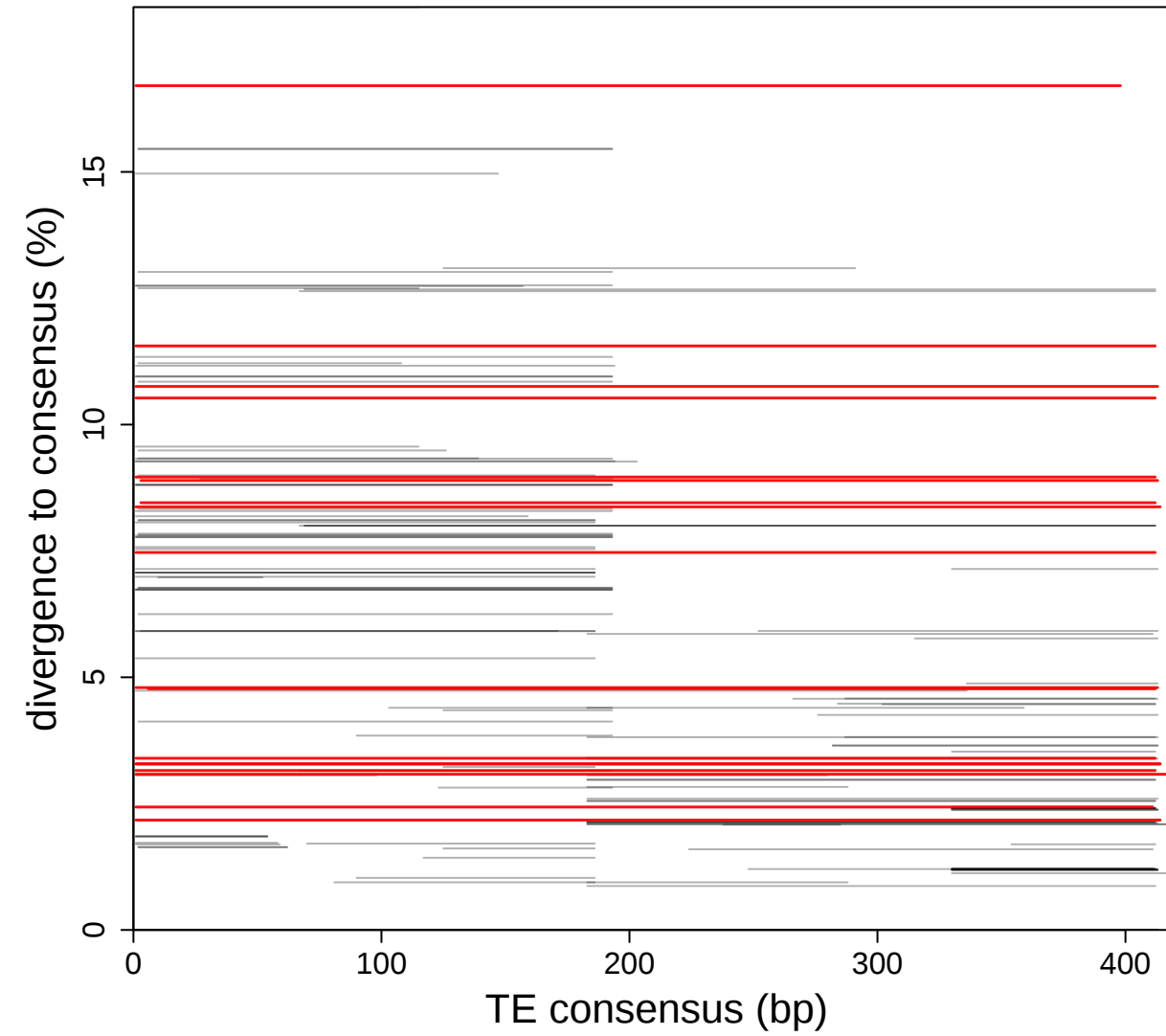


MSA length =



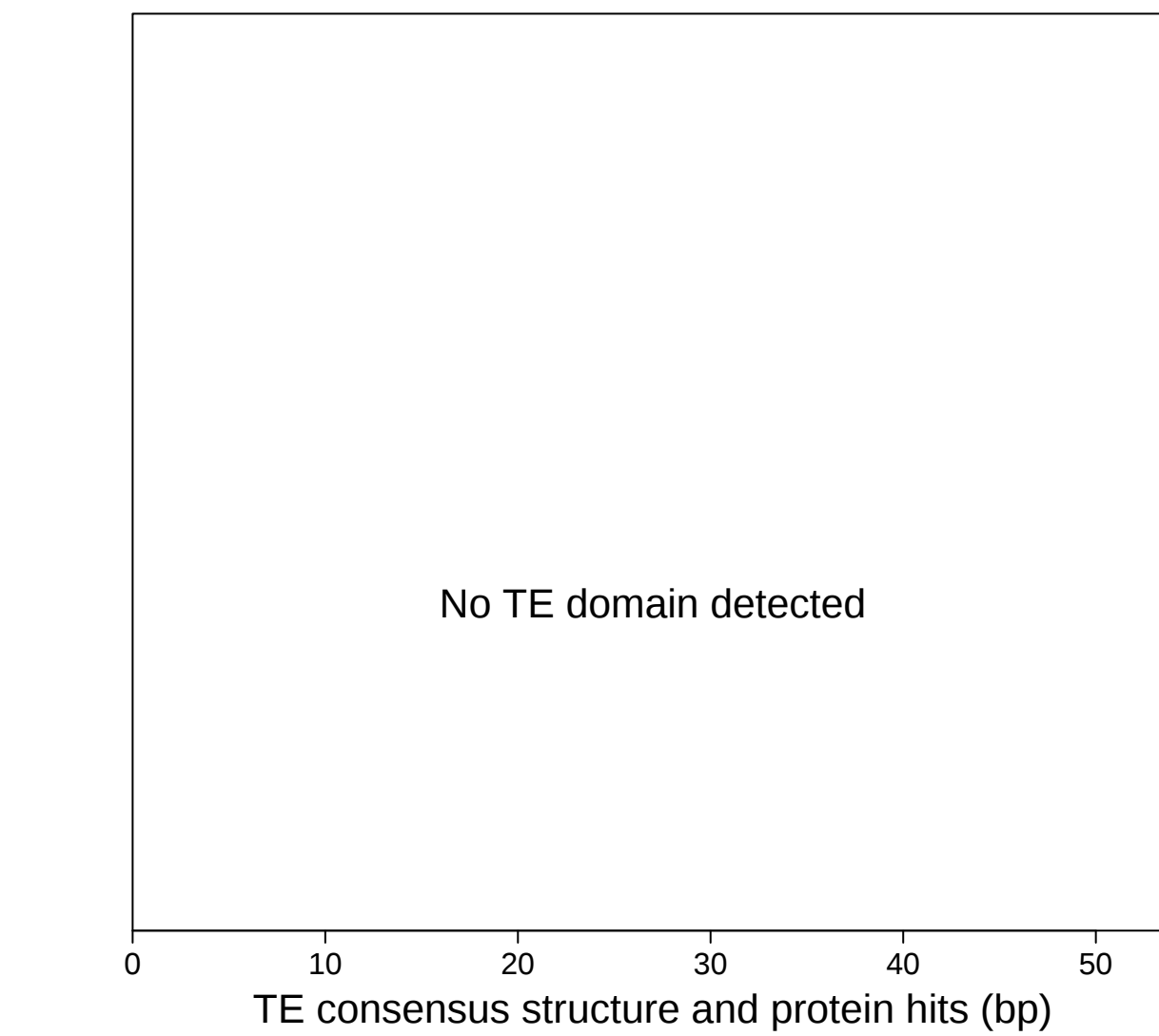
