

Start crop Point

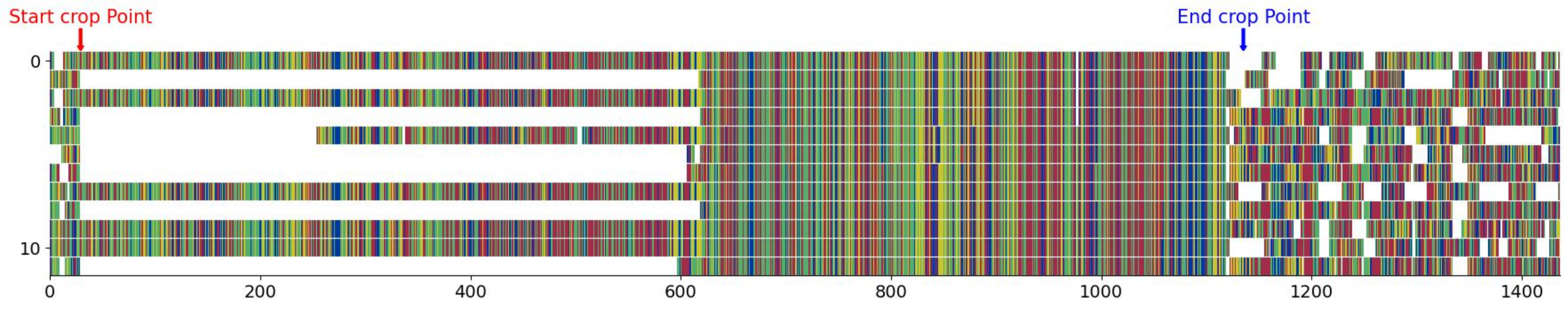


