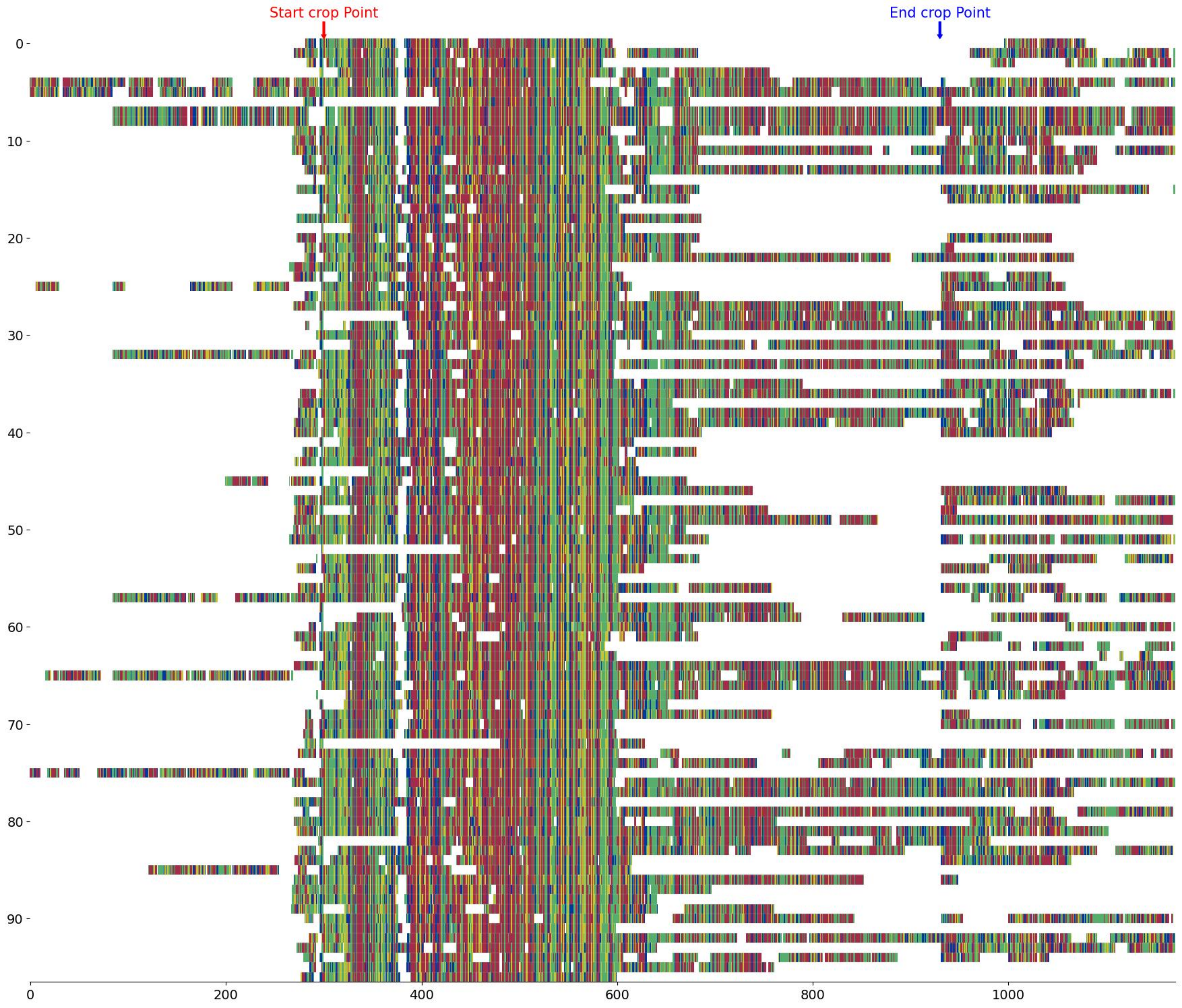
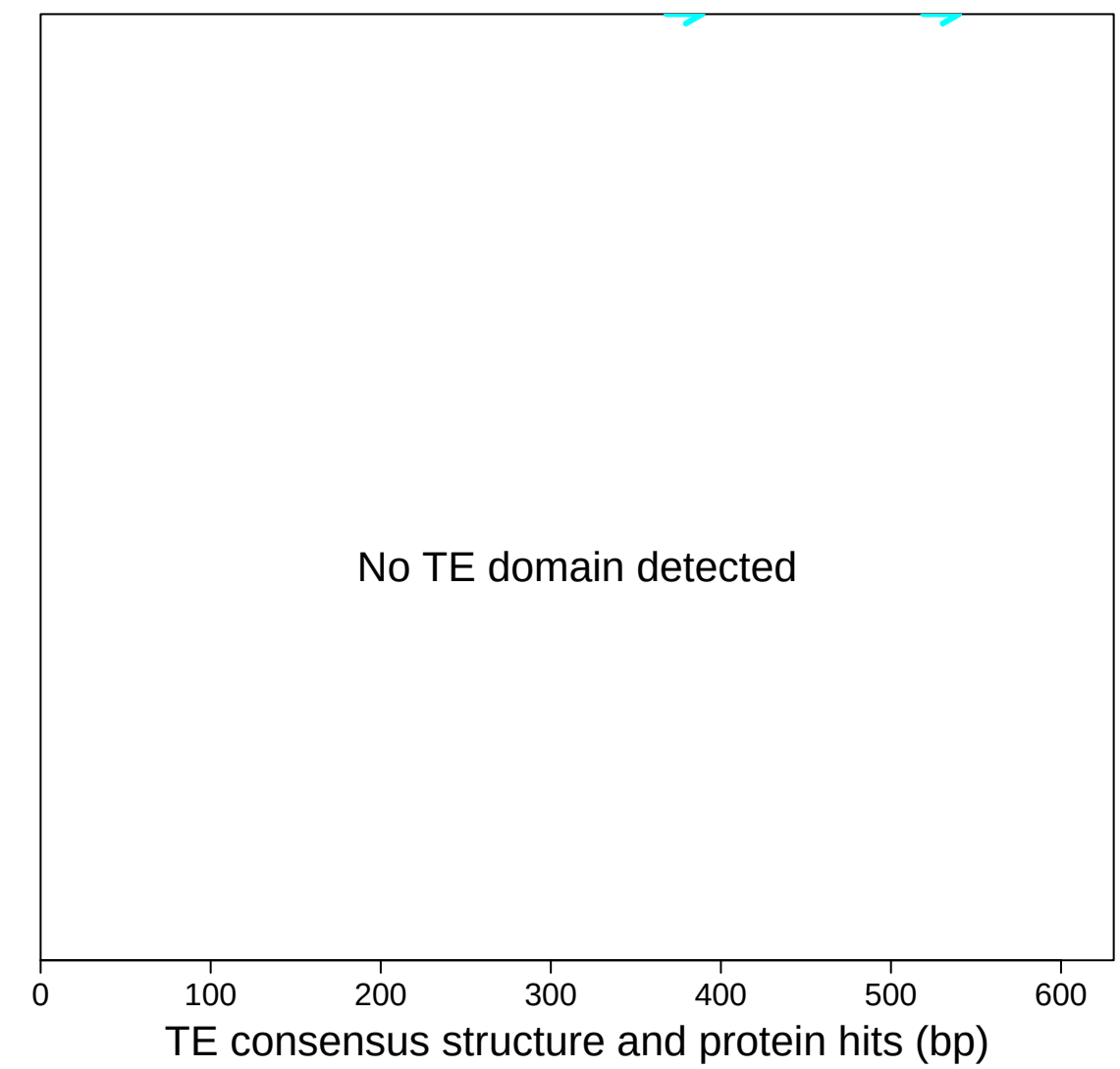
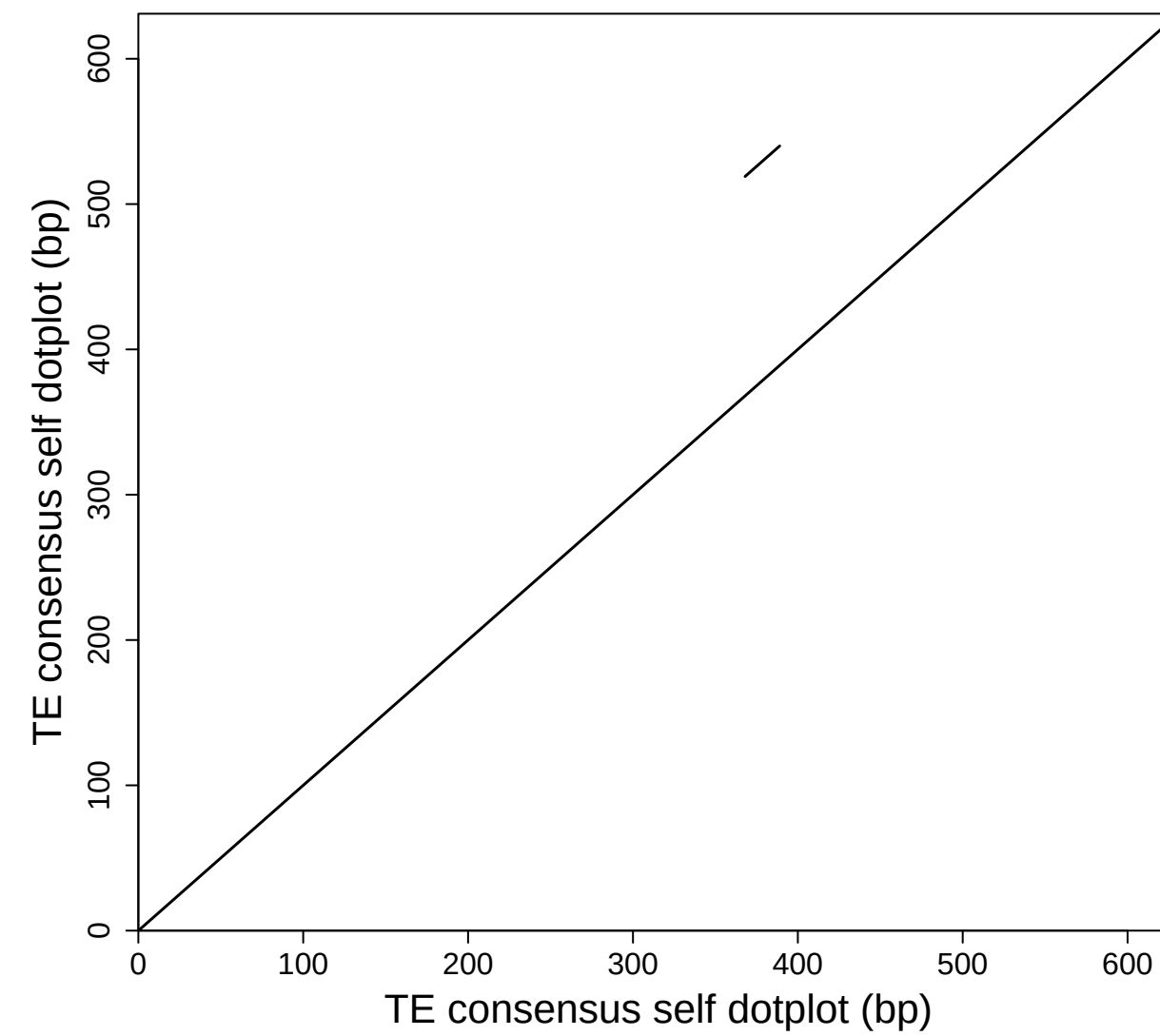