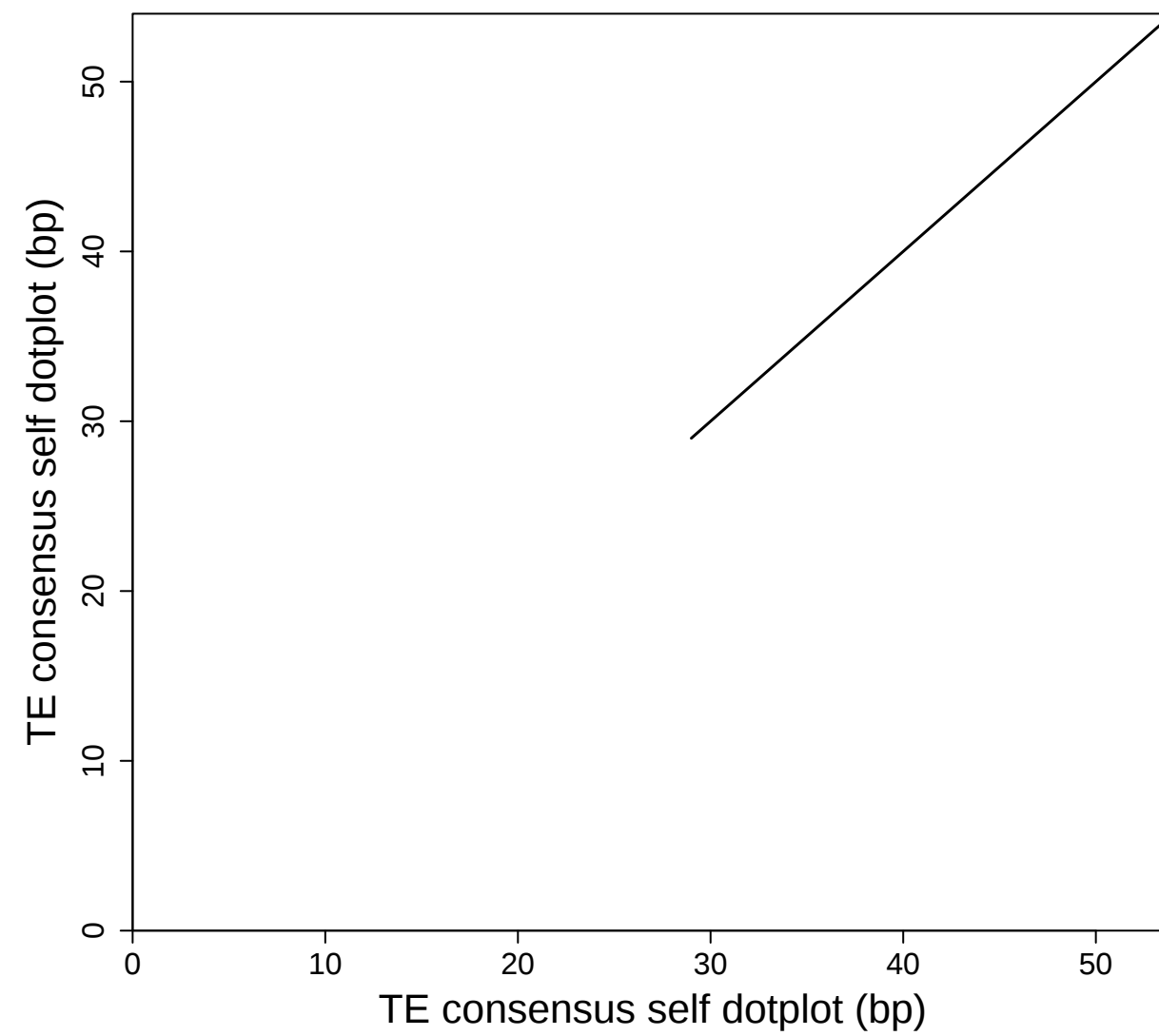
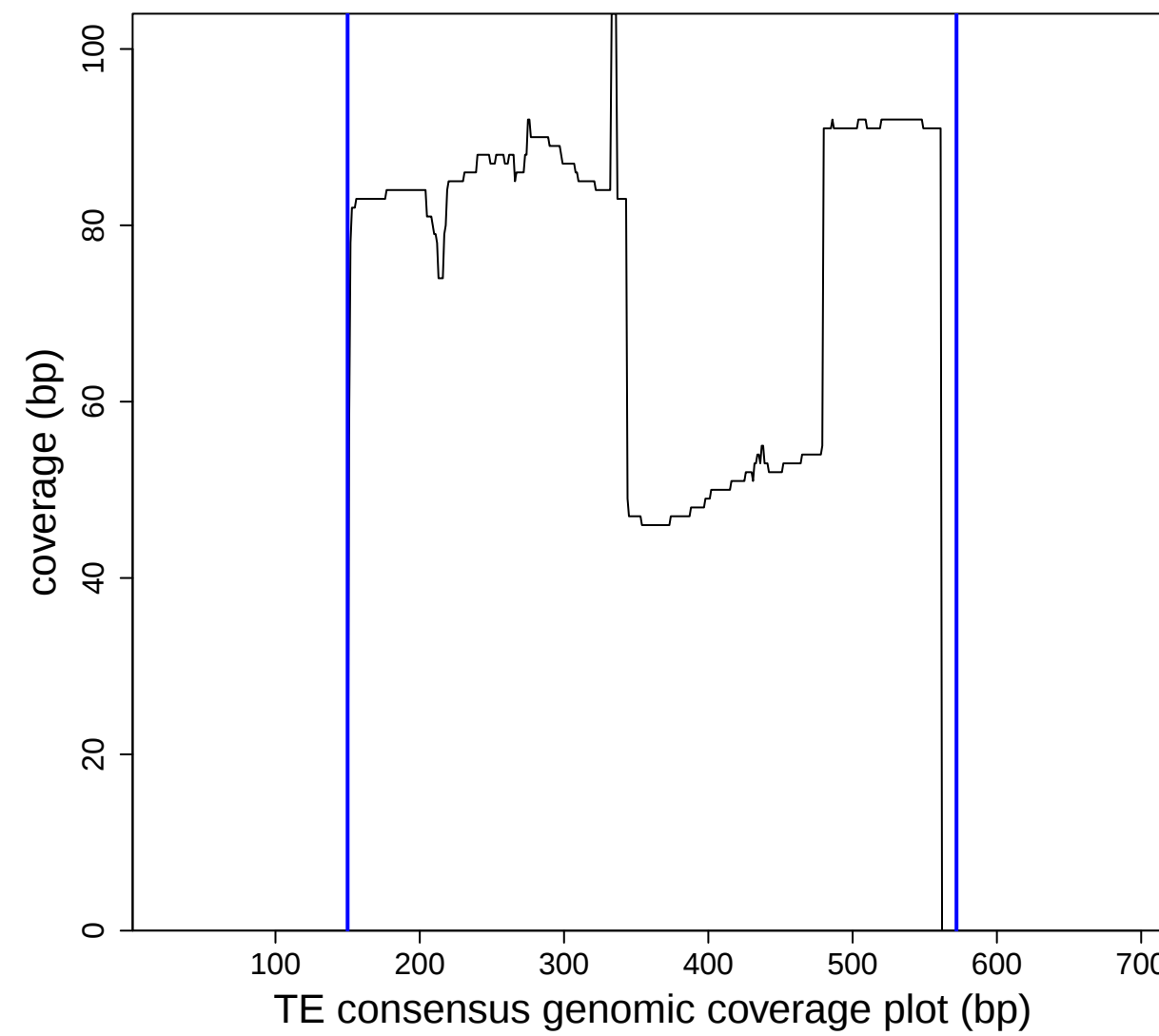
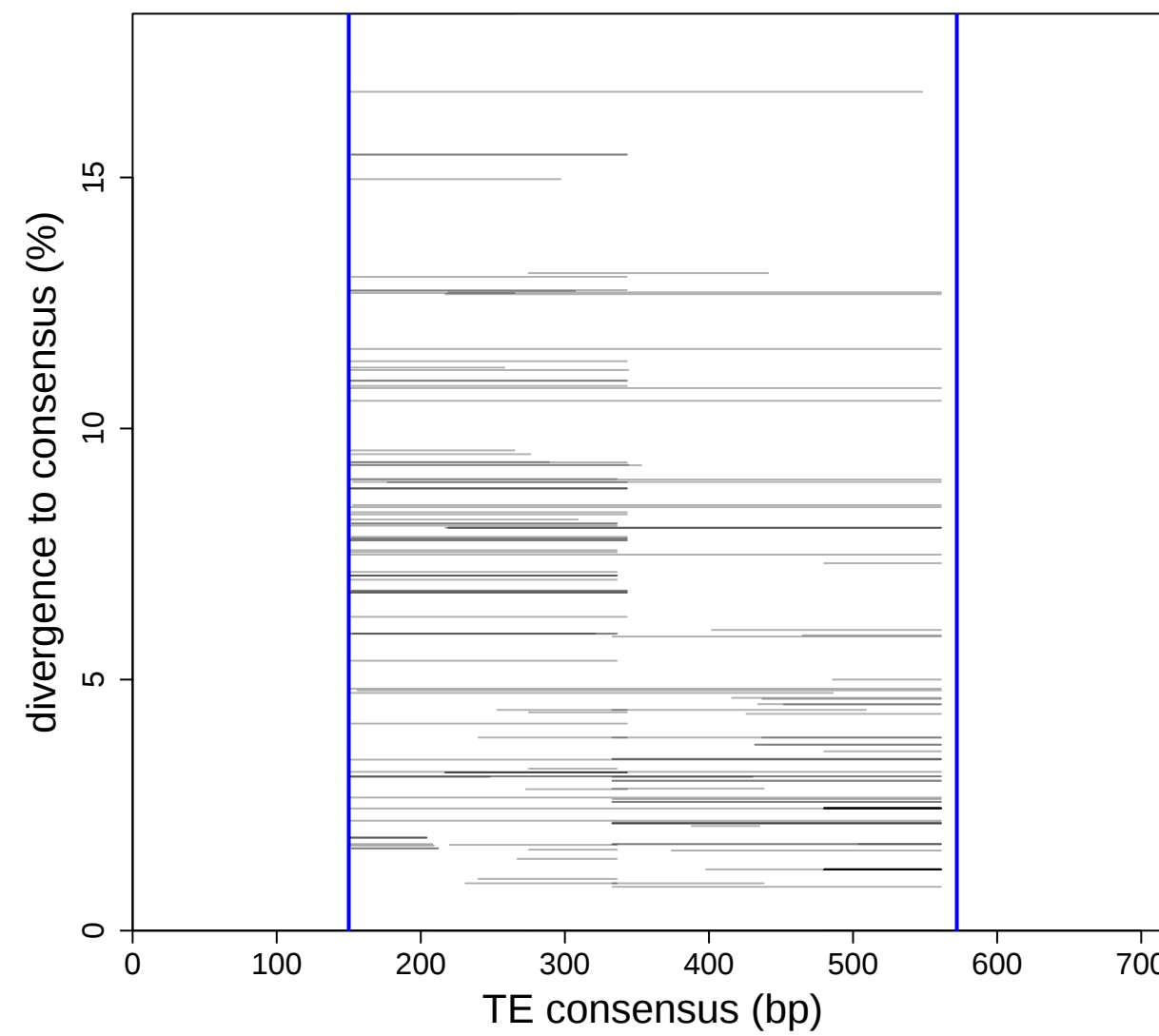
