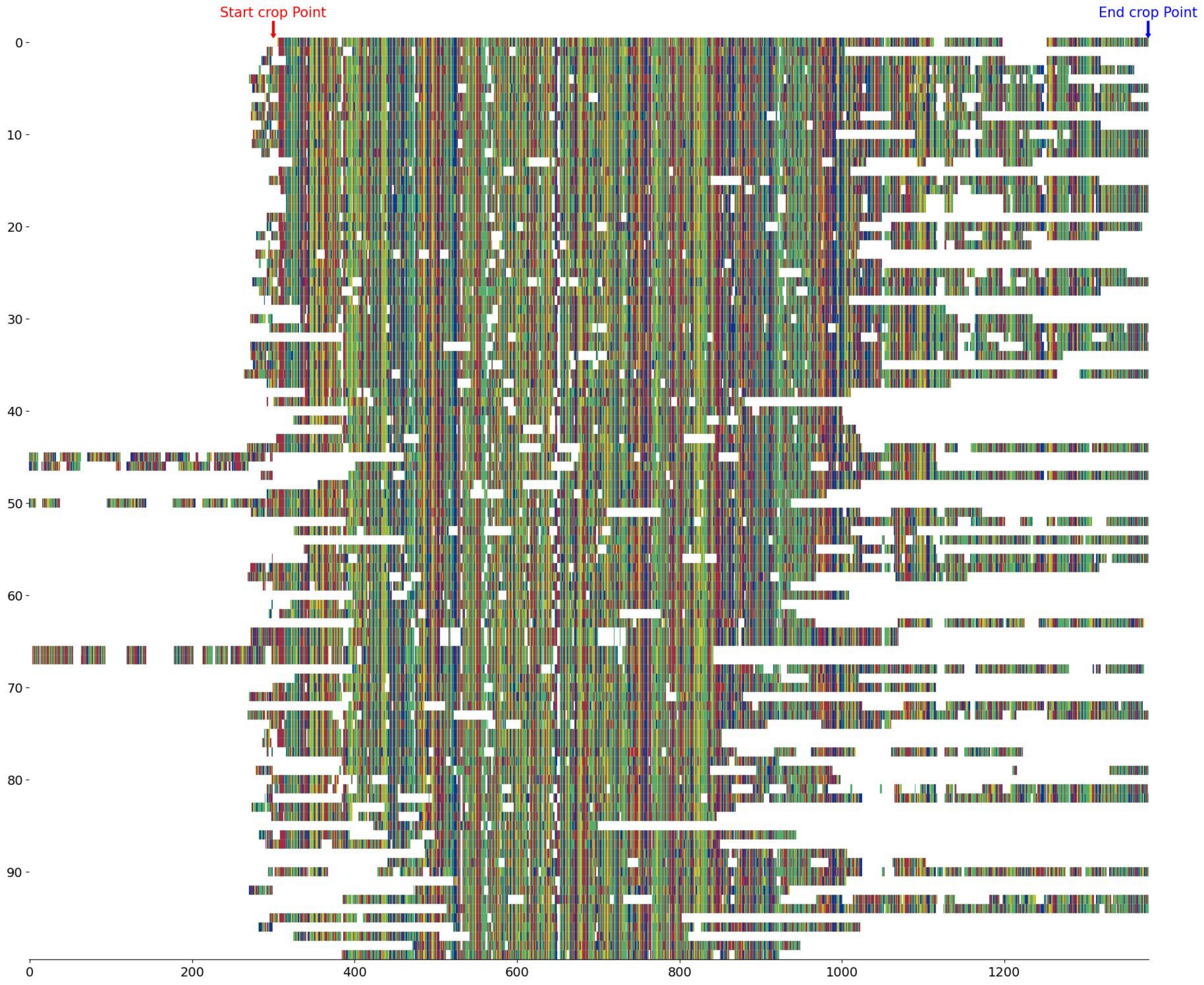


Start crop Point

End crop Point

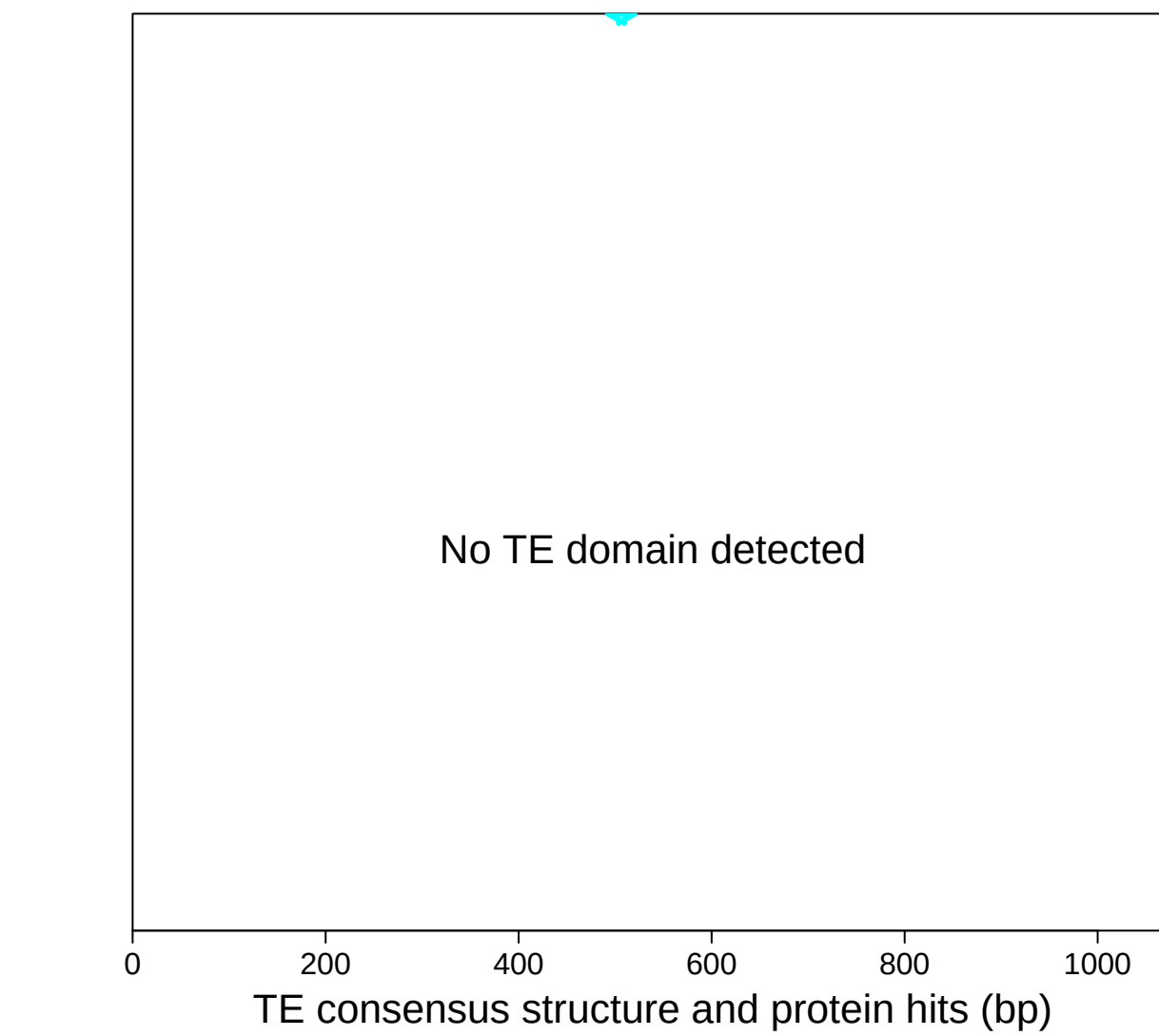