End crop Point



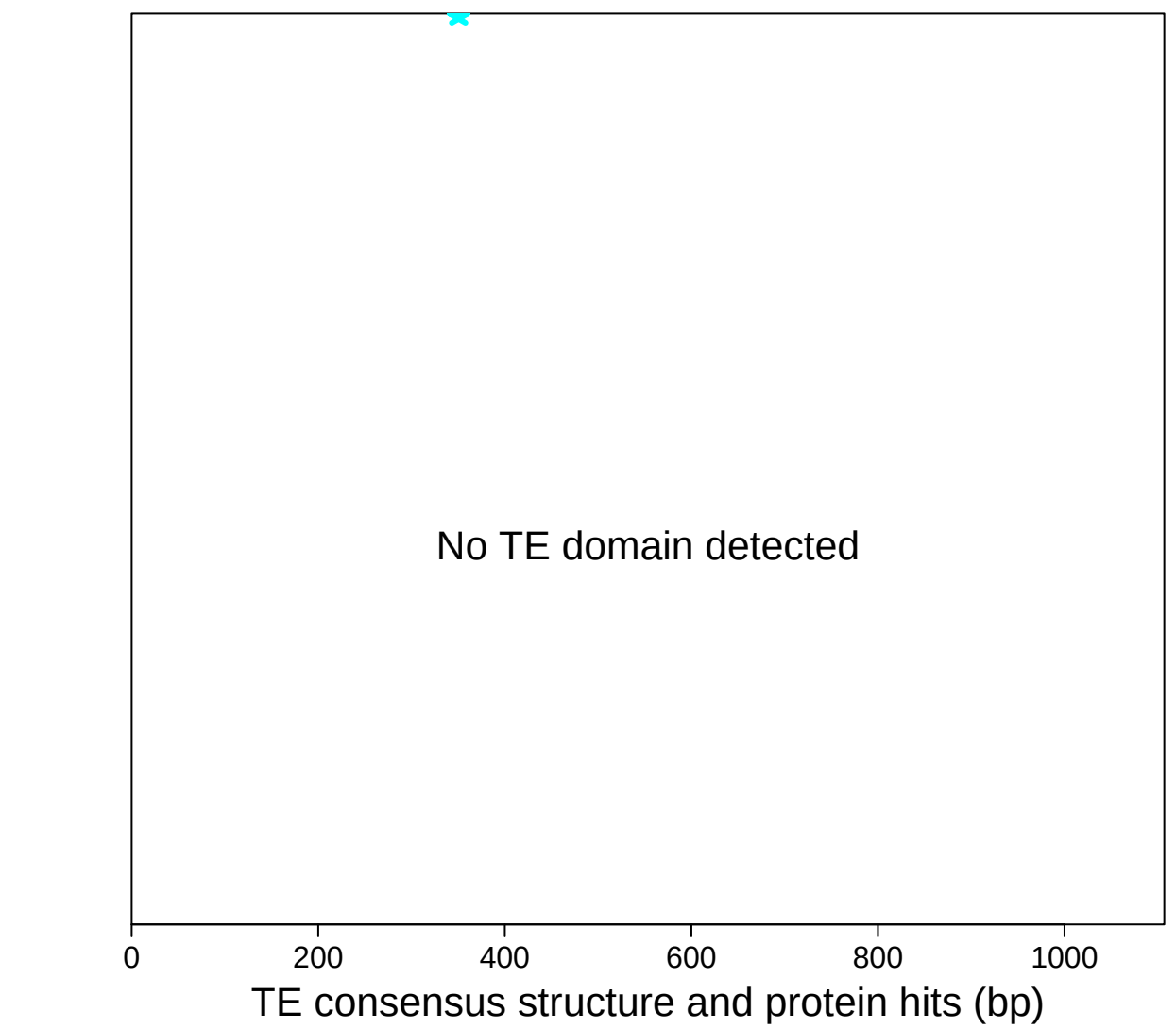
