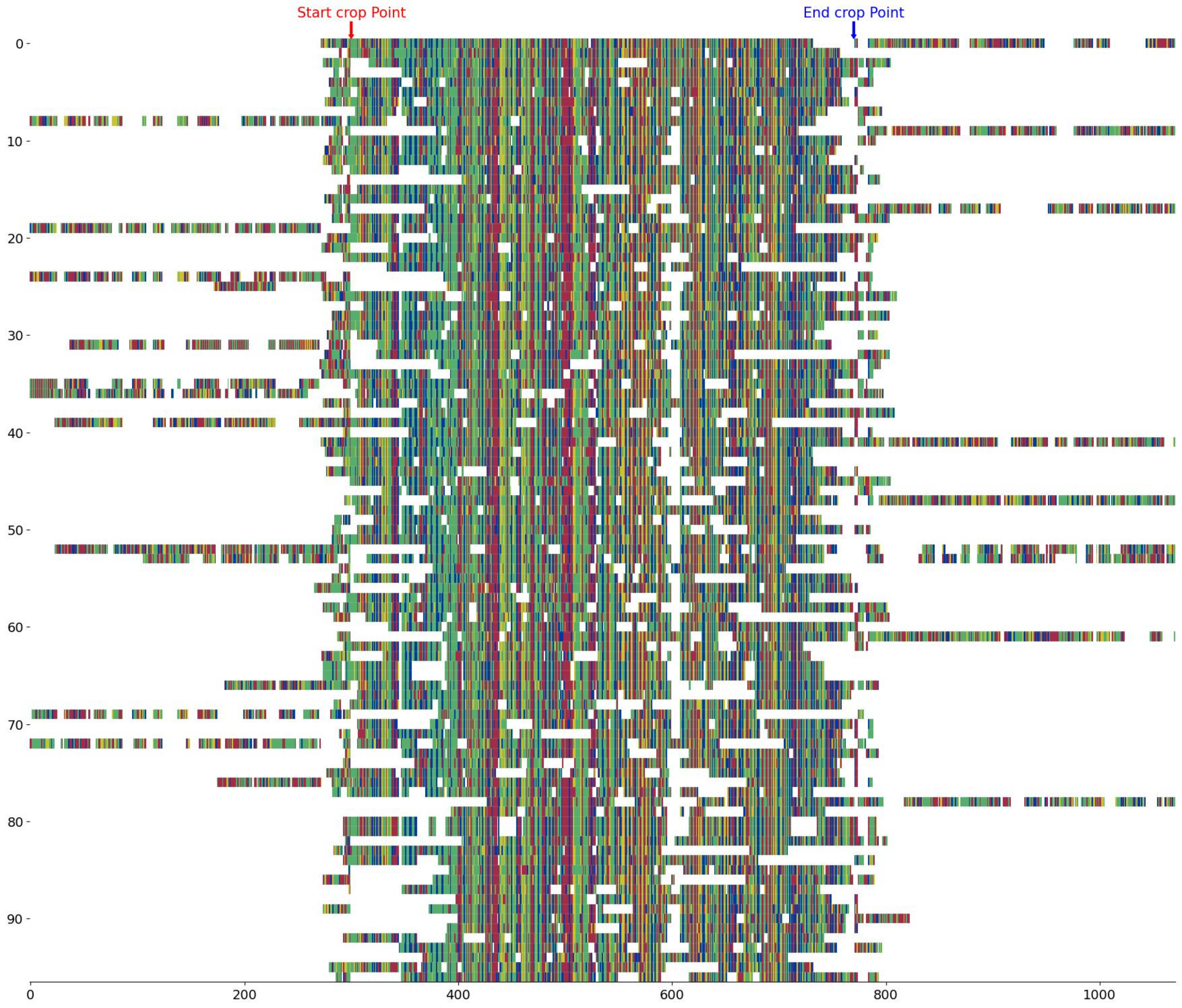


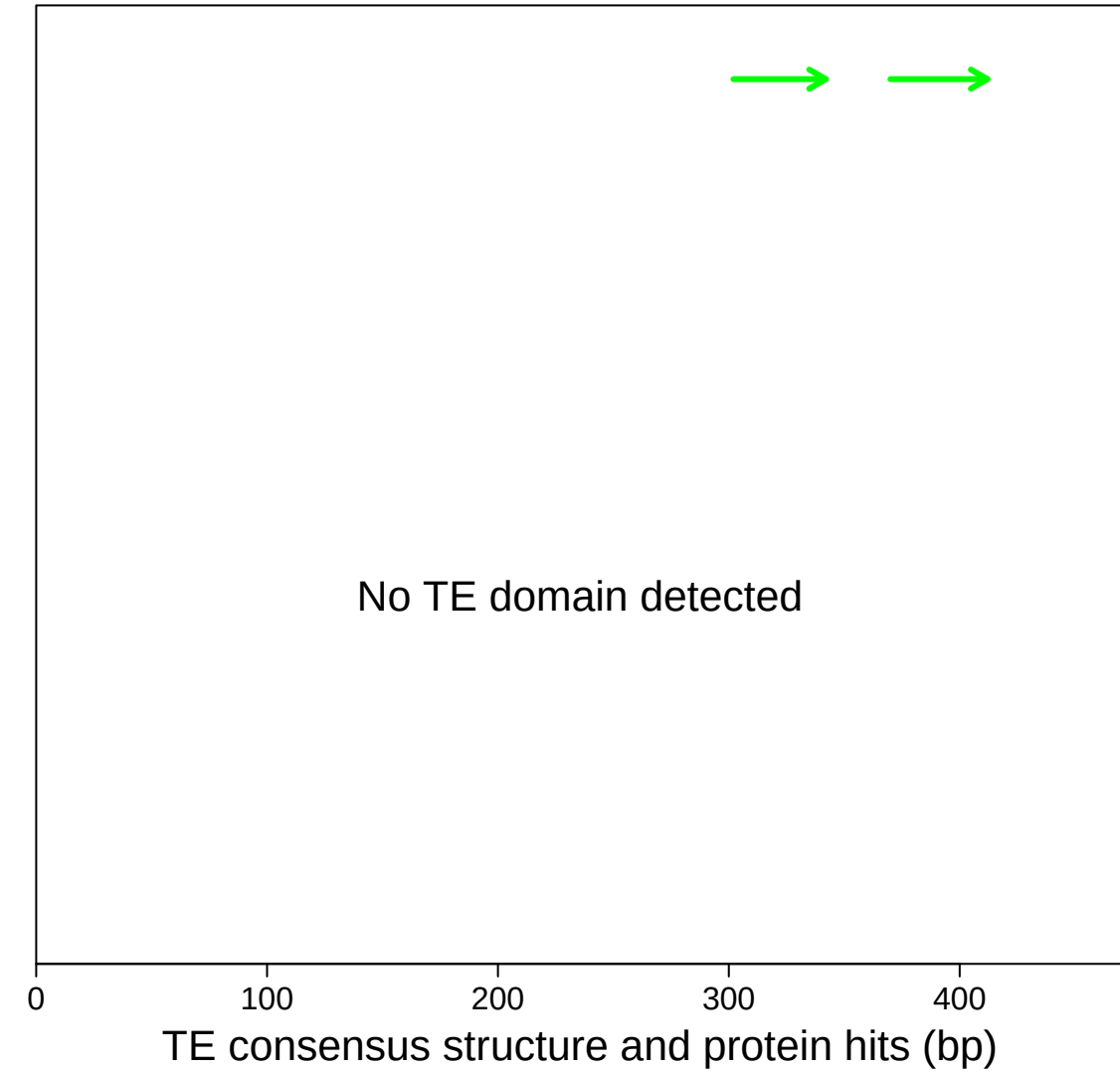
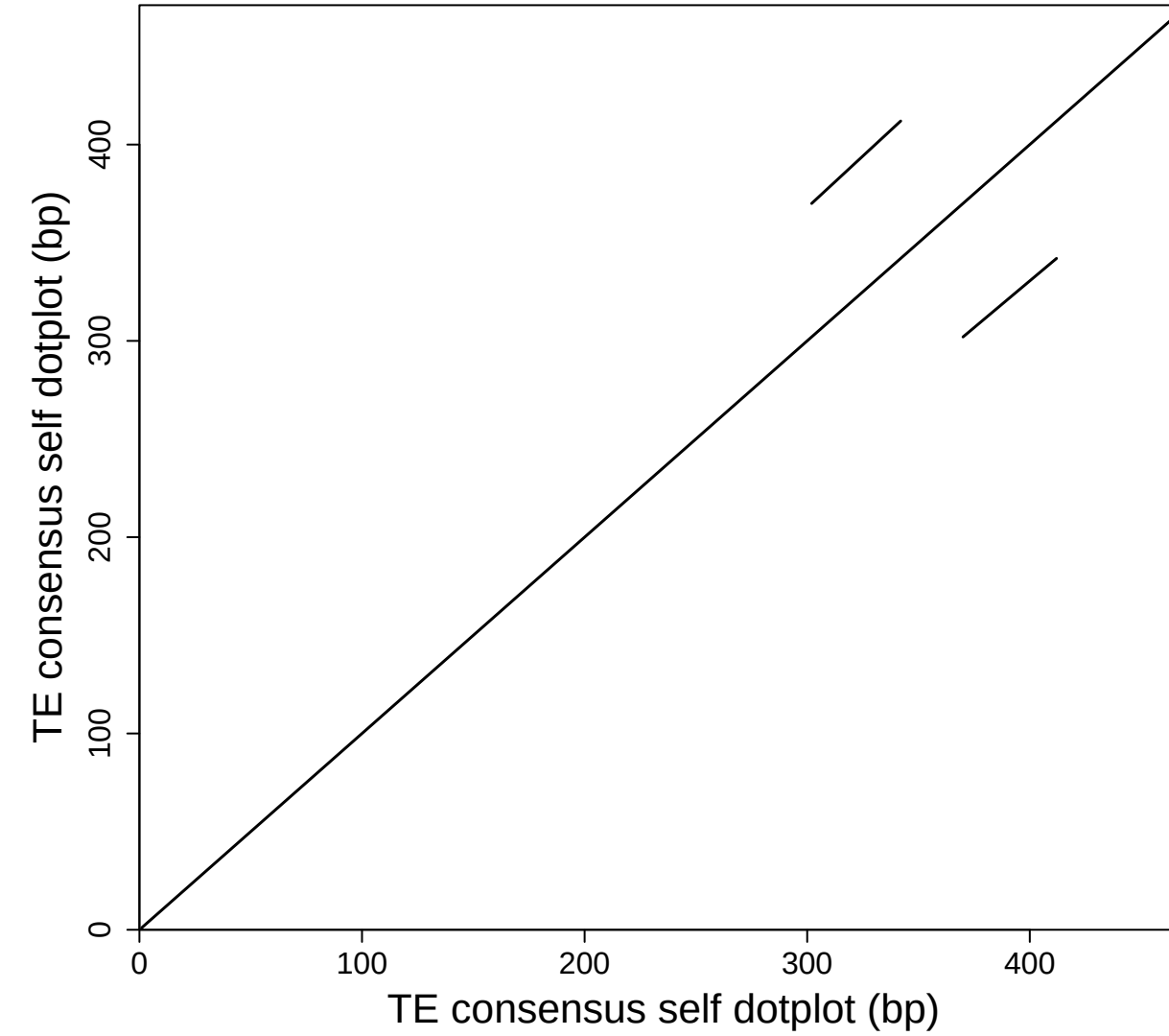