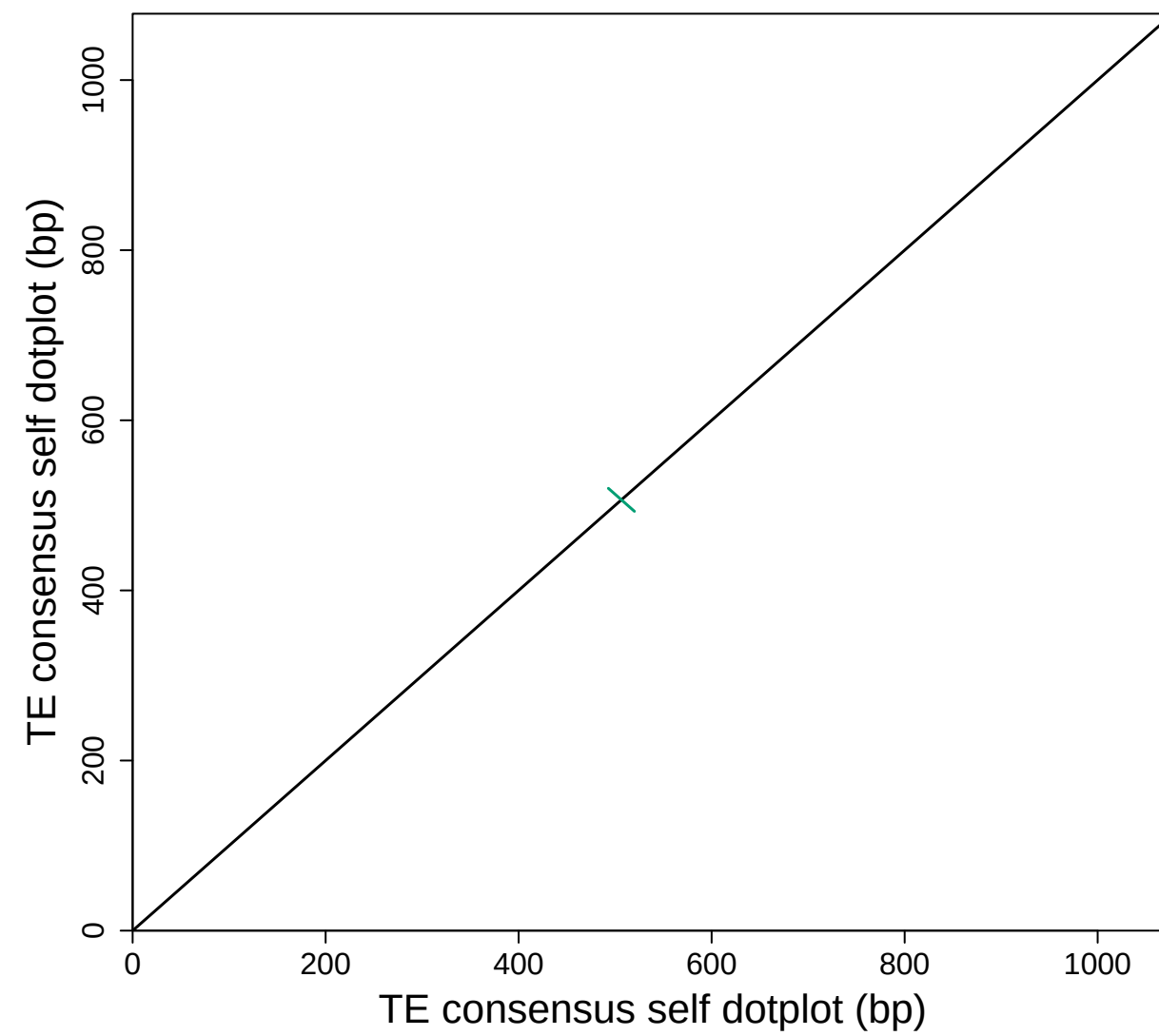
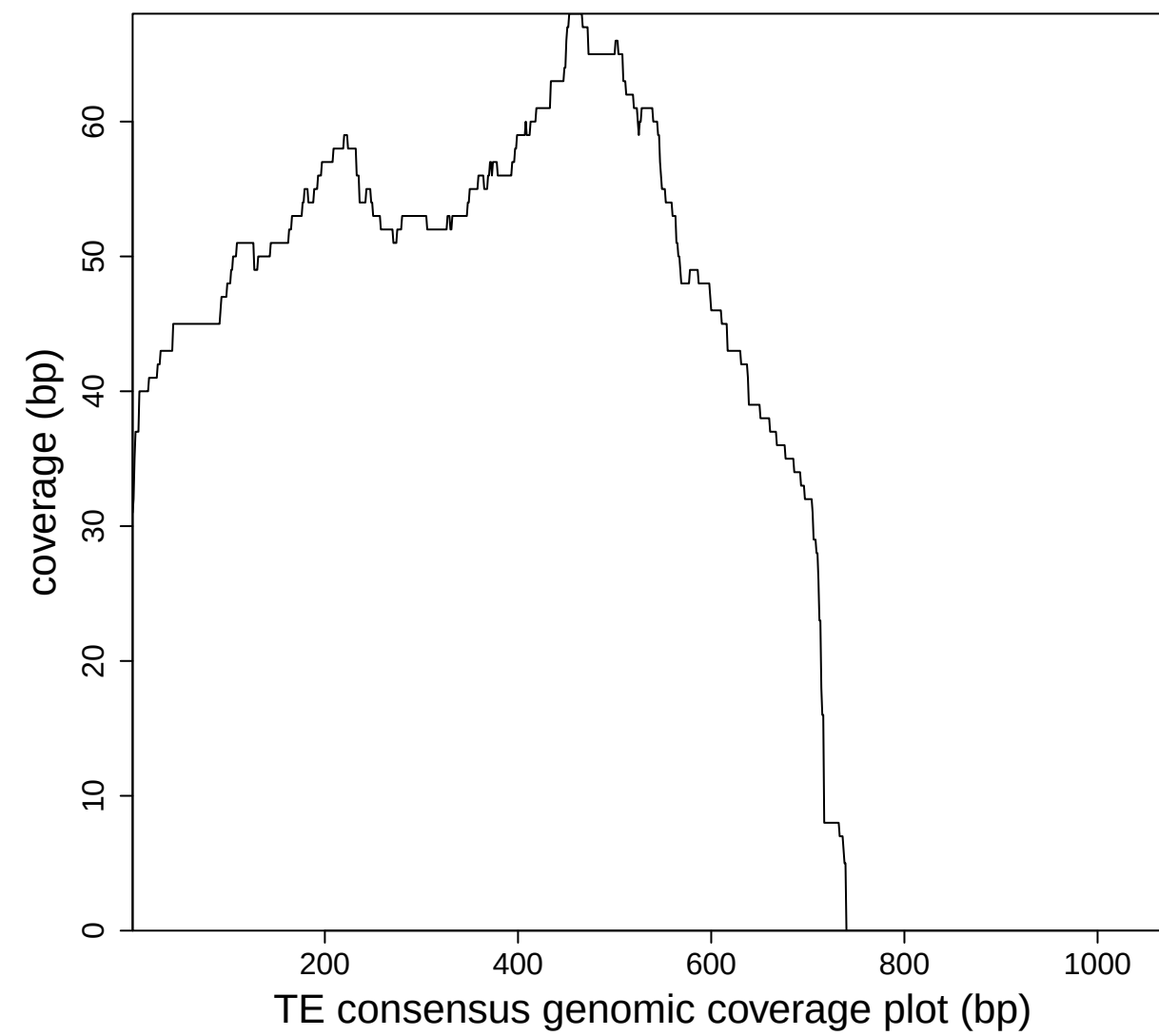
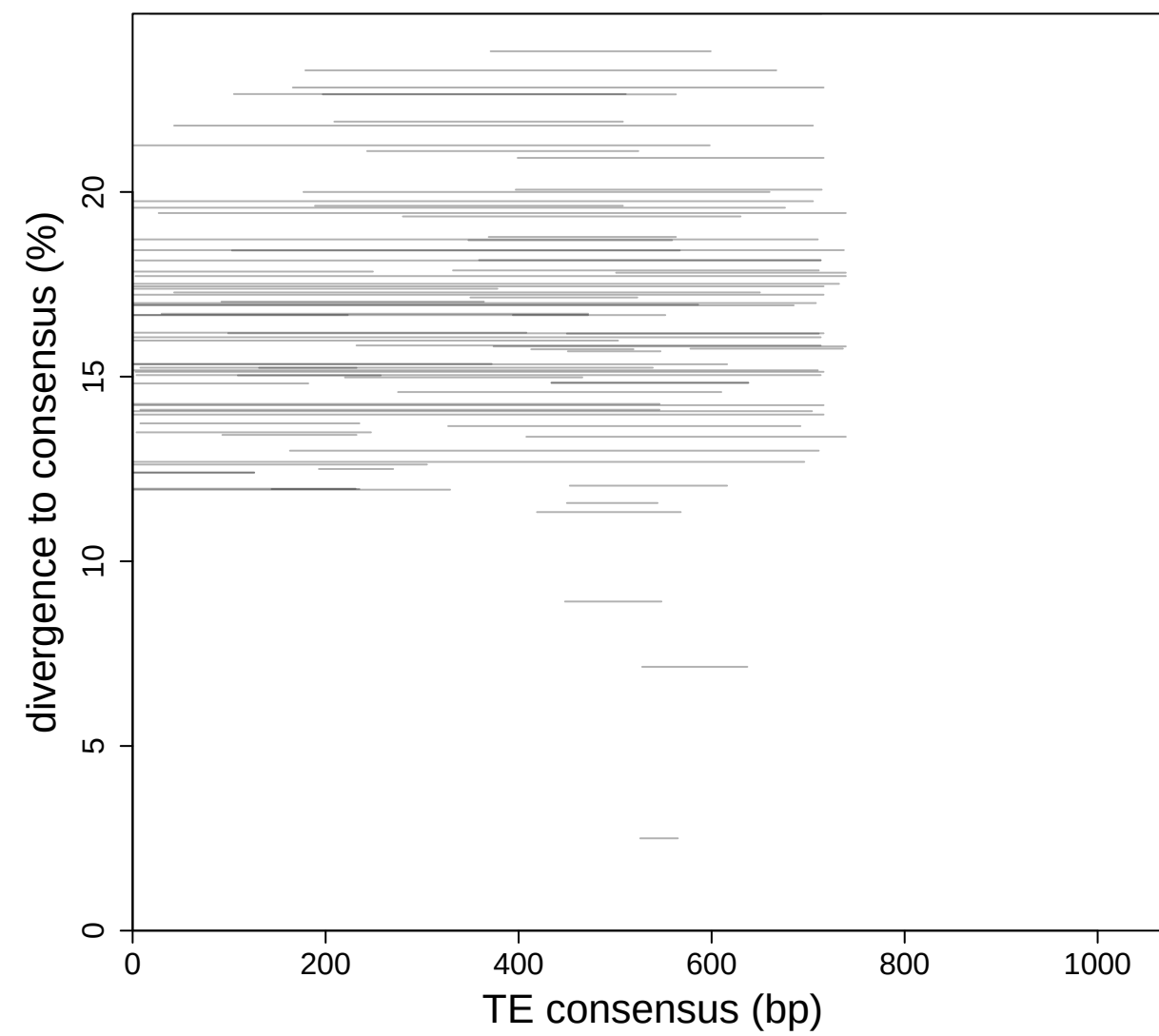
MSA length = 1078