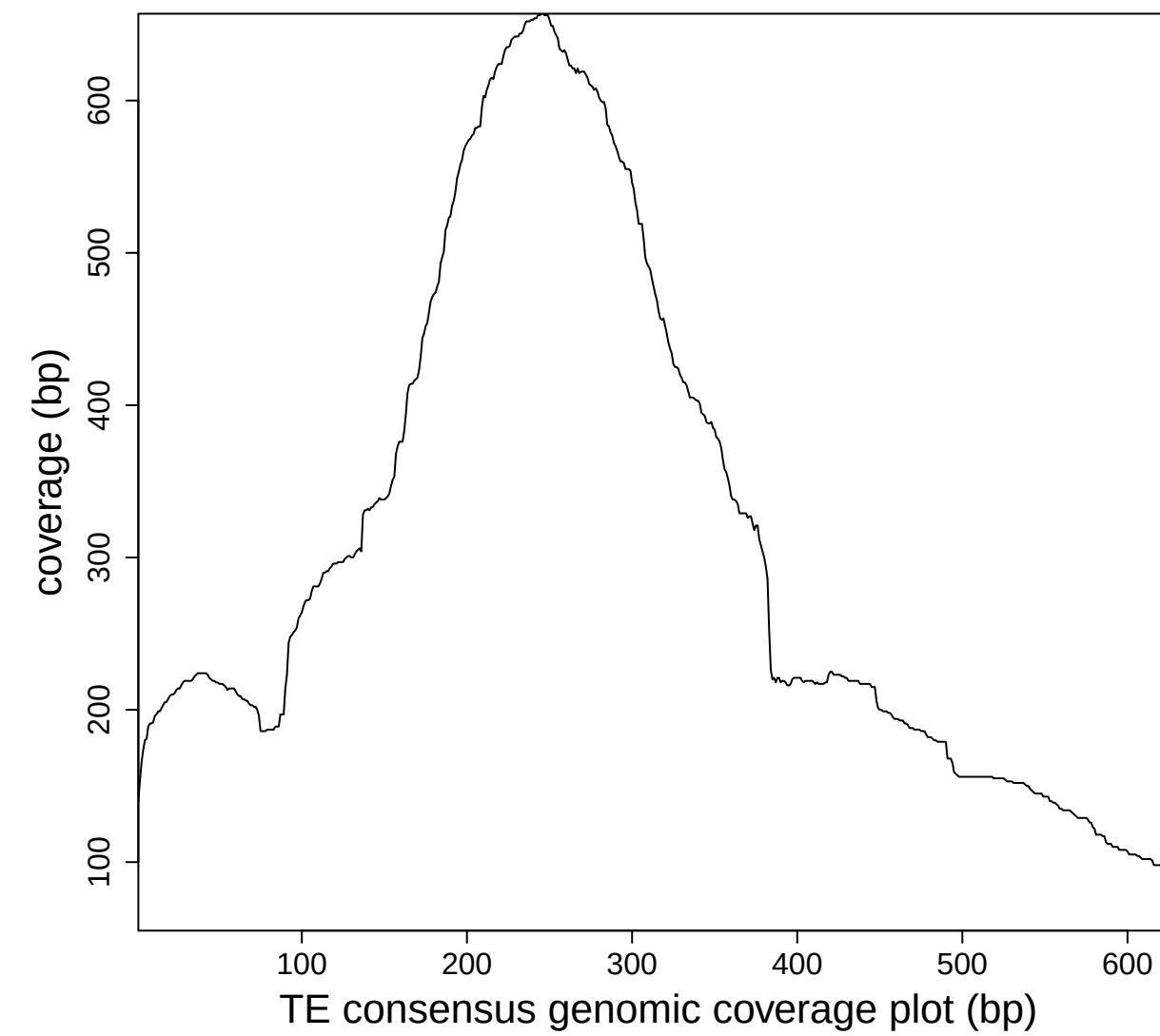
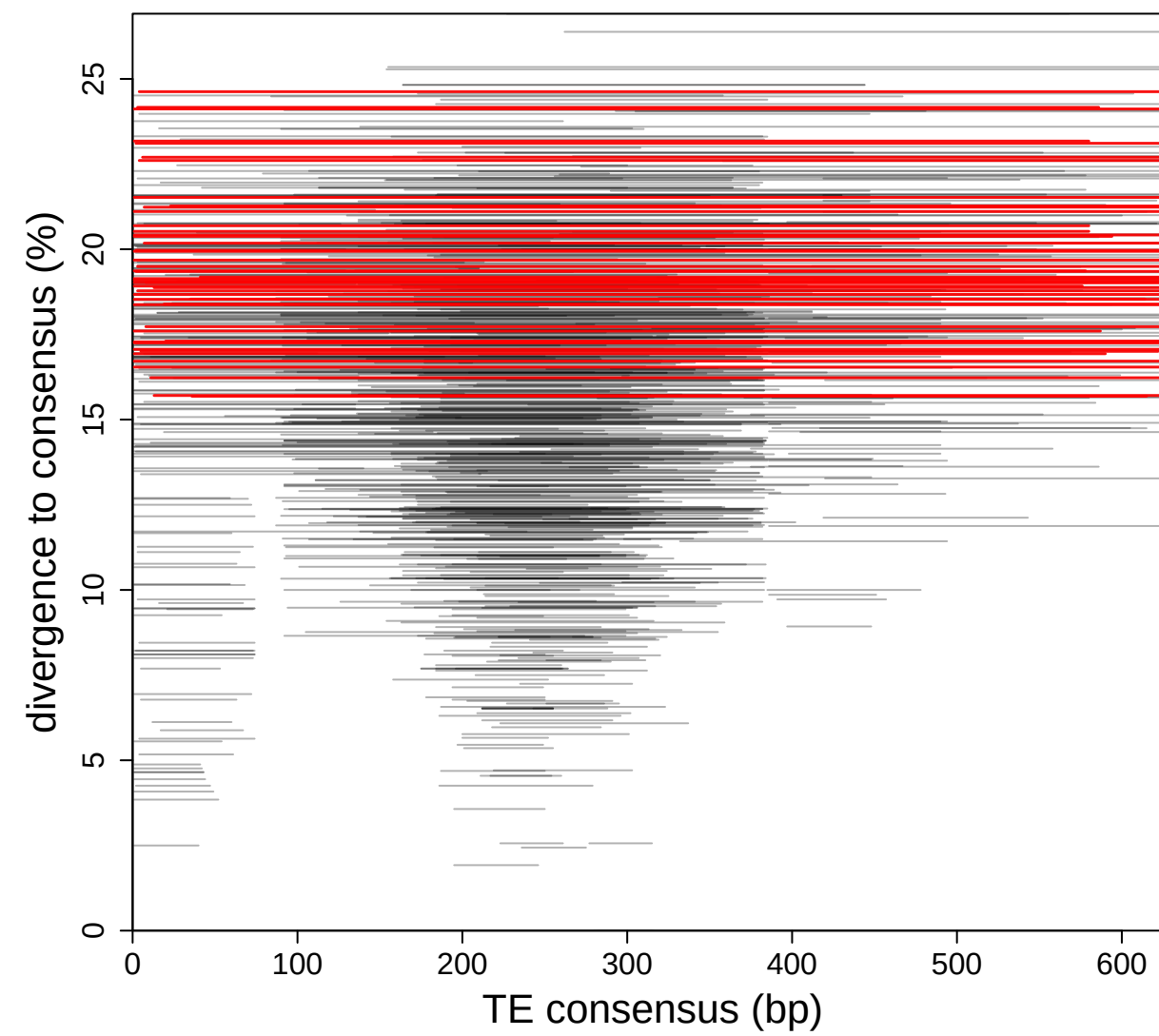






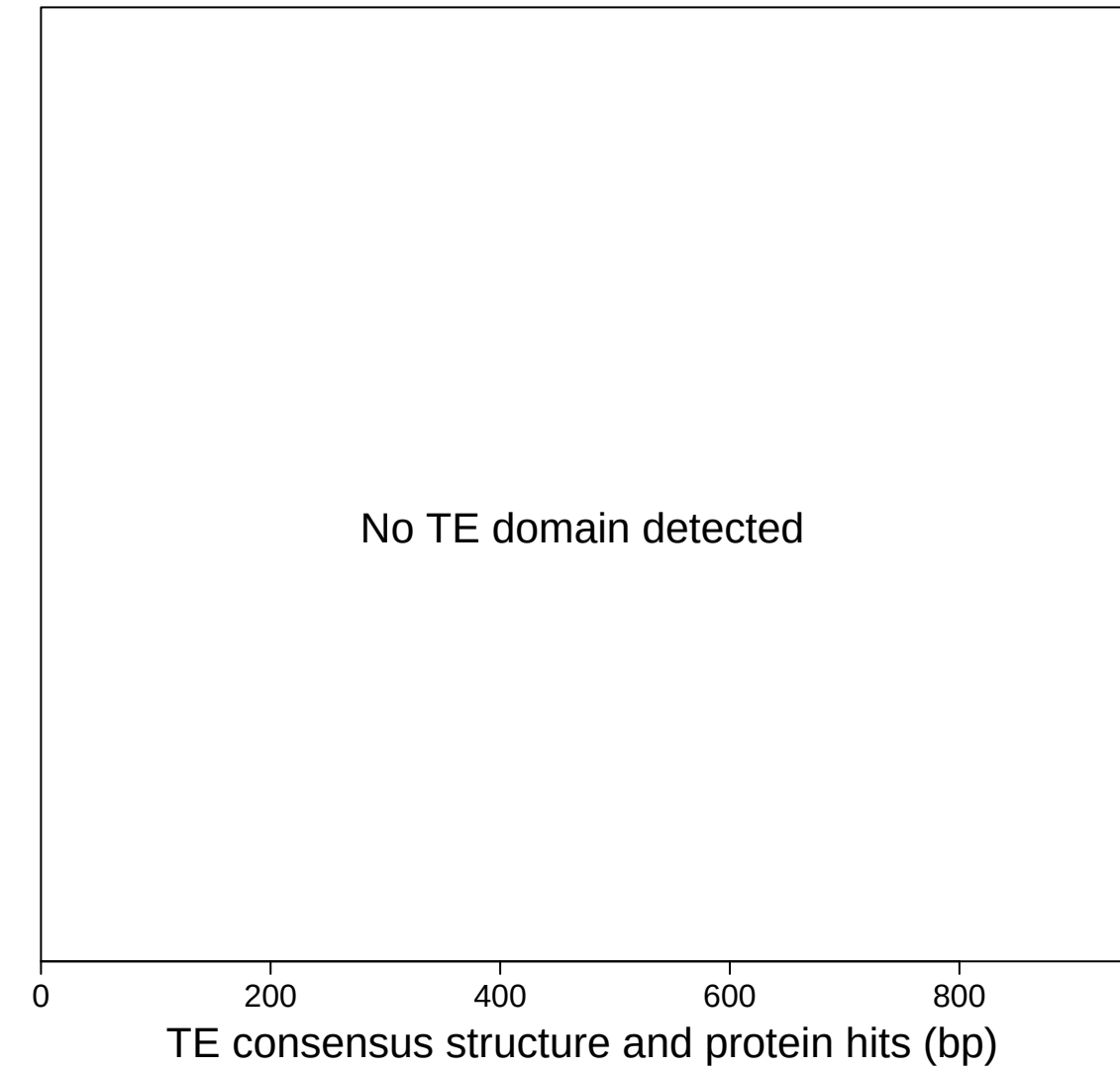
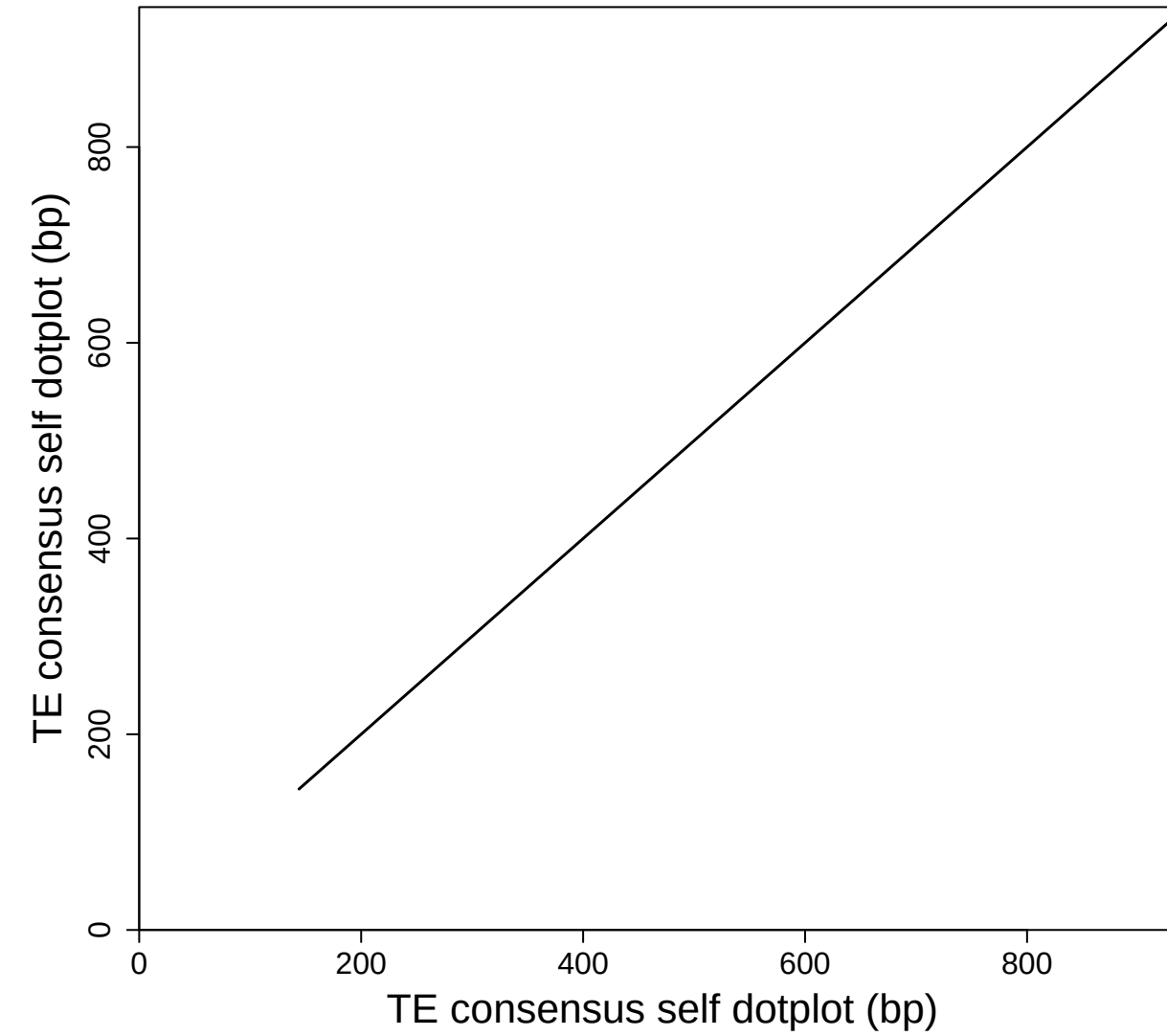