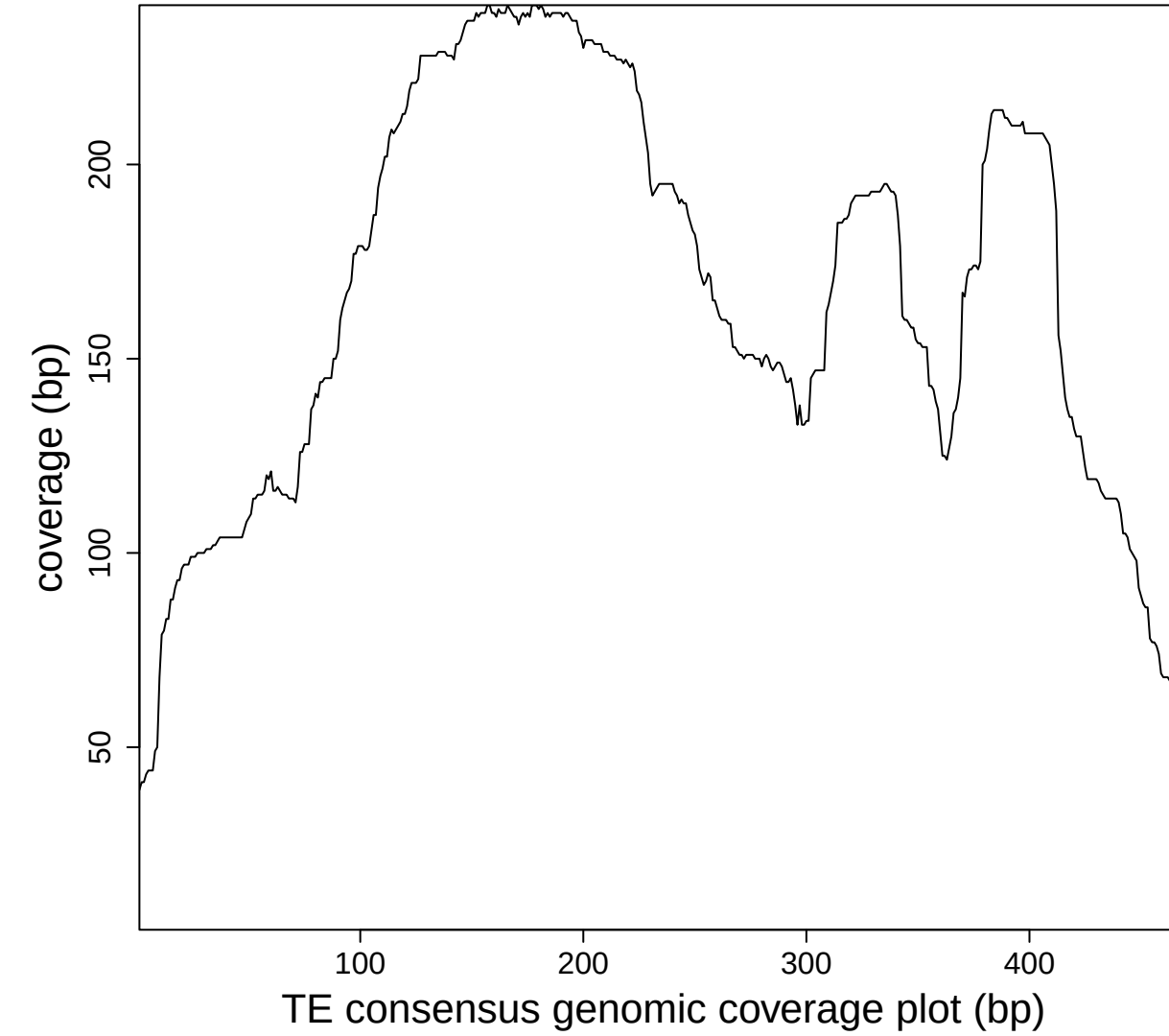
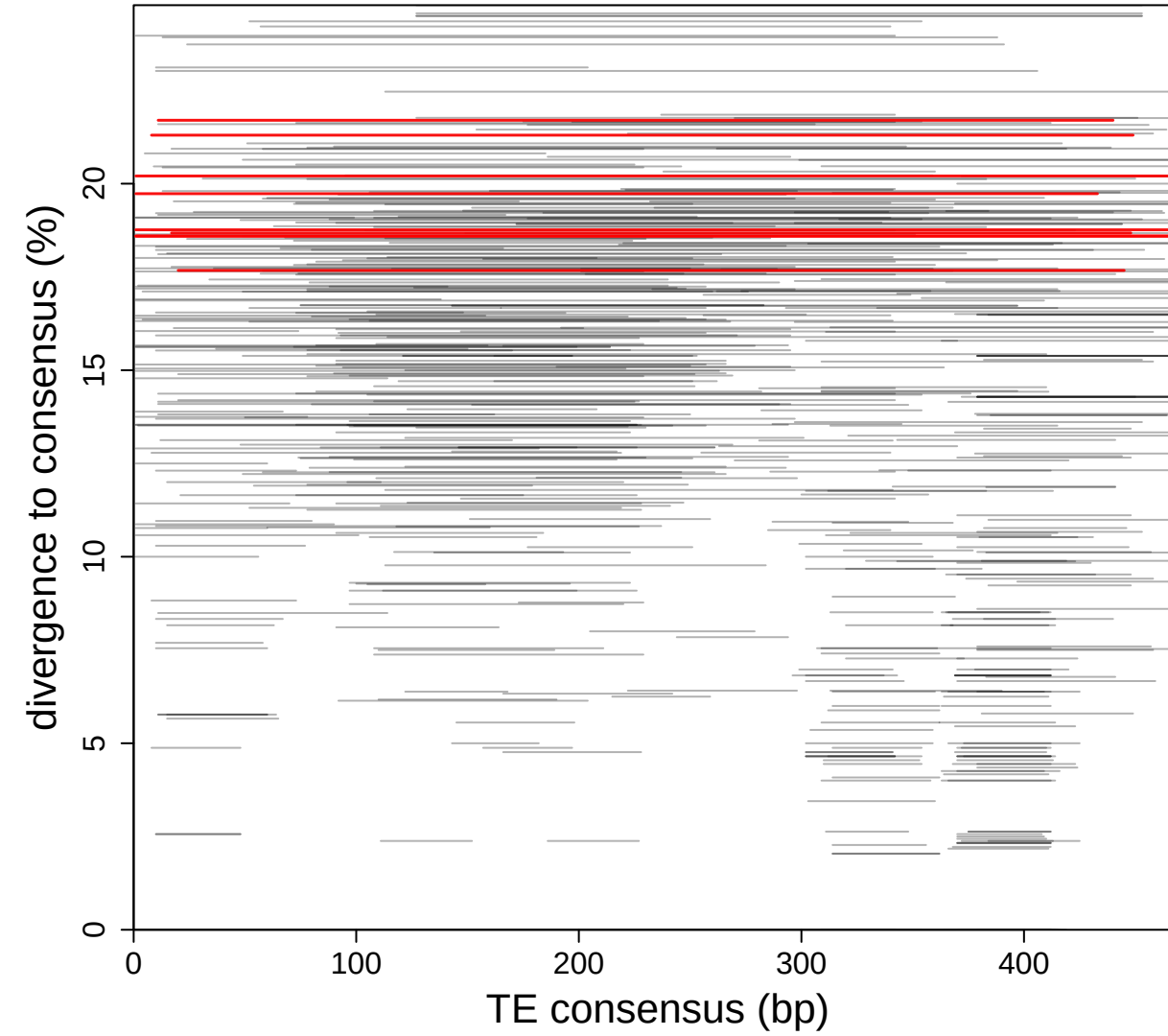
MSA length = 471



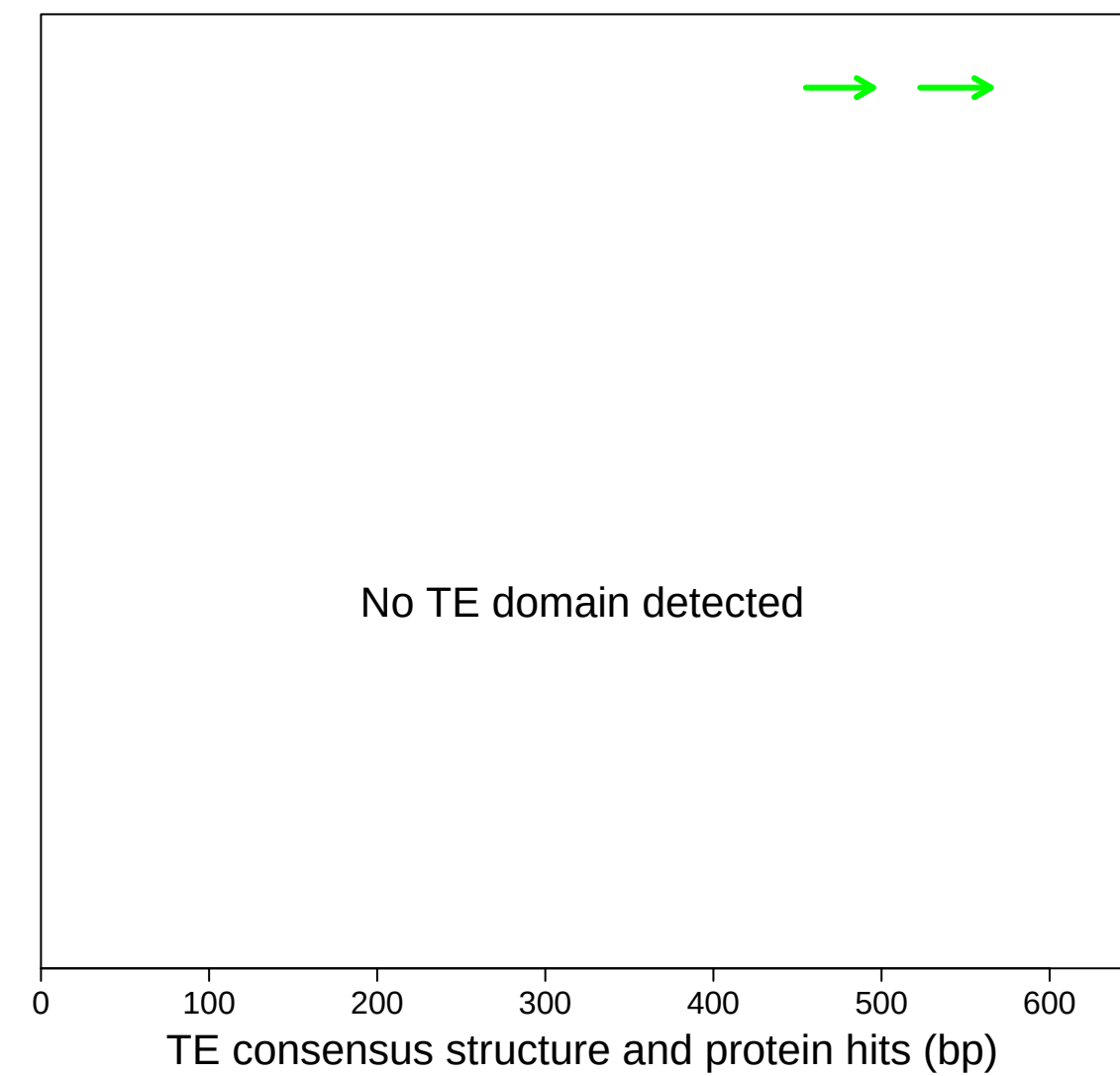