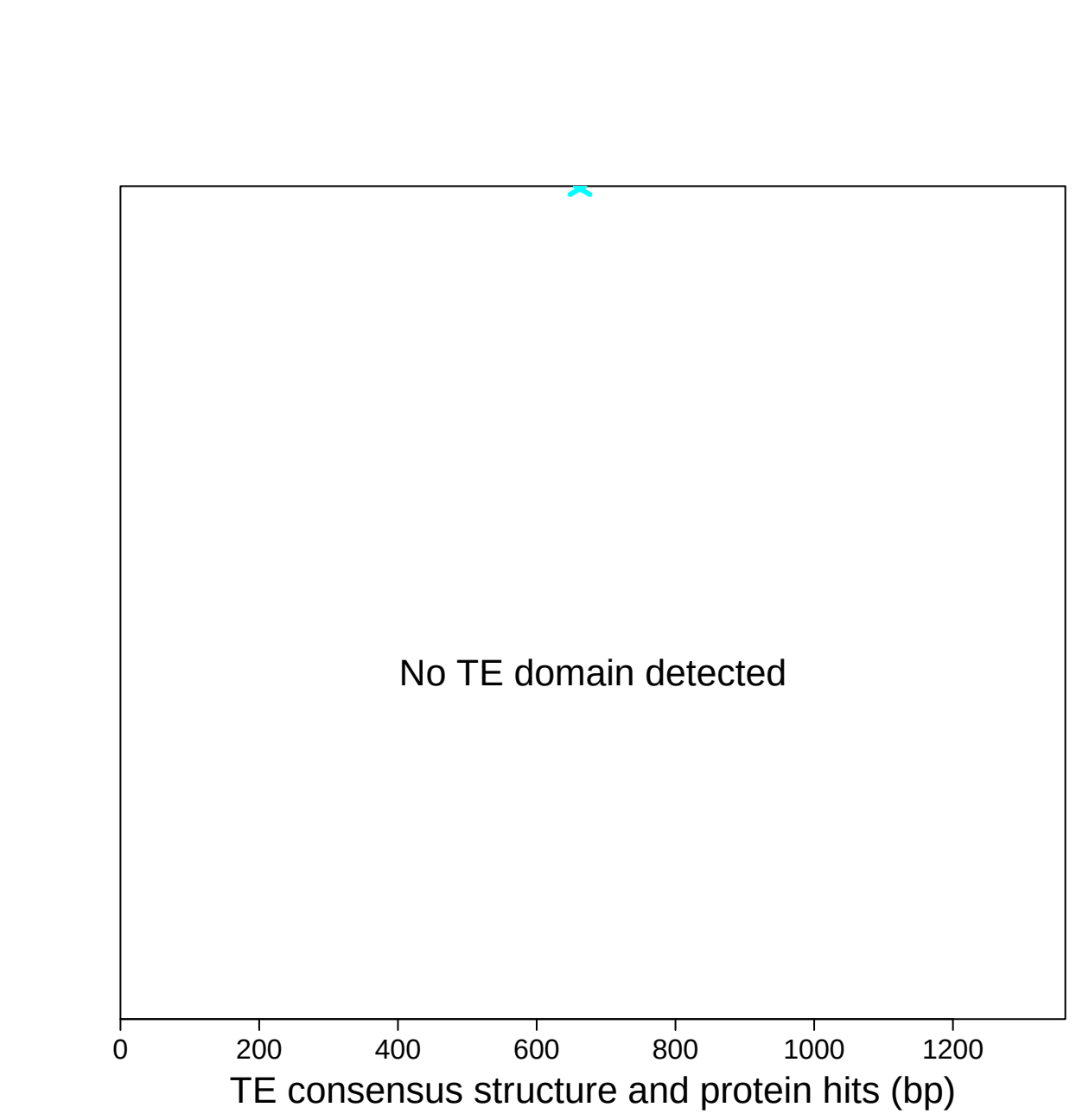
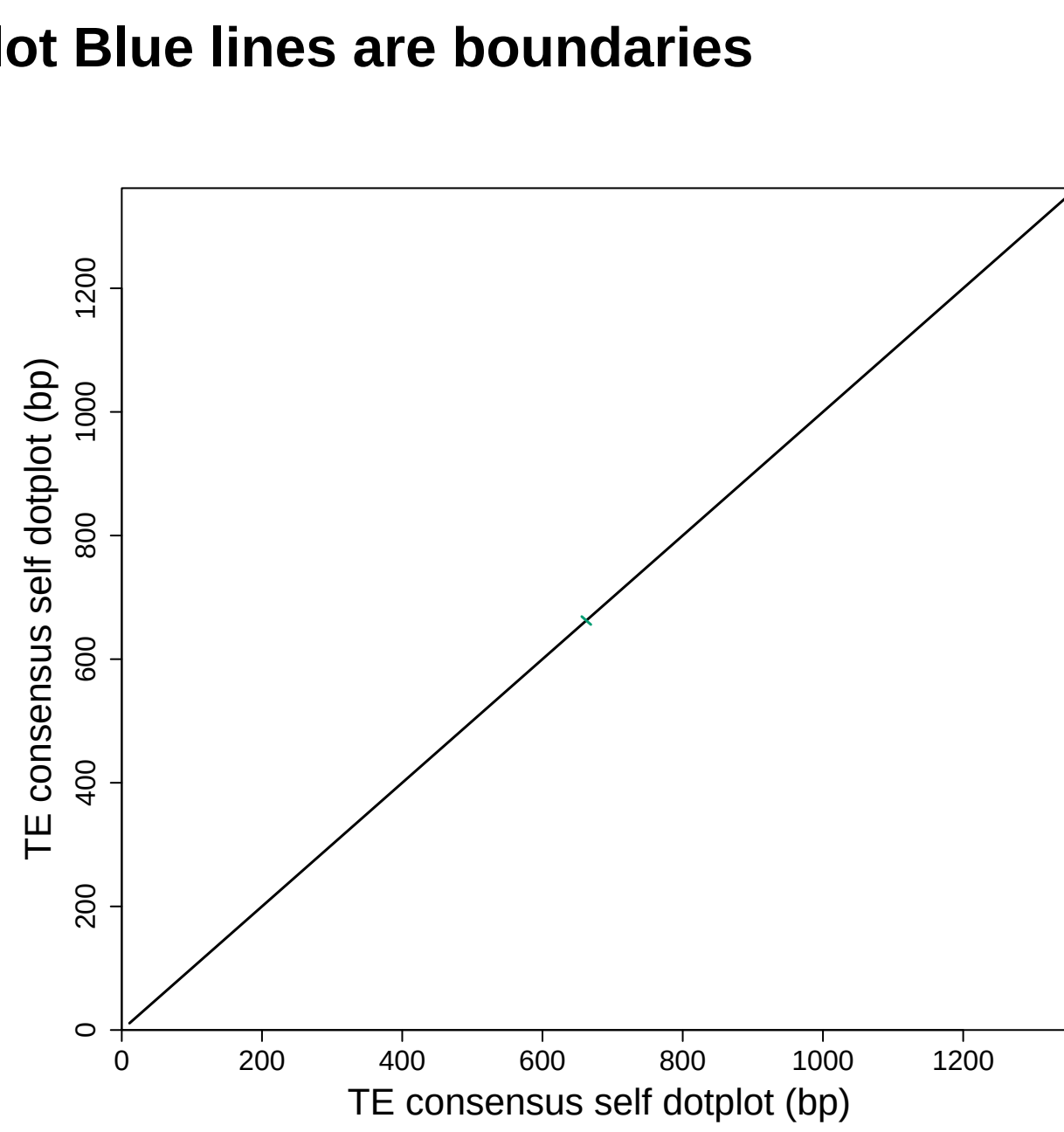
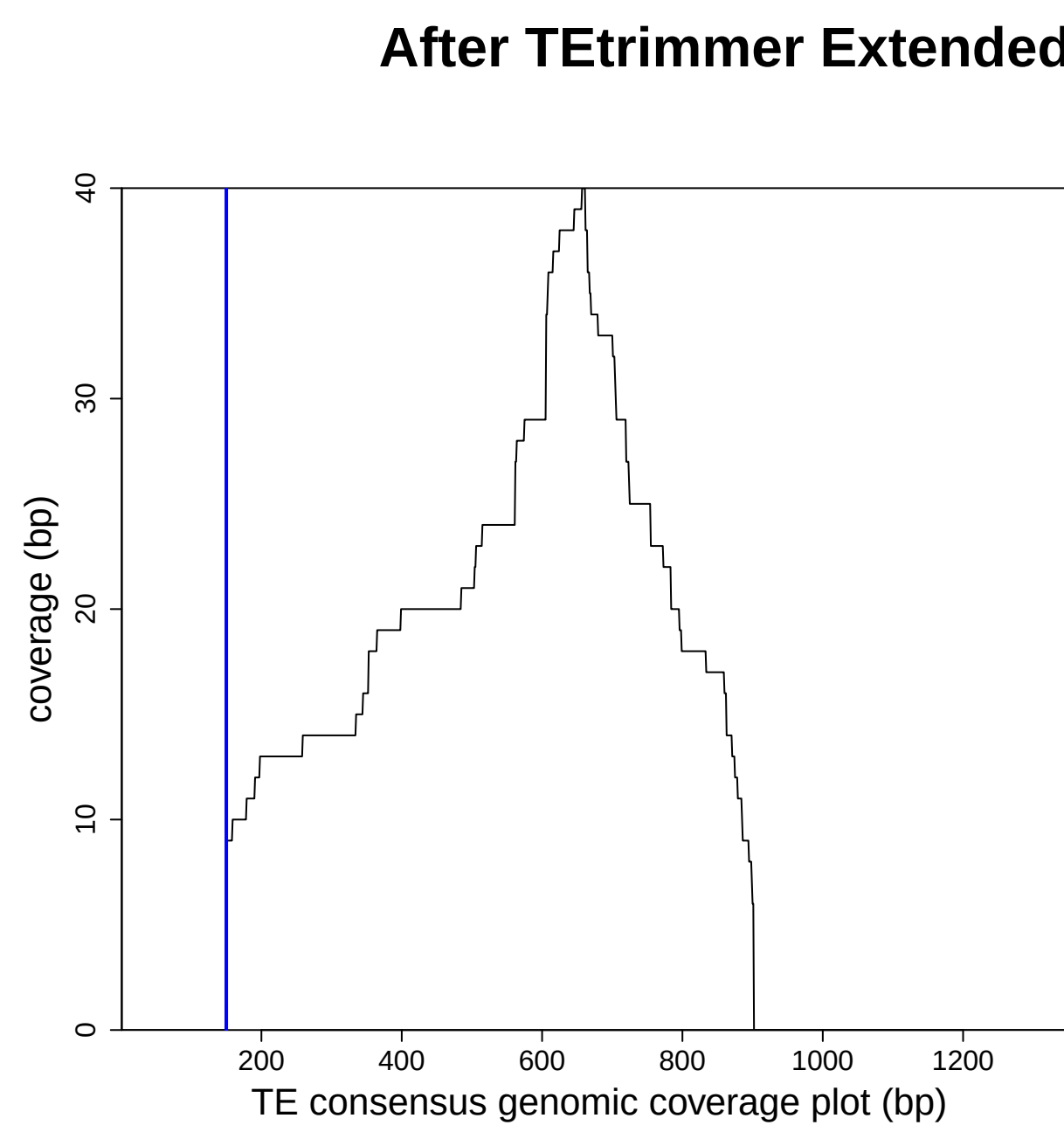