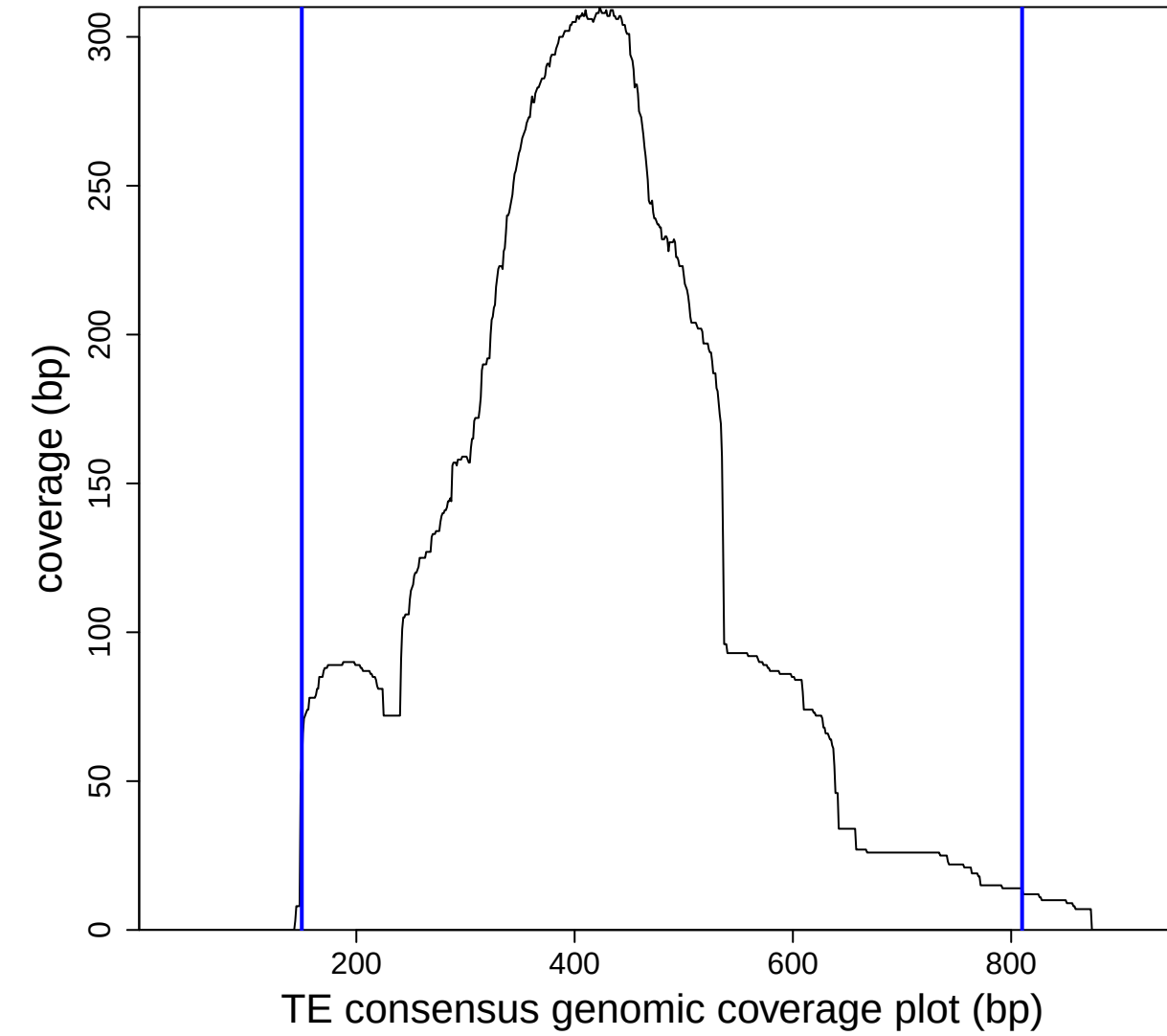
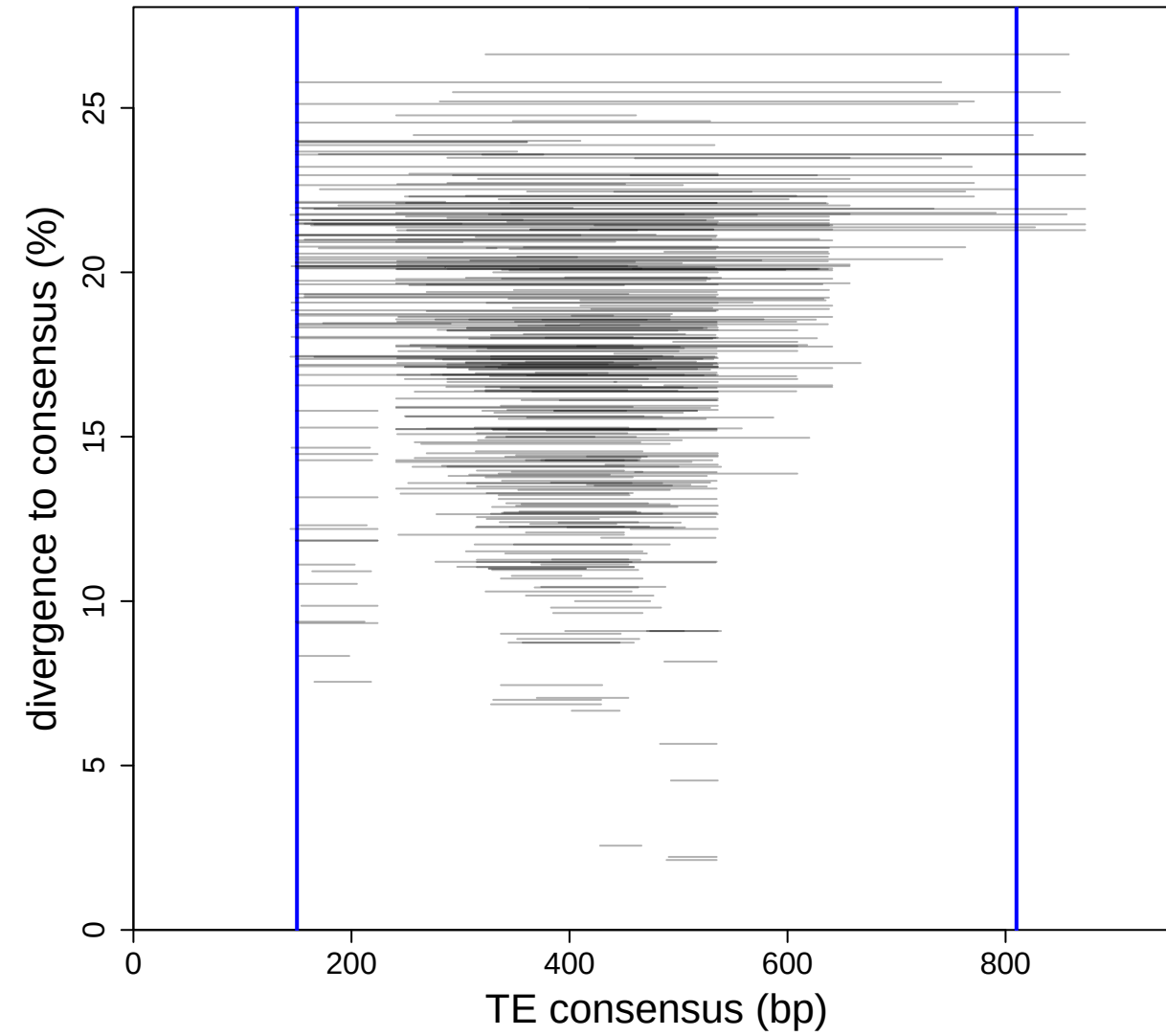
a.b.bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.f