a.b.bed_uf.bed_g_3.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.f
size: 422bp; fragments: 181; full length: 18 (≥ 379.8 bp)

After TEtrimmer 422 bp

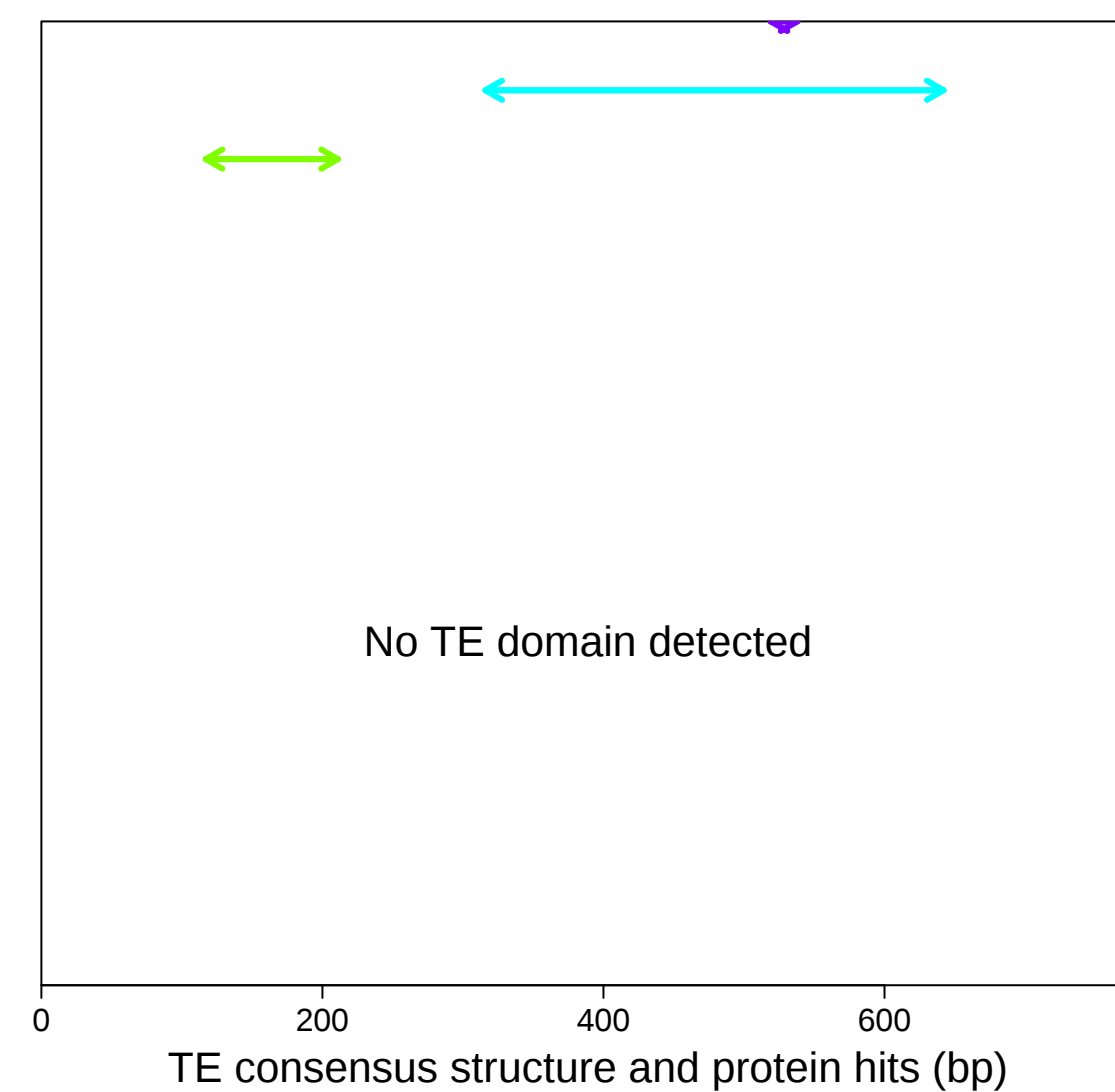