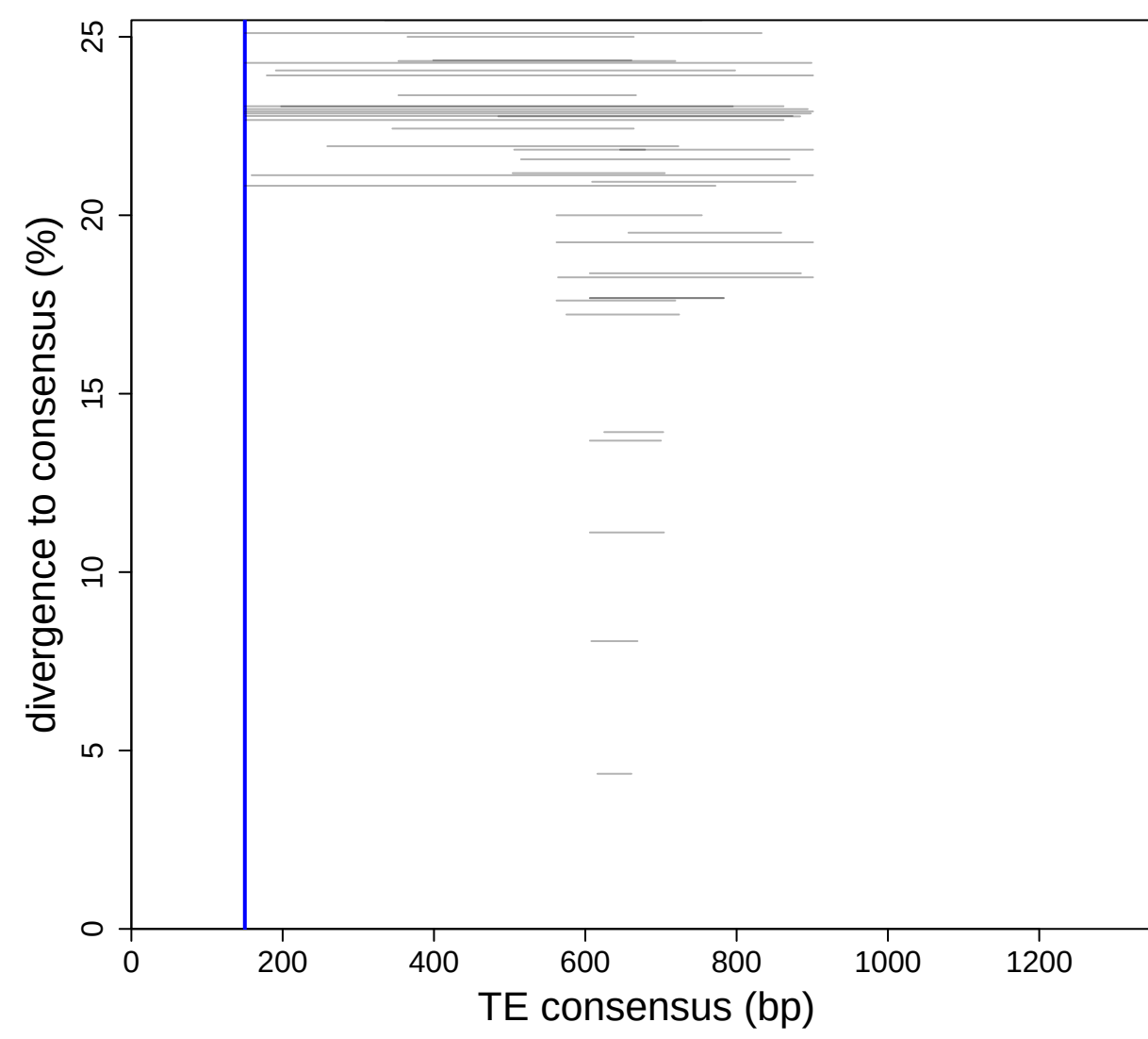


a.b.bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.f
size: 1078bp; fragments: 91; full length: 0 (>=970.2bp)