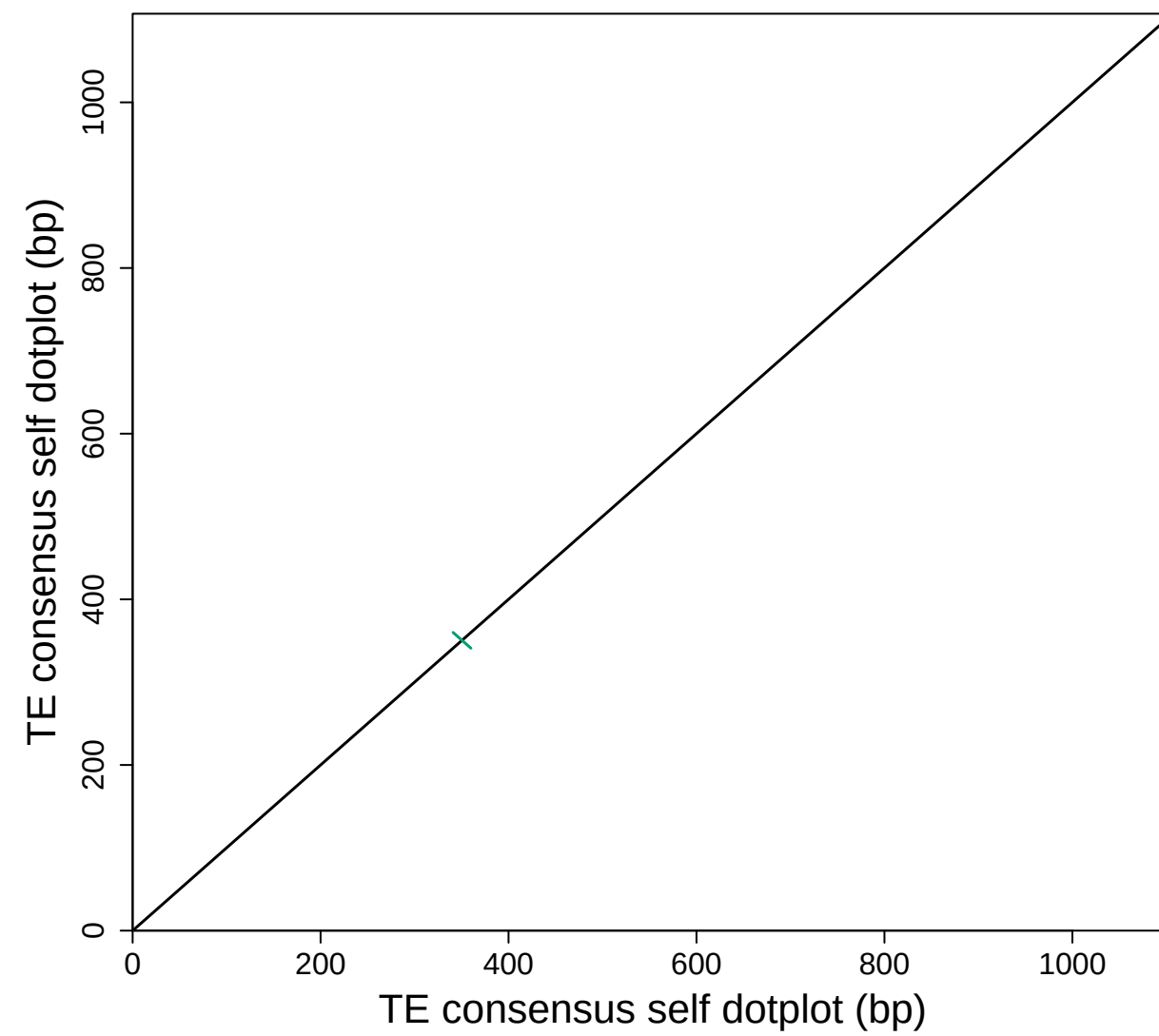