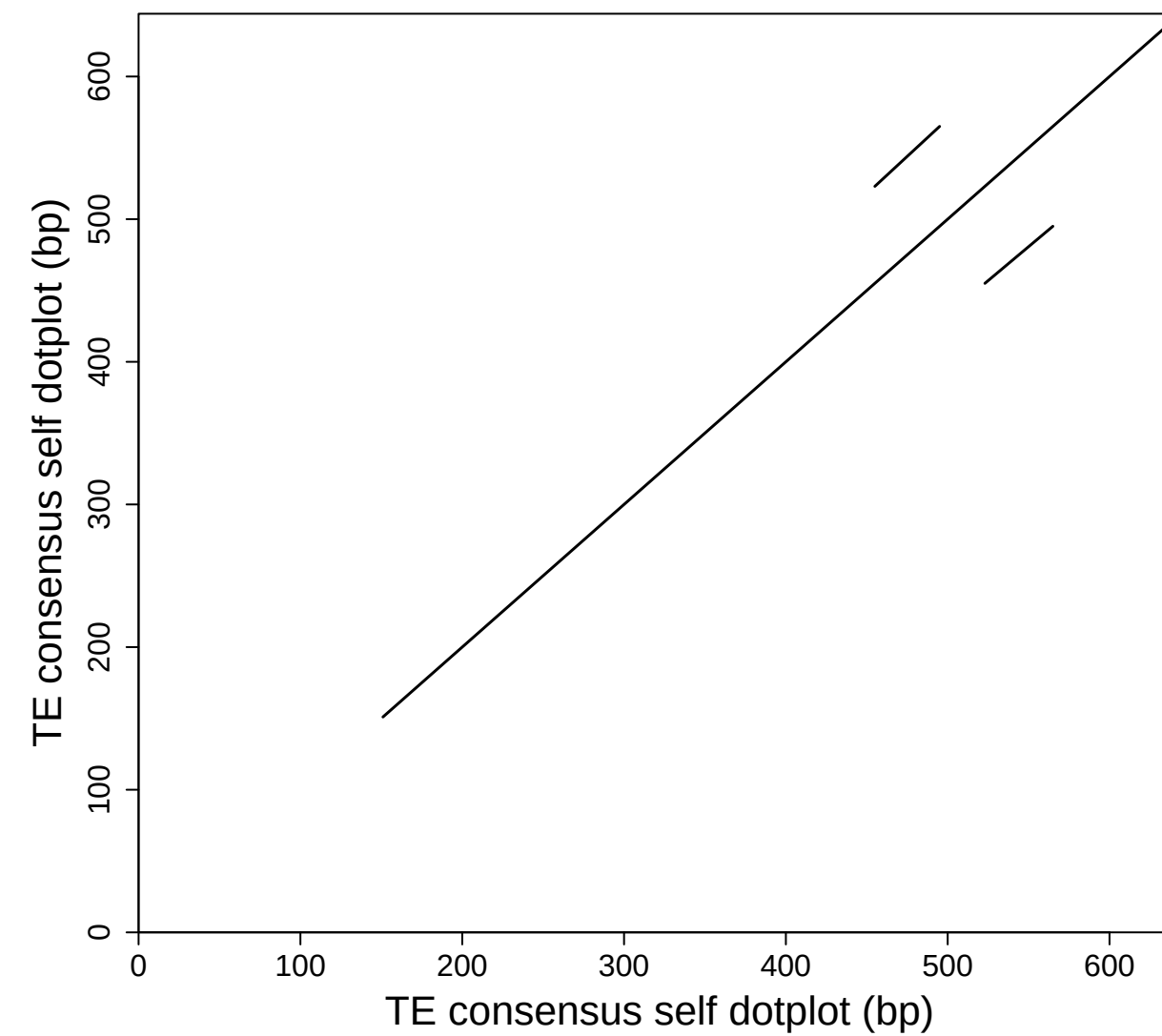
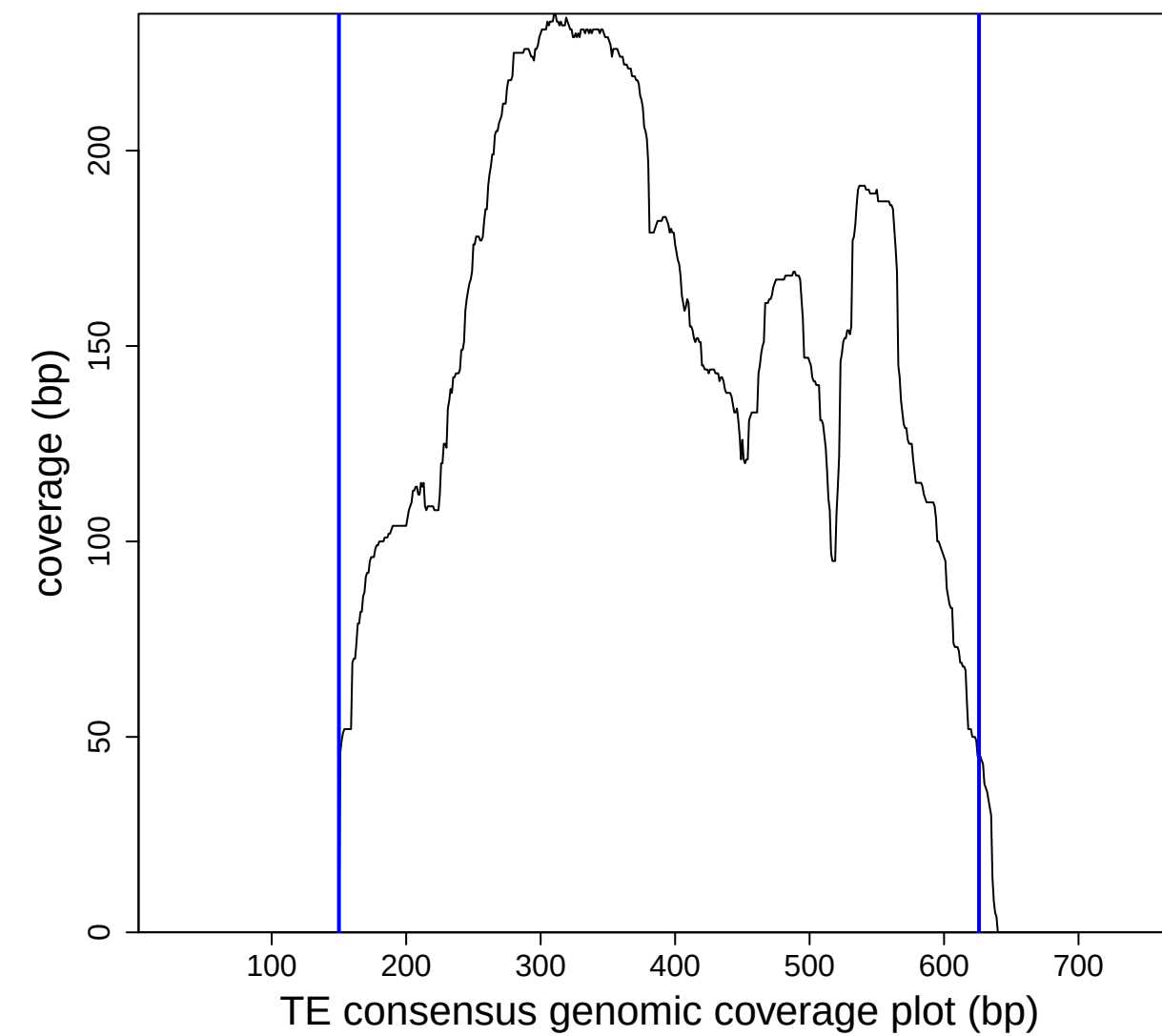
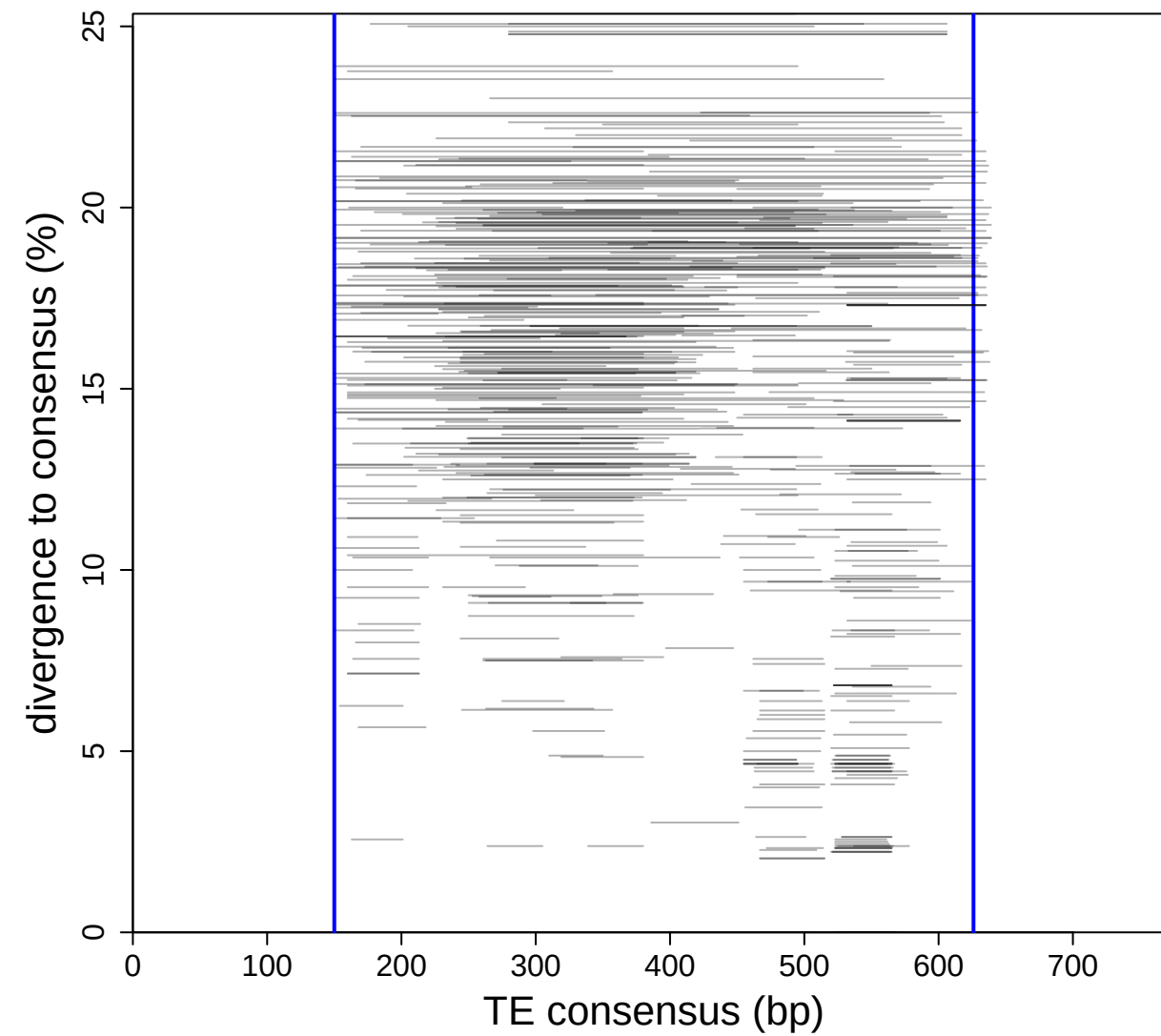