After TEtrimmer 1078 bp

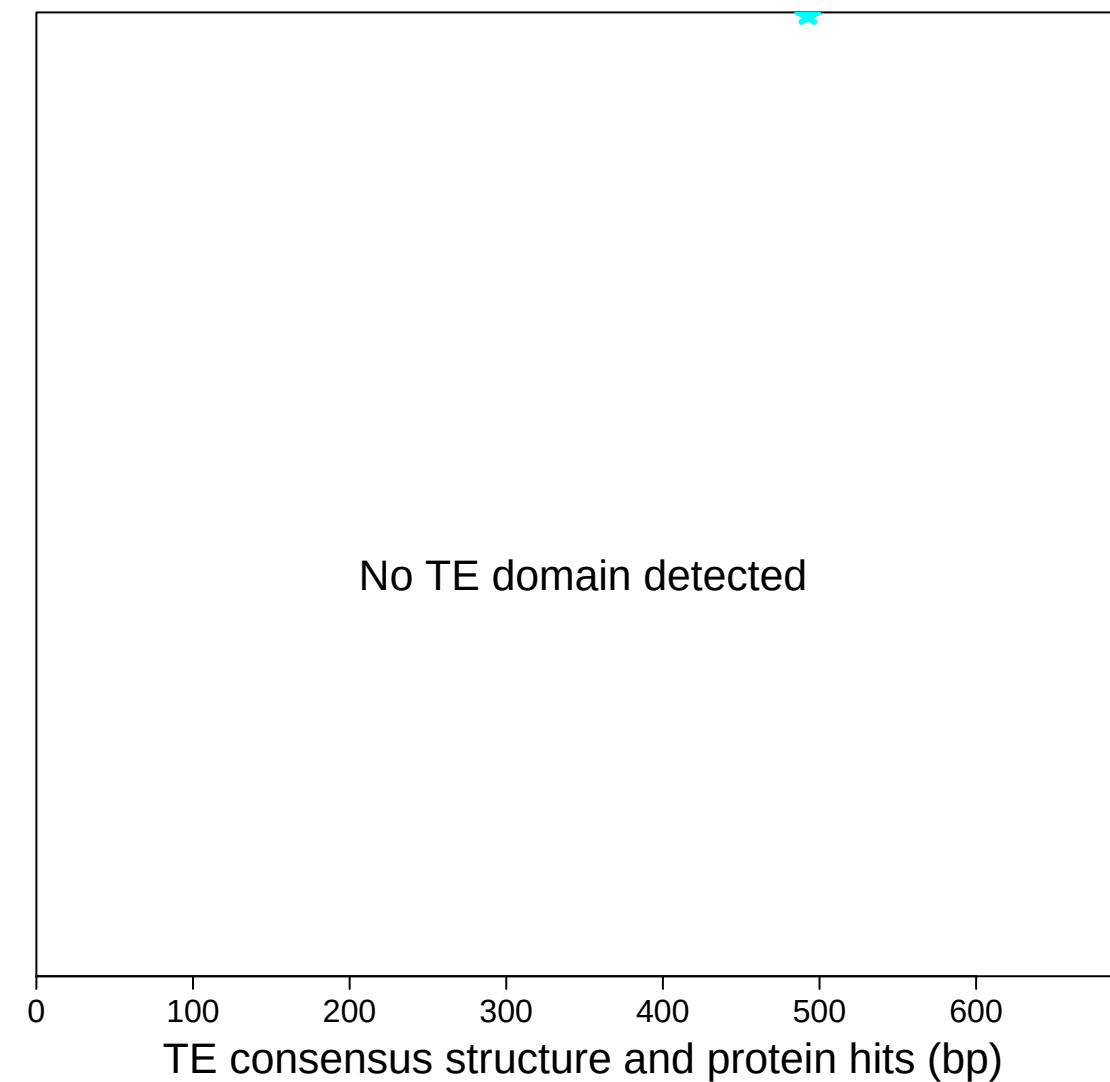
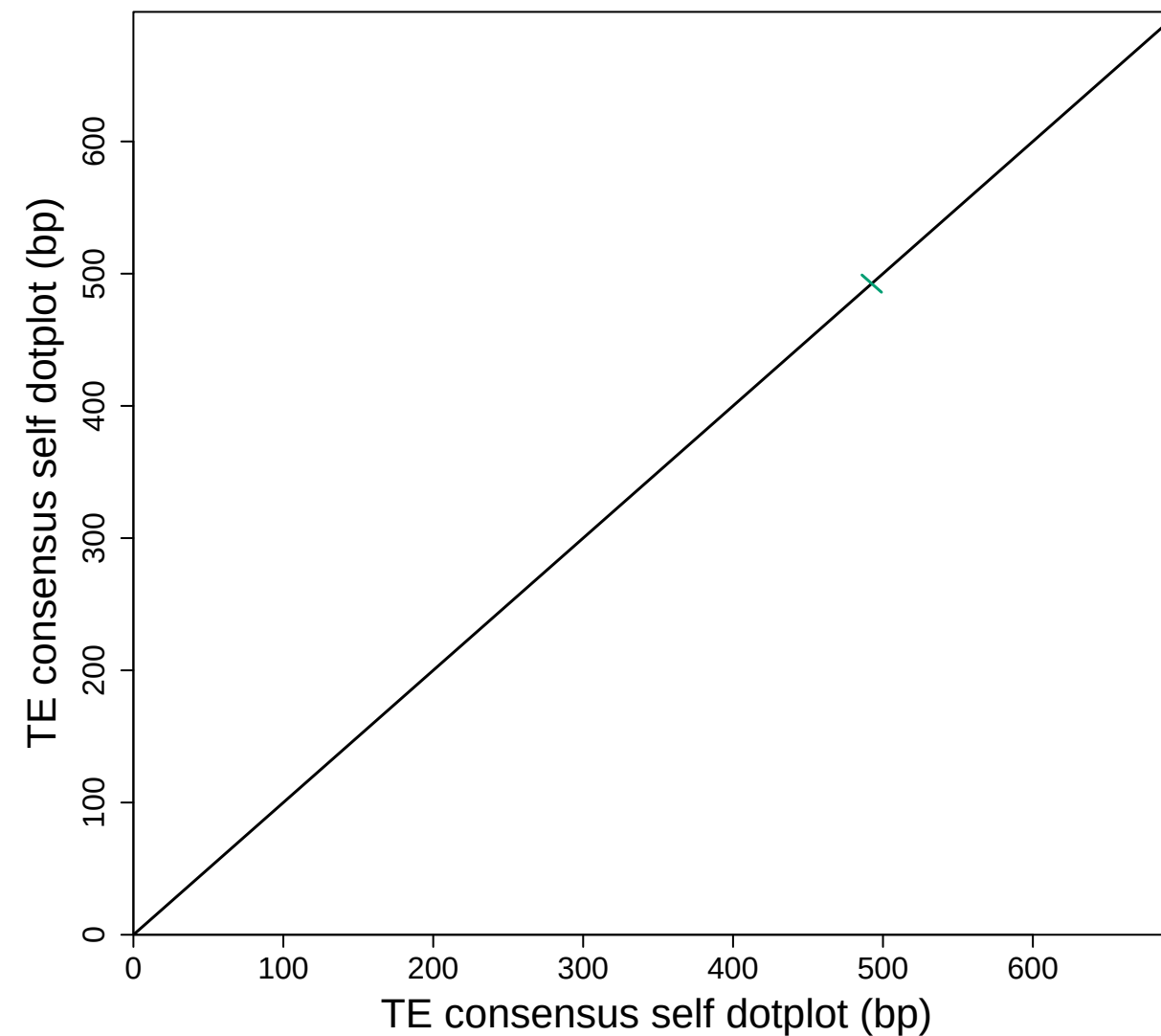