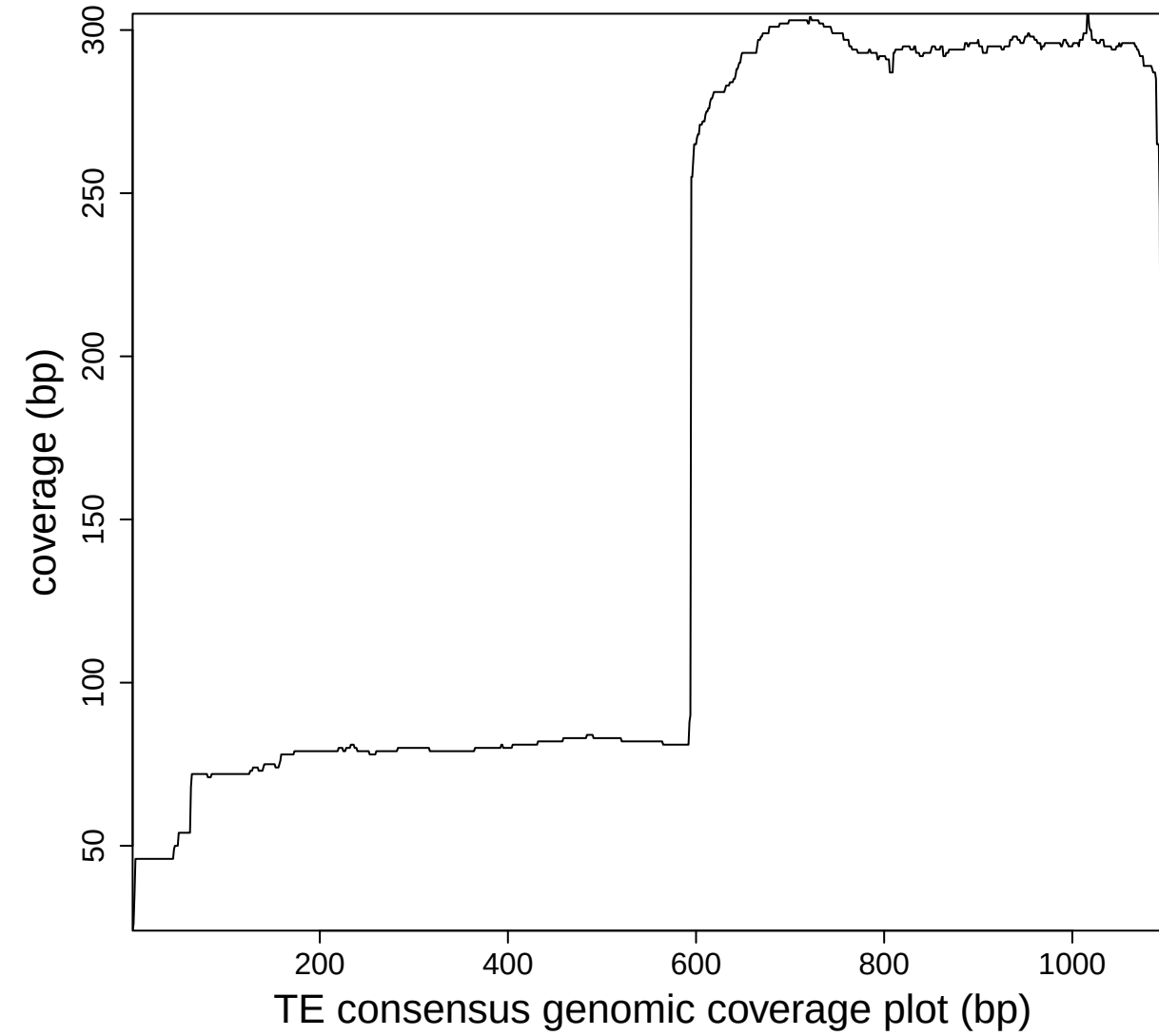