.l.b.bed uf.bed g_1.bed fm_1.bed 0 0 bcIn.fa aln.fa cl.fa gs.fa
size: 471bp; fragments: 526; full length: 9 (>=423.9bp)

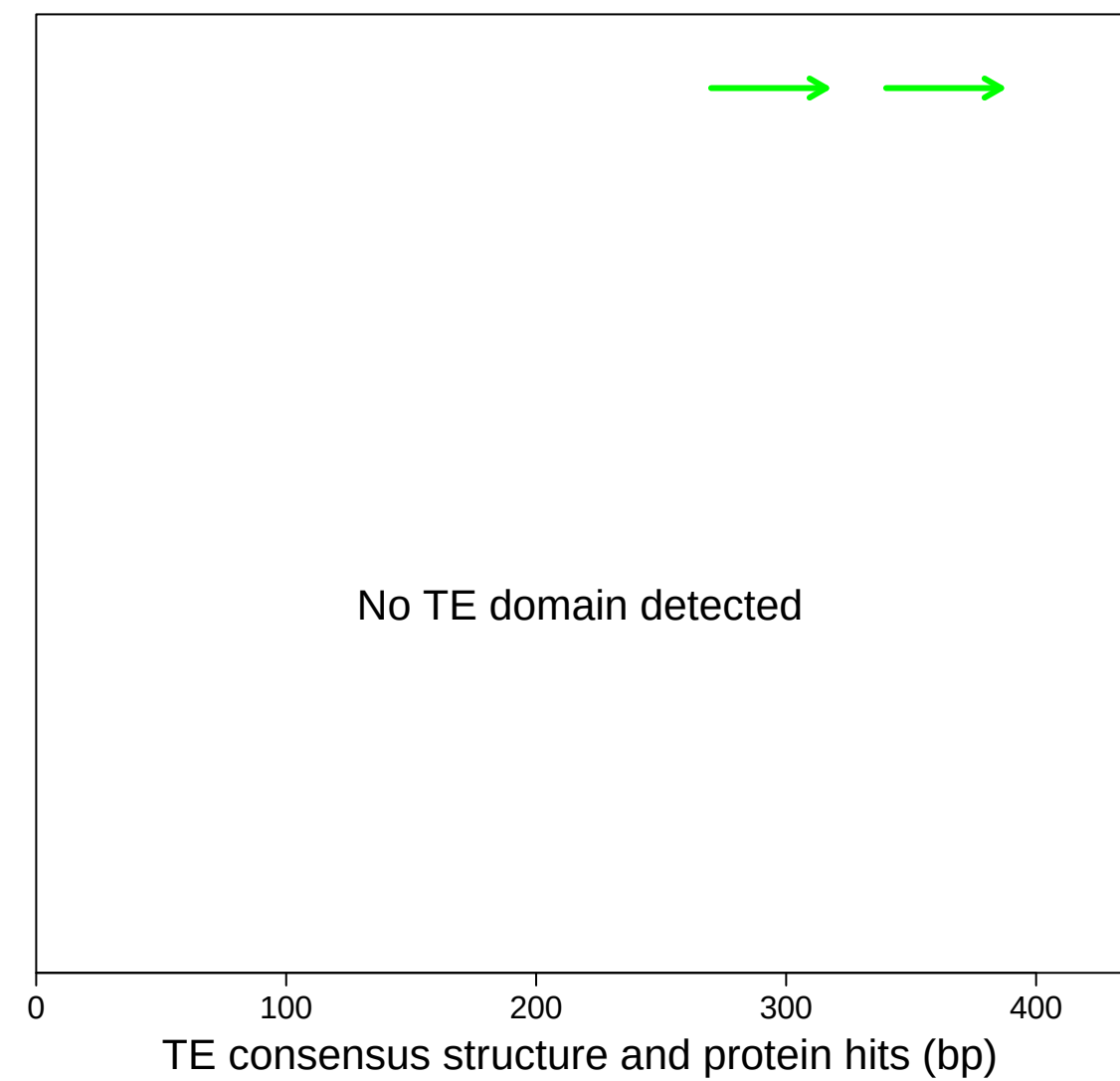
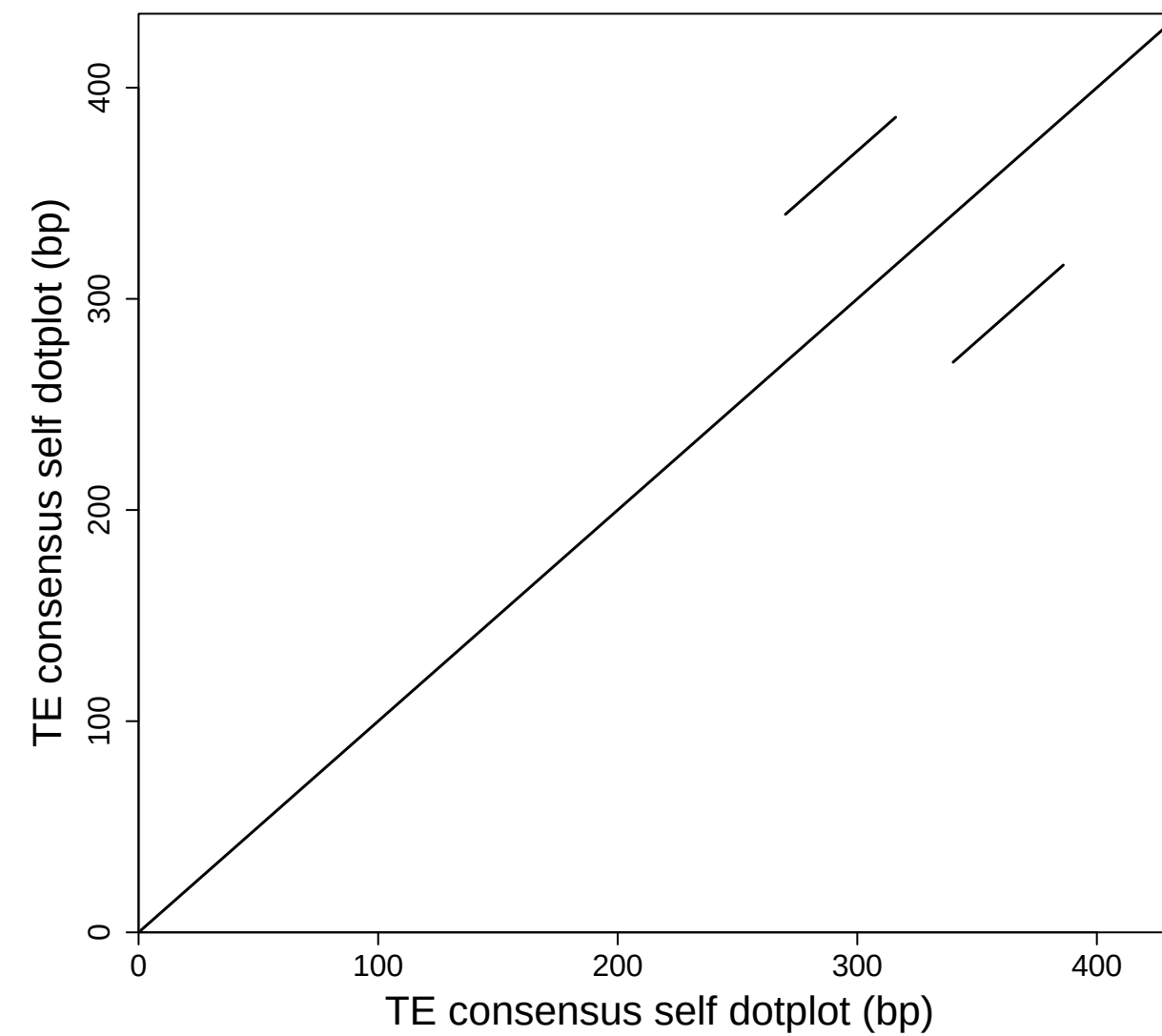
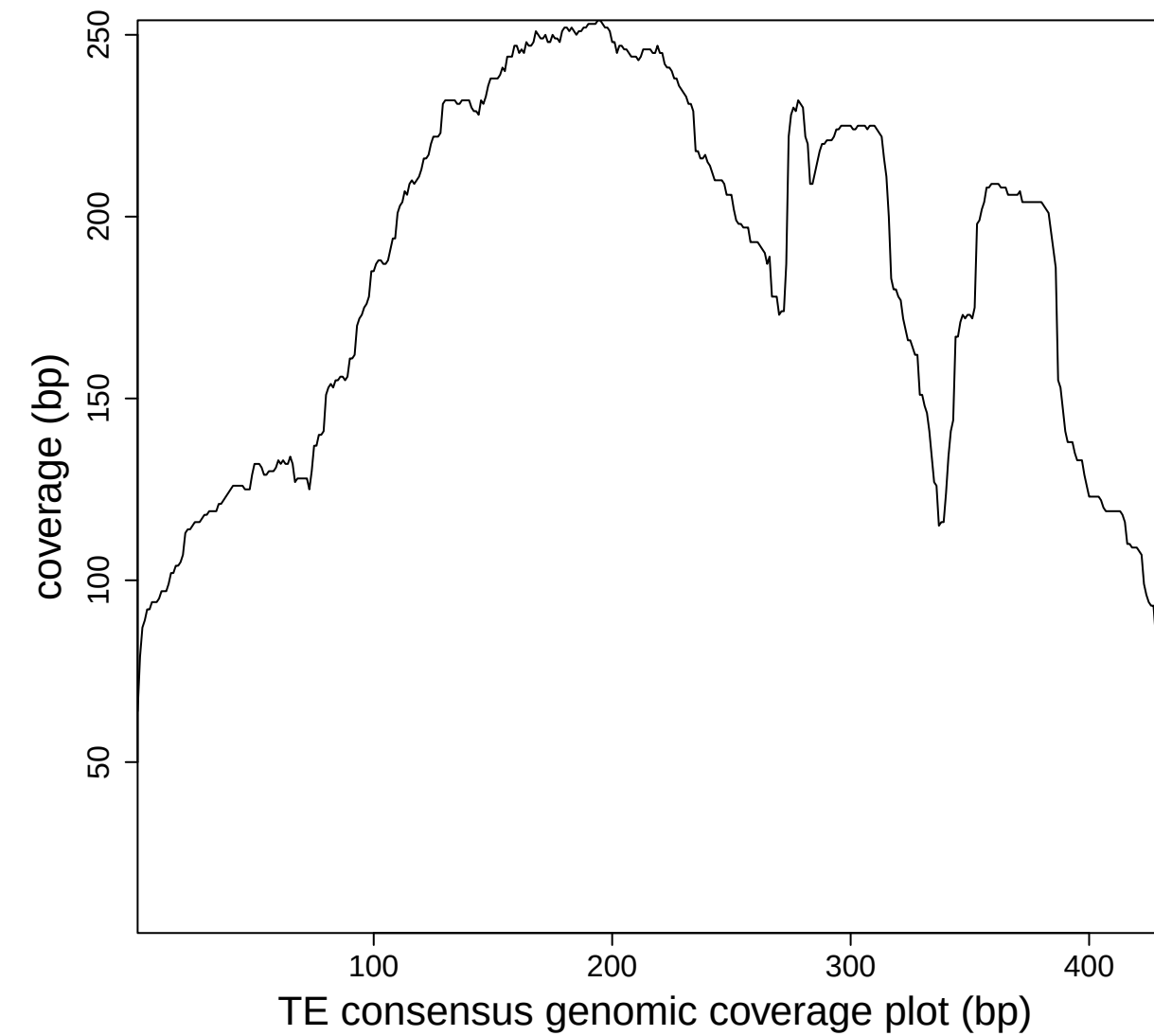