size: 631bp; fragments: 803; full length: 48 (≥ 567.9 bp)

After TEtrimmer 631 bp



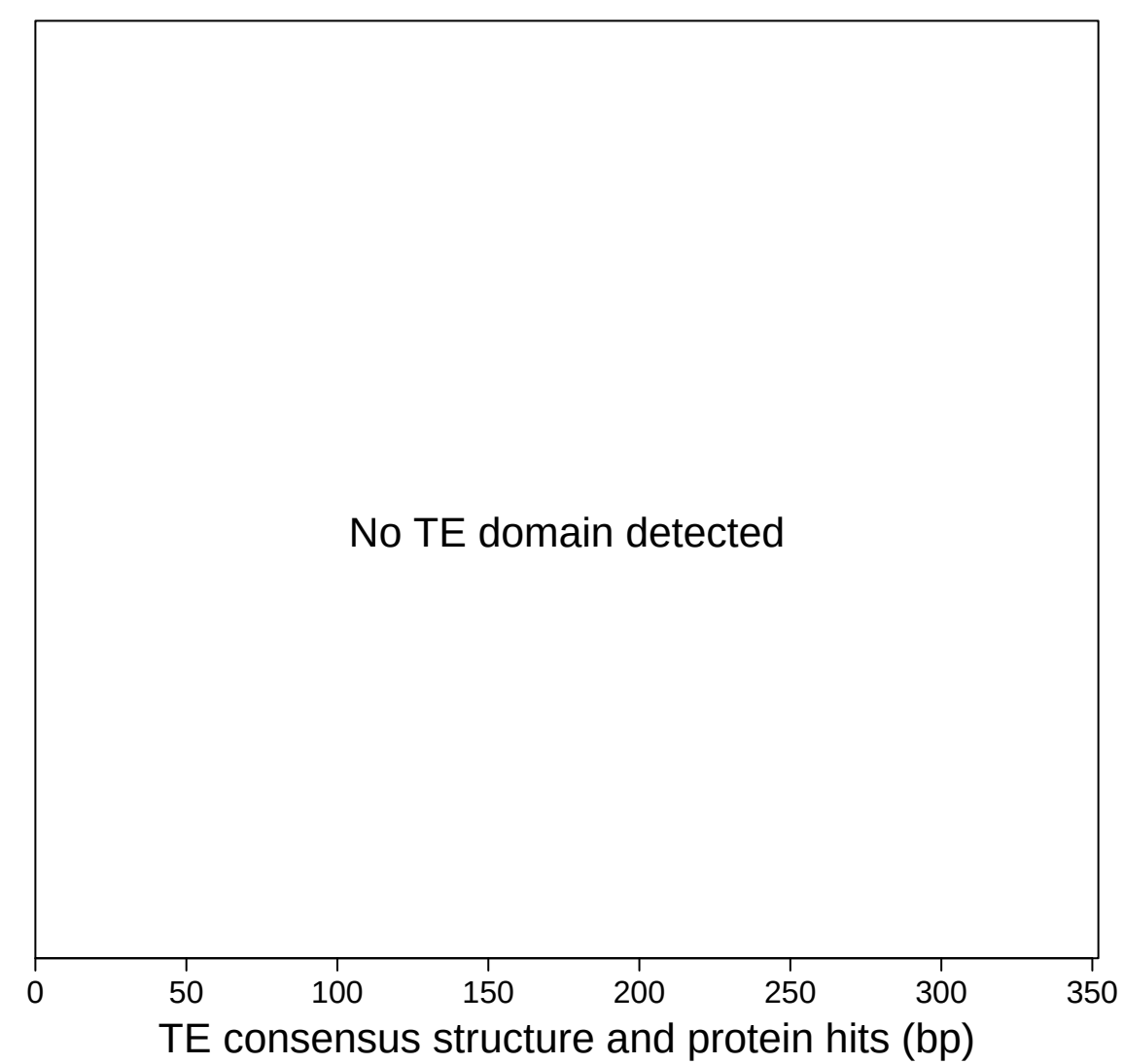