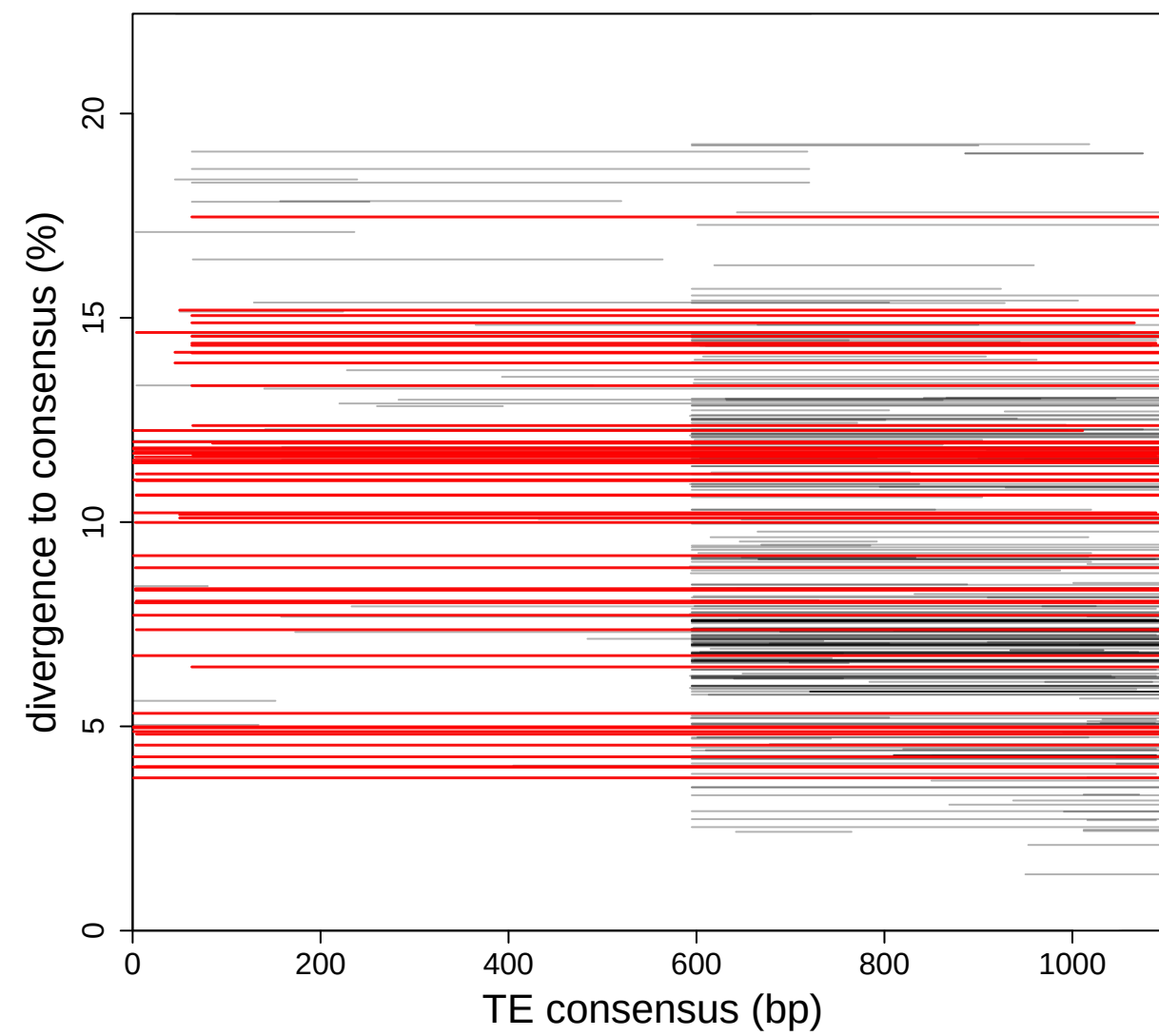
MSA length = 1107