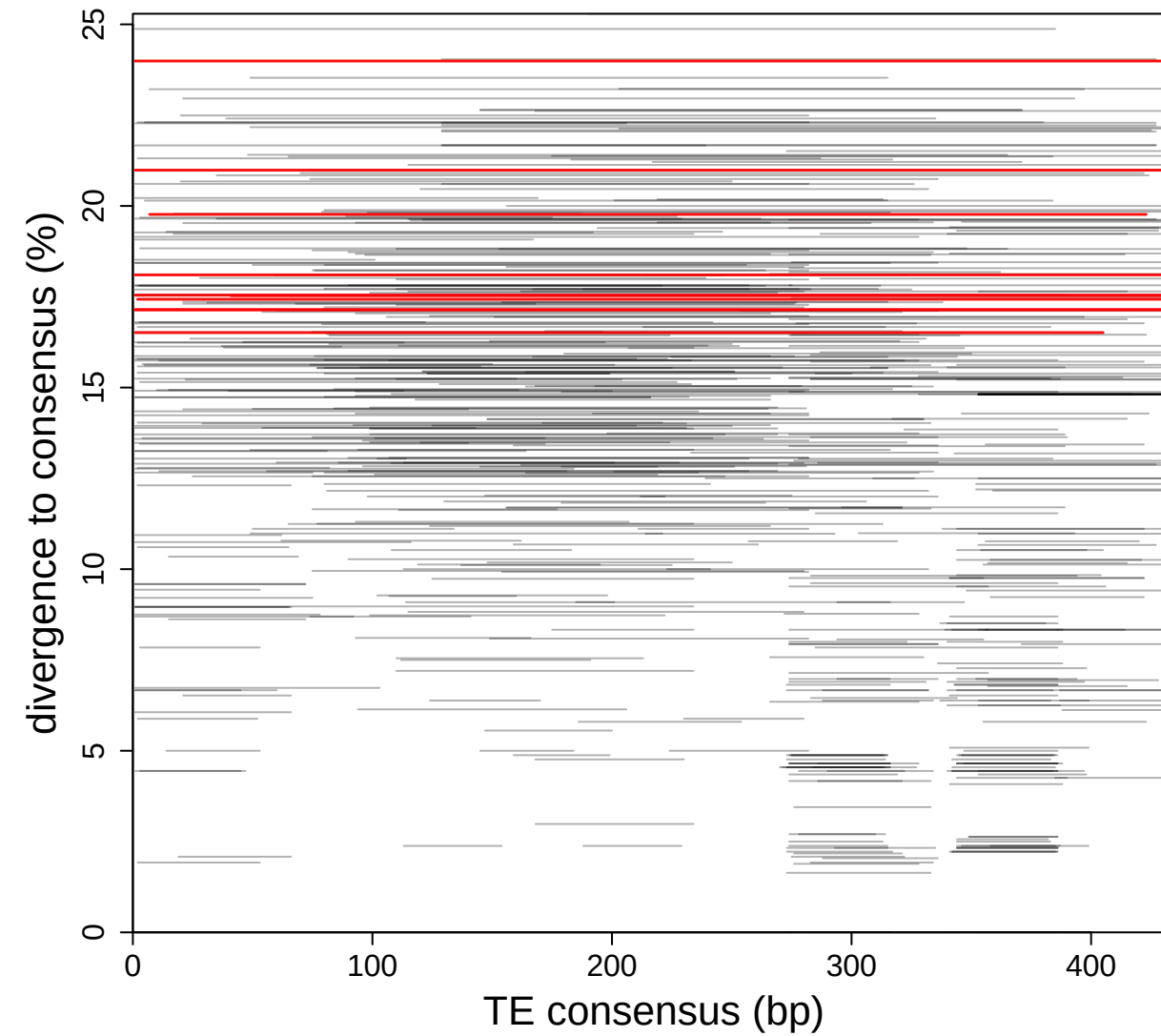
After TEtrimmer 471 bp



6.fasta.b.bed uf.bed g 1.bed fm 1.bed 0 0 bcln.fa aln.fa cl.fa
size: 776bp; fragments: 479; full length: 0 (>=698.4bp)



TE: rnd_5_family_806
size: 435bp; fragments: 563; full length: 10 (≥ 391.5 bp)



Before TEtrimmer 435 bp

Dotplot

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