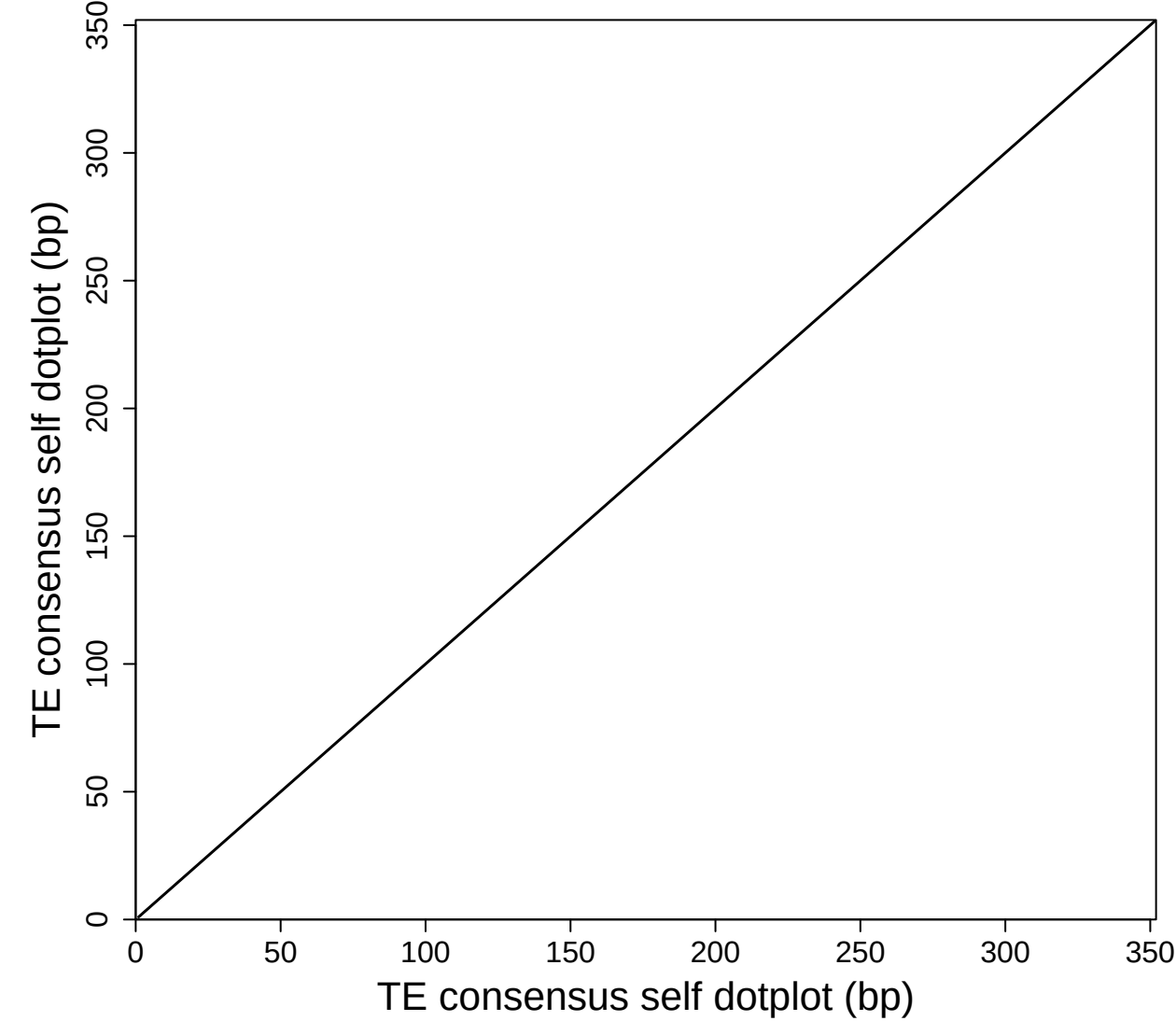
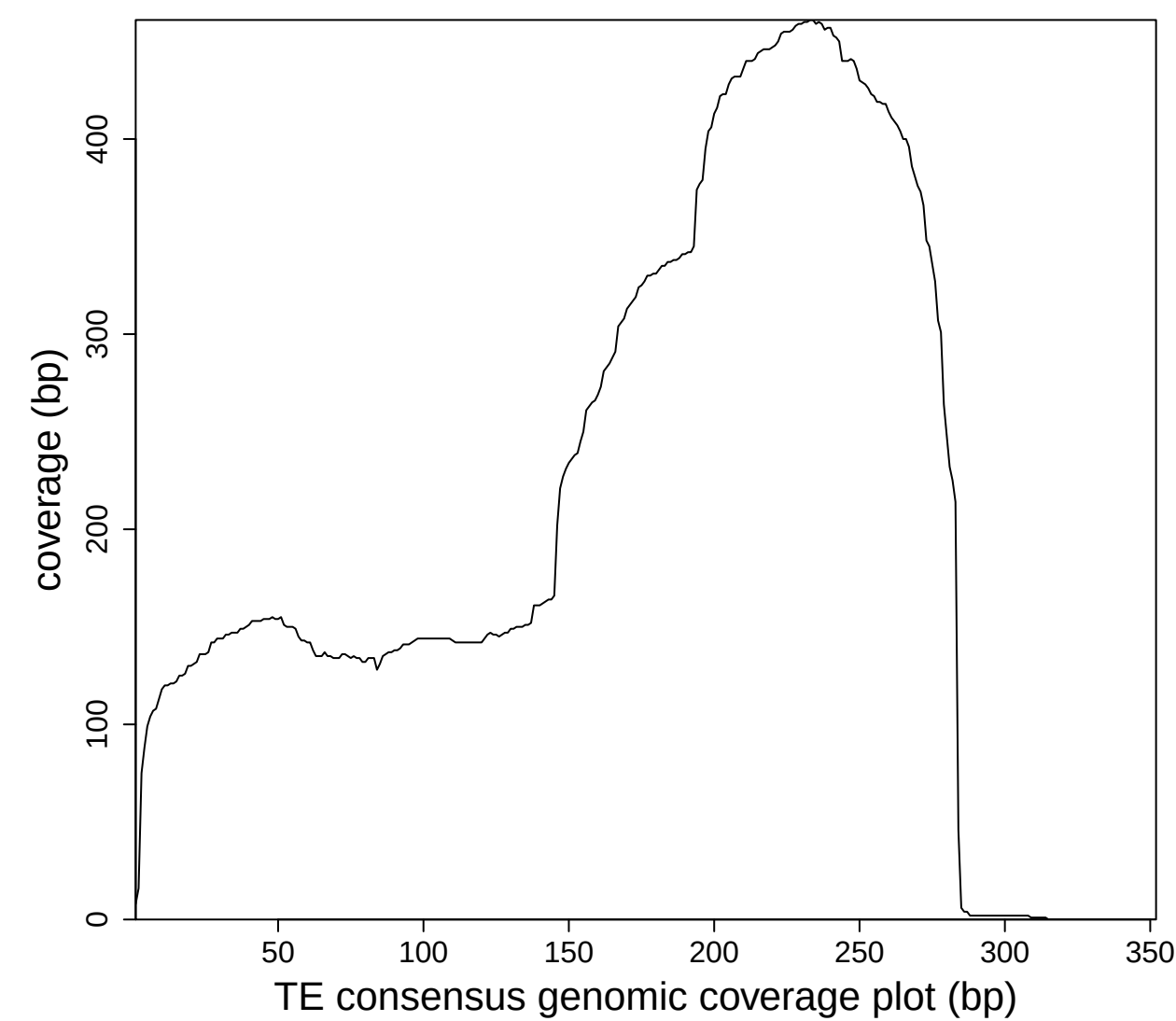
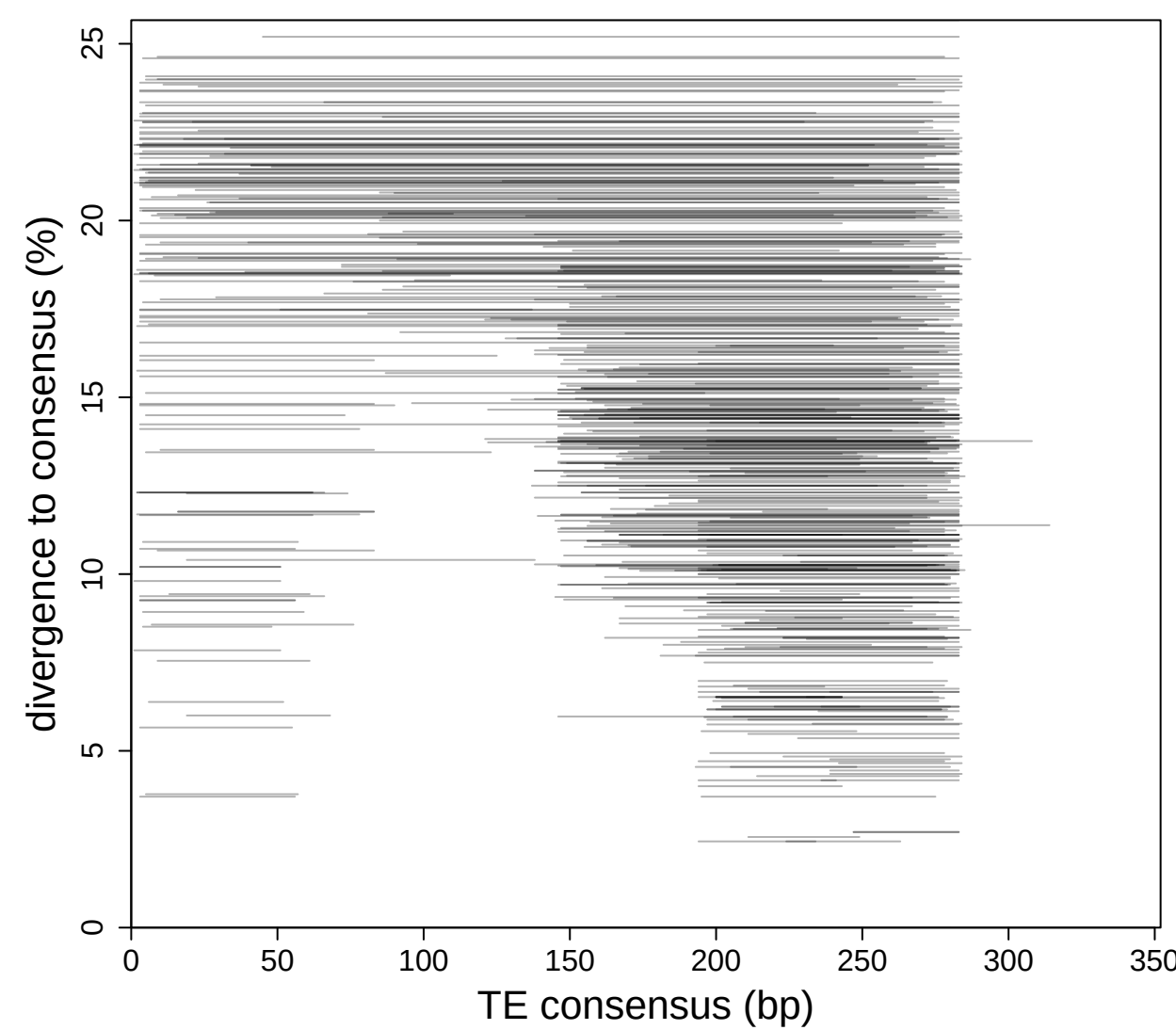