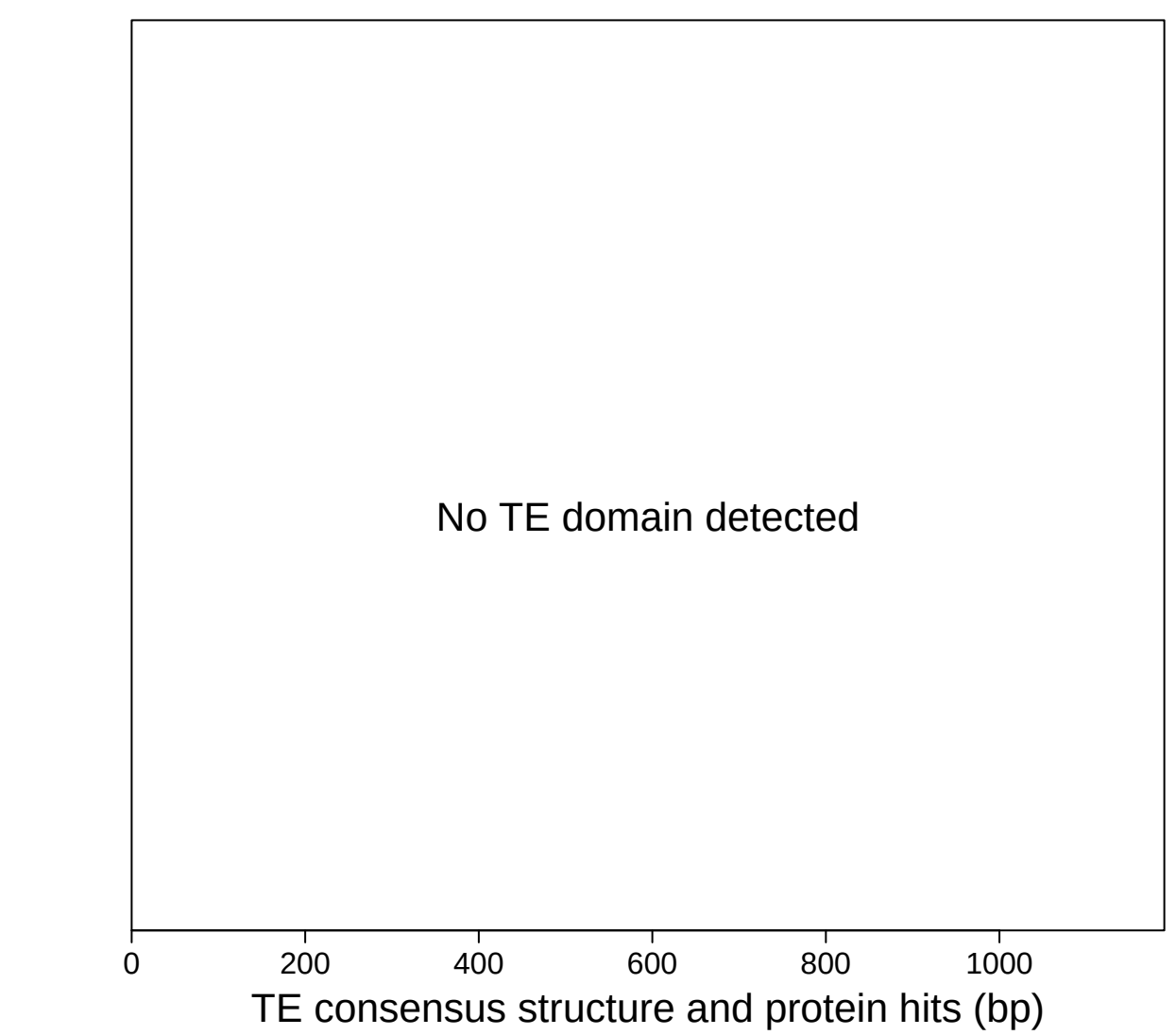
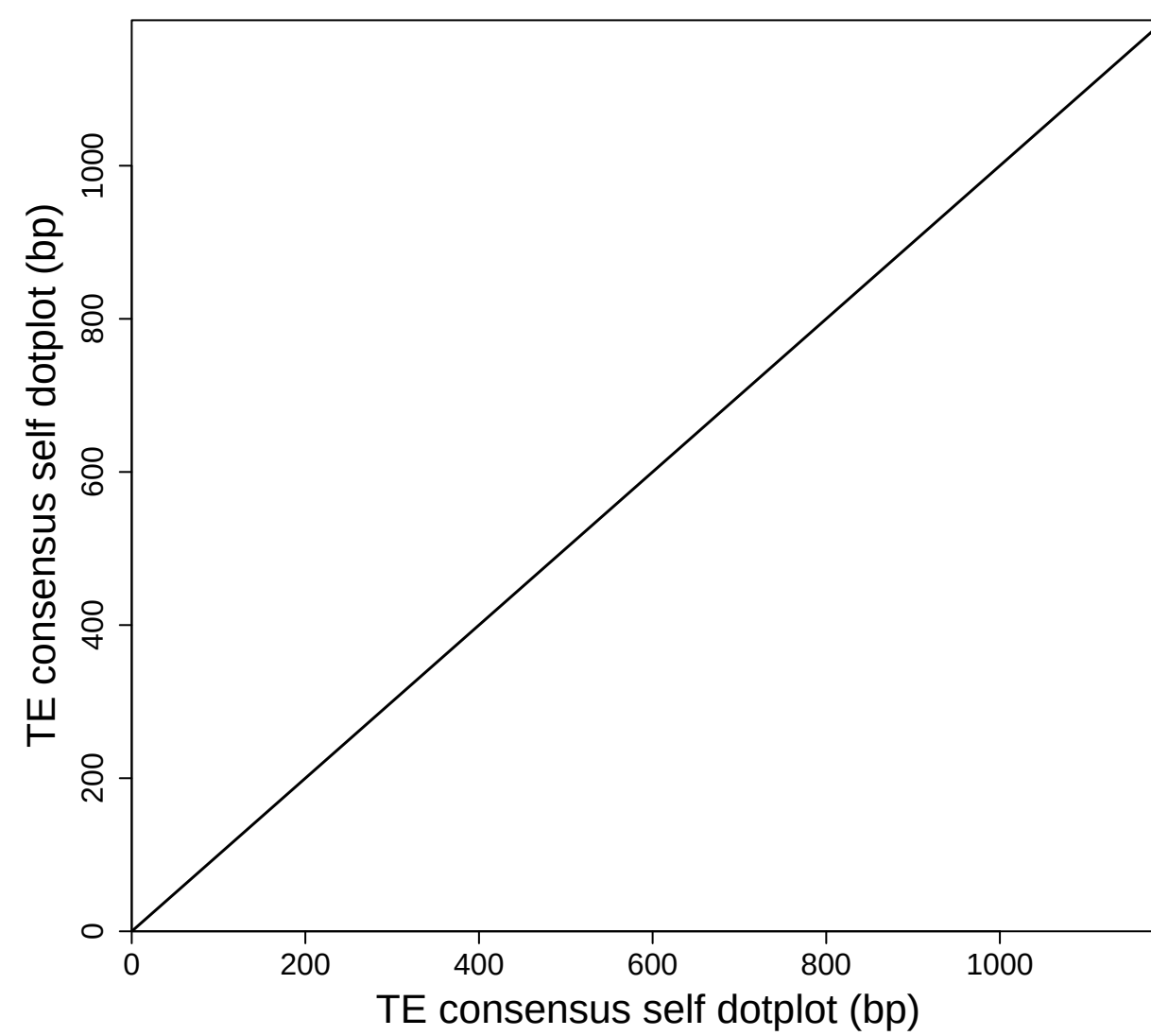