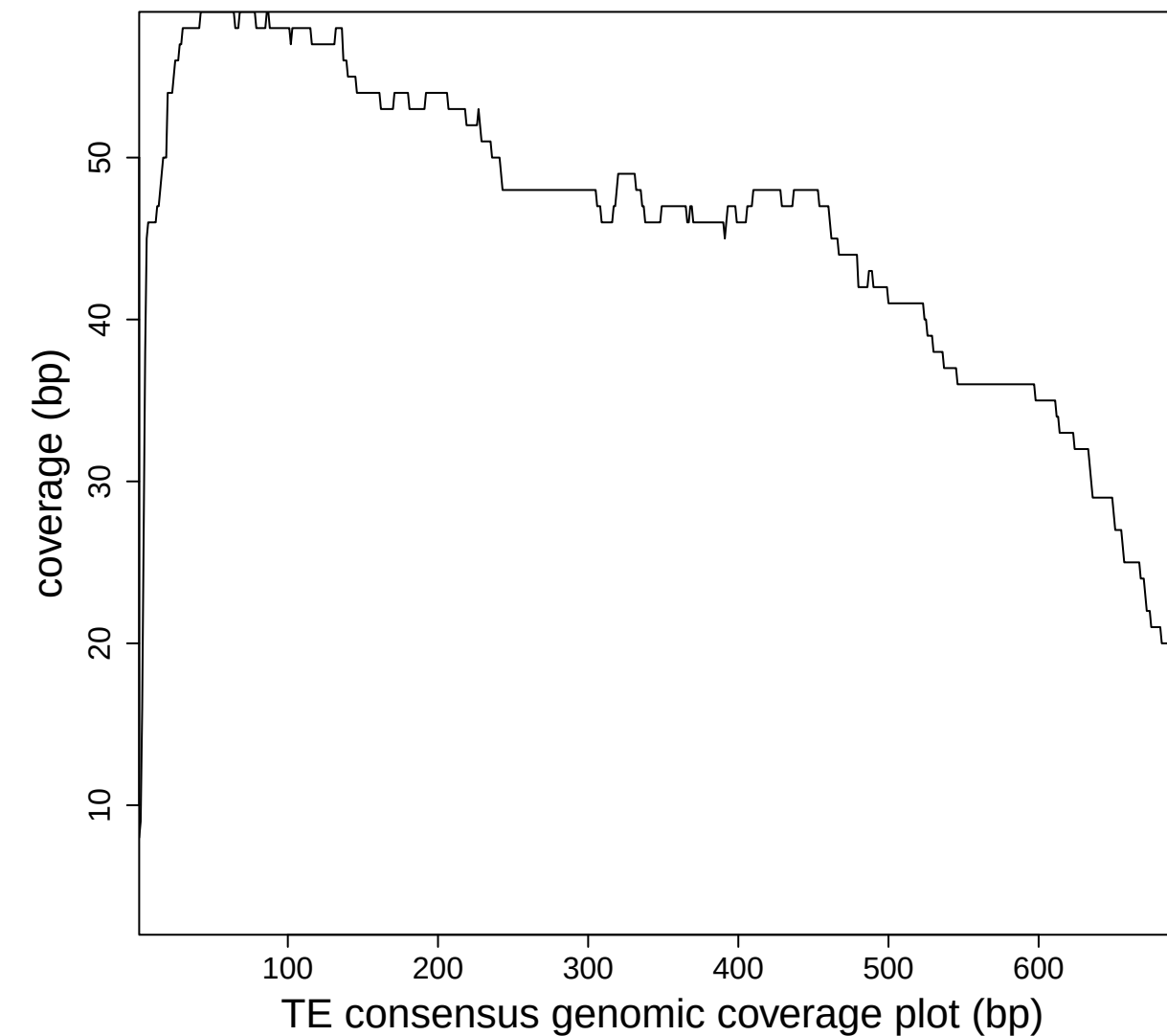
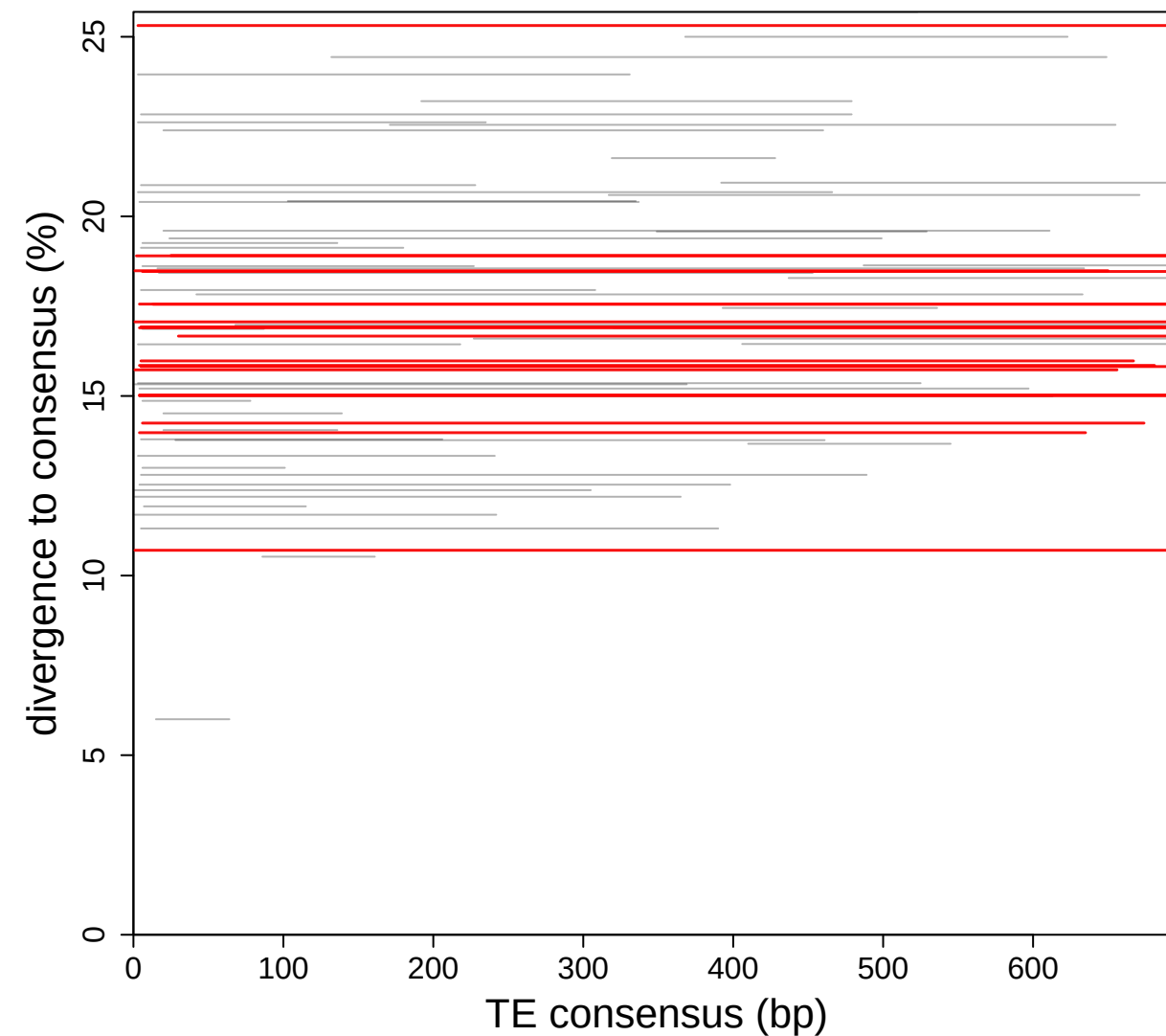


3.fasta.b.bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.f