3.fasta.b.bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.f
size: 960bp; fragments: 371; full length: 0 (>=864bp)

After TEtrimmer Extended plot Blue lines are boundaries

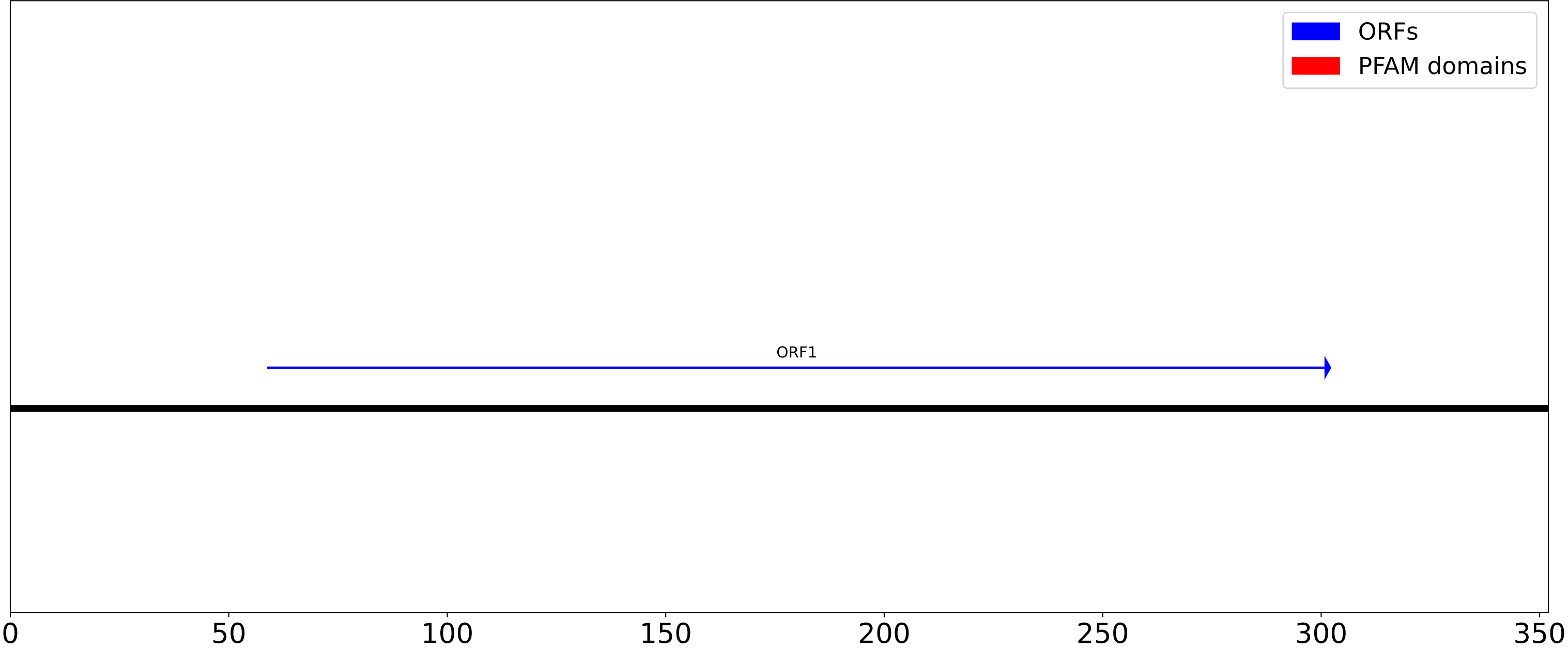


Before TEtrimmer 352 bp

TE: rnd_4_family_1553