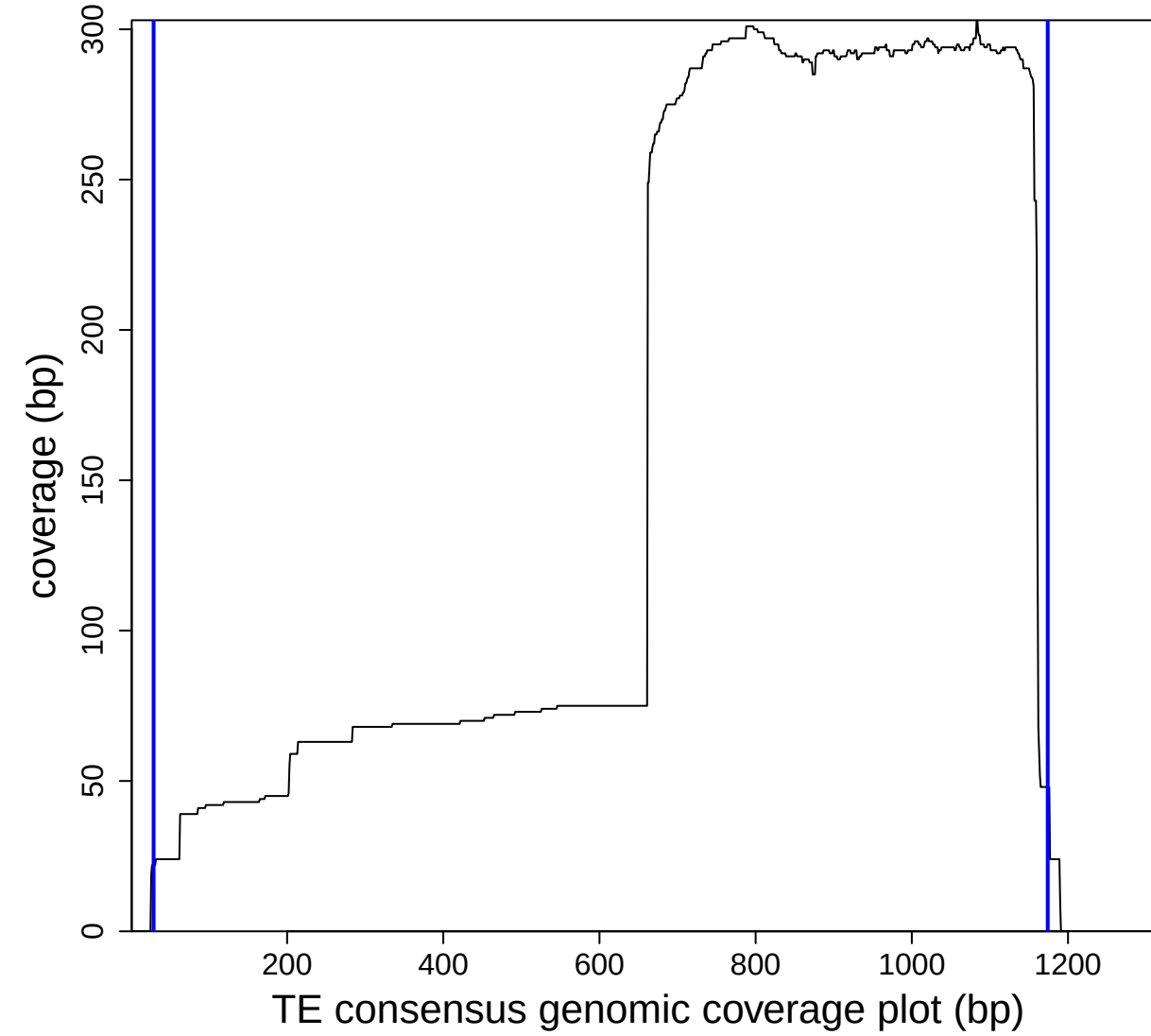