size: 1362bp; fragments: 40; full length: 0 (>=1225.8bp)



Before TEtrimmer 698 bp

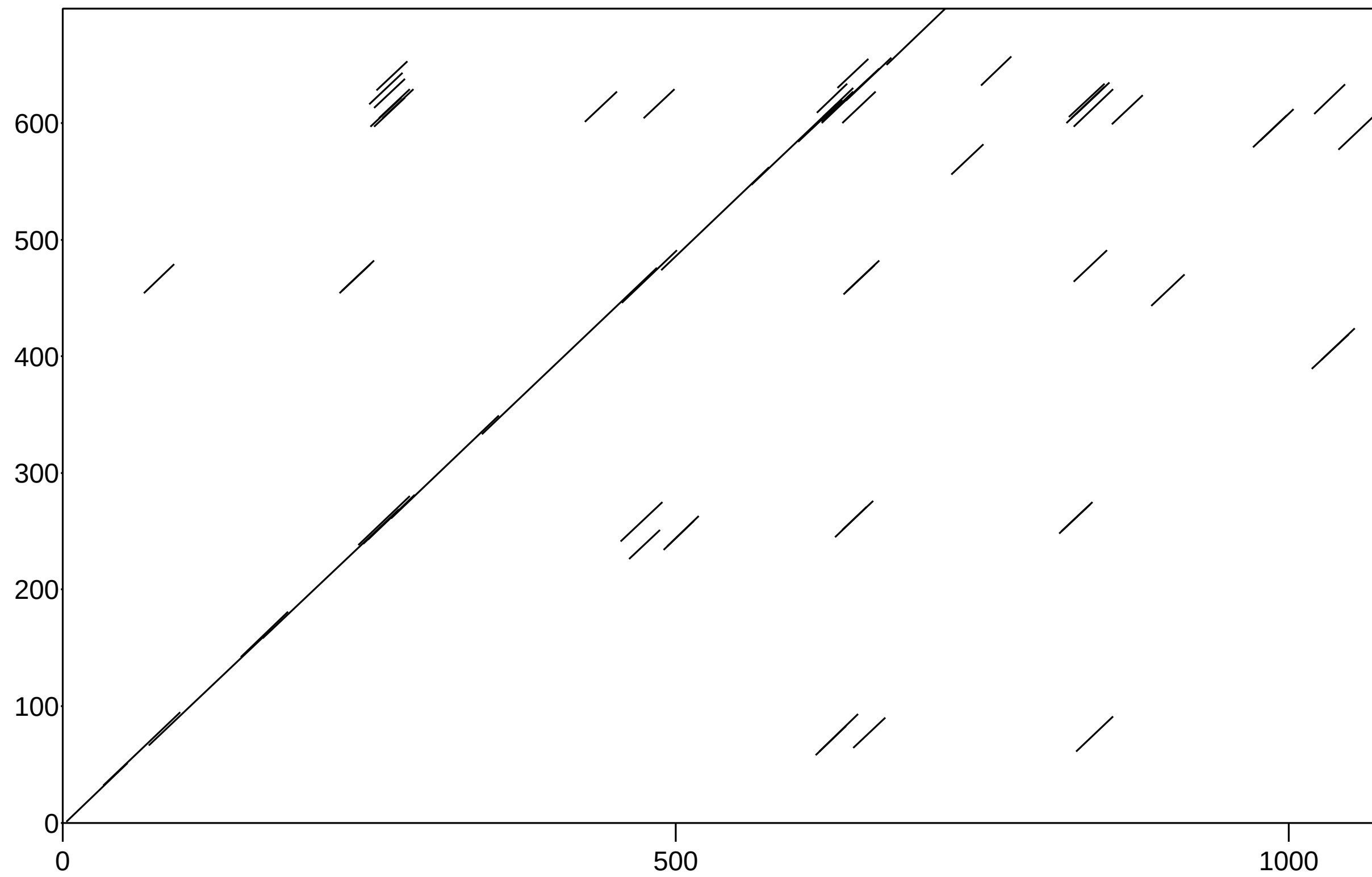
TE: rnd_4_family_2863
size: 698bp; fragments: 77; full length: 20 (≥ 628.2 bp)



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)