size: 352bp; fragments: 516; full length: 0 (≥ 316.8 bp)



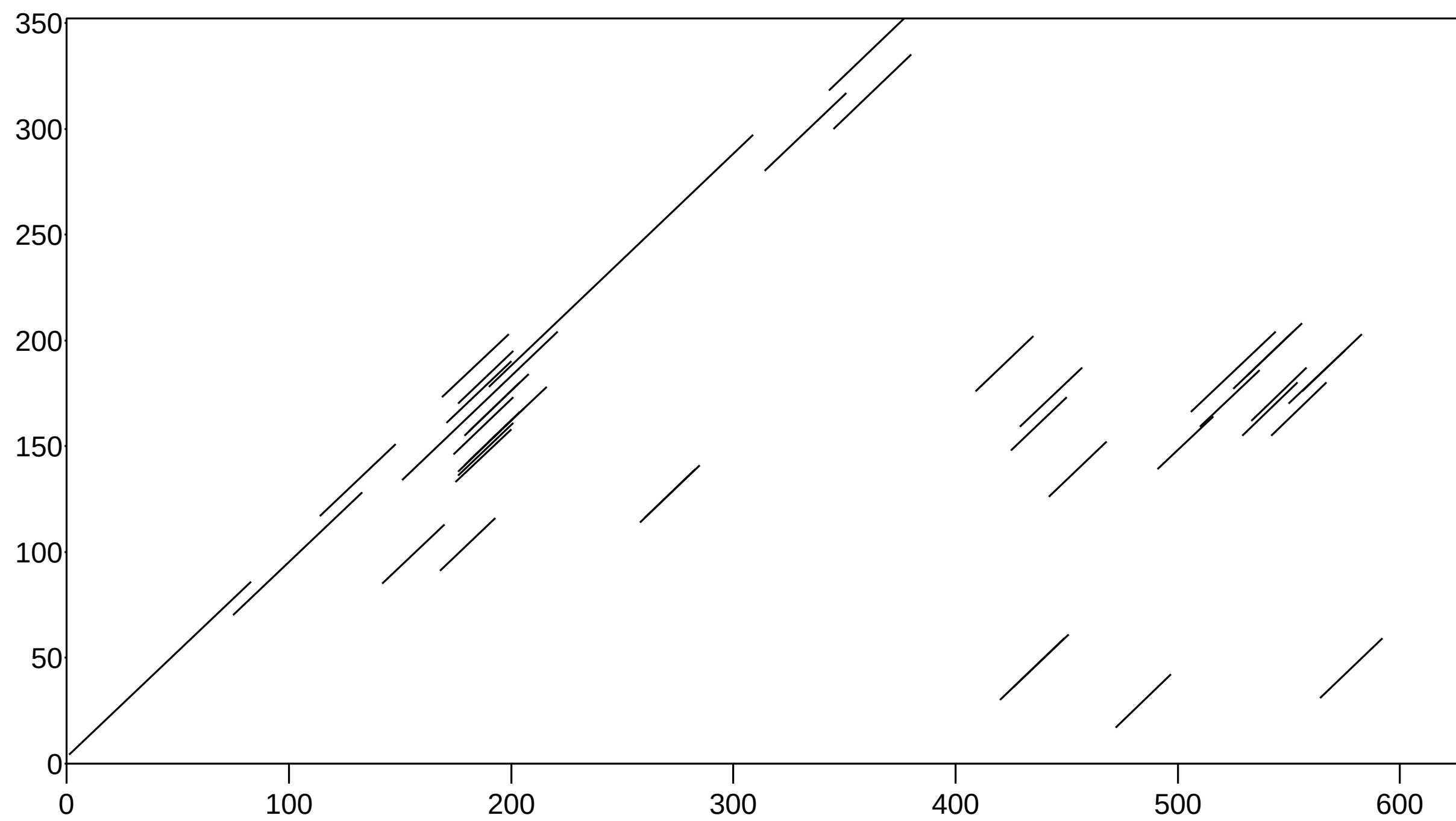
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)