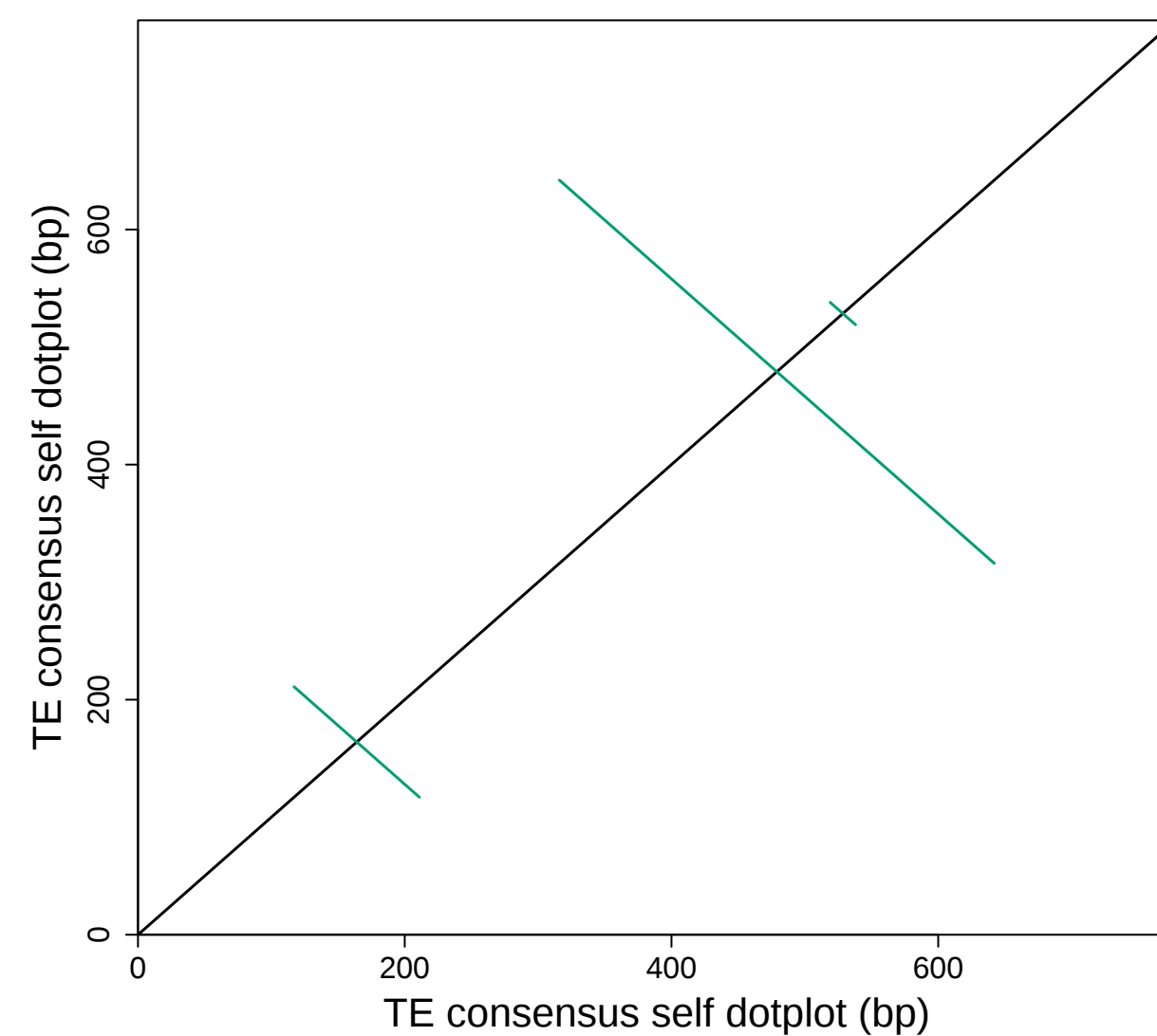
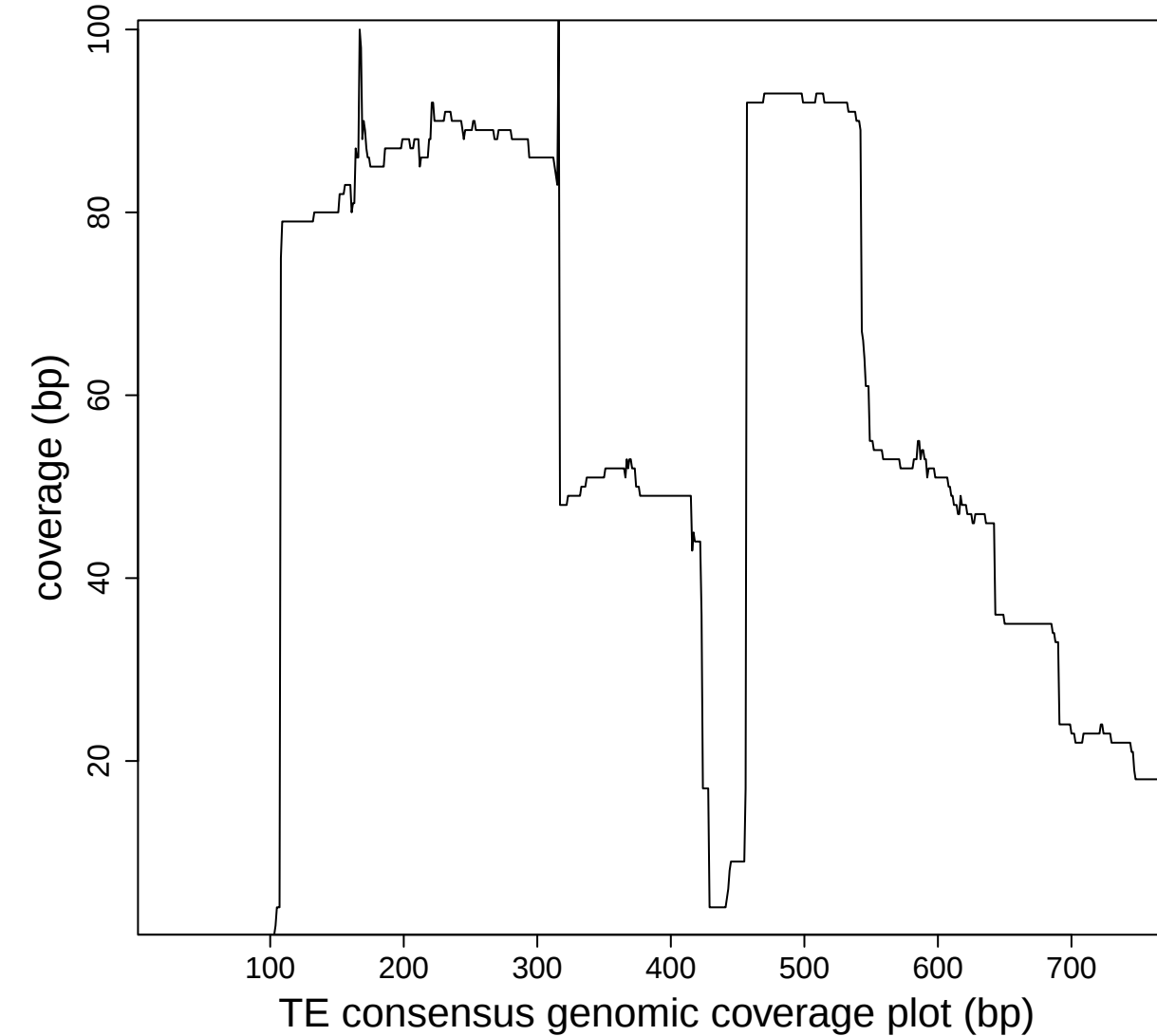
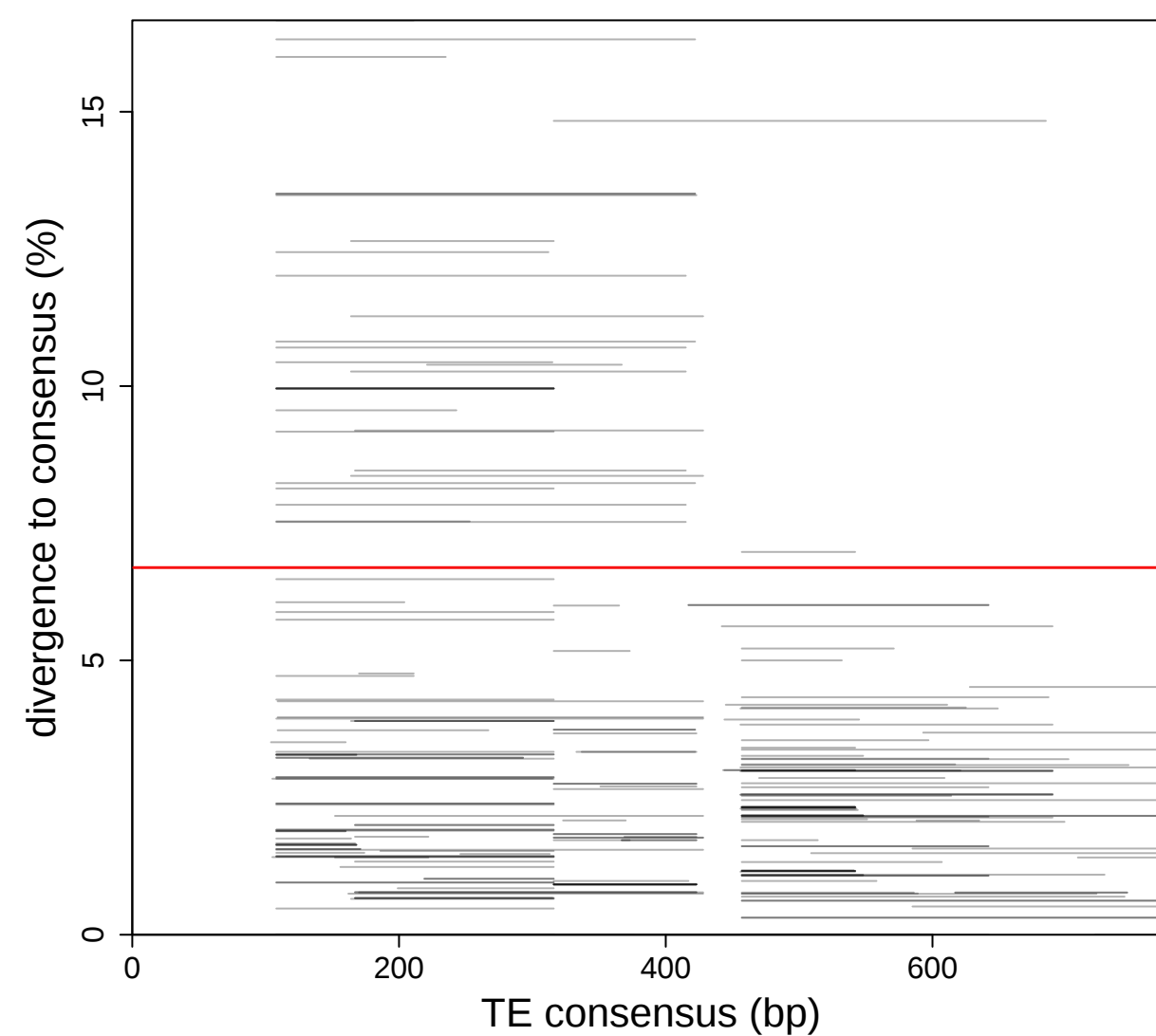


0.fasta.b.bed_uf.bed_g_3.bed_fm_1.bed_0_0_bcIn.fa_aln.fa_cl.f
size: 722bp; fragments: 180; full length: 0 (>=649.8bp)

After TEtrimmer Extended plot Blue lines are boundaries



TE: rnd_5_family_1850
size: 778bp; fragments: 251; full length: 1 (≥ 700.2 bp)



Dotplot

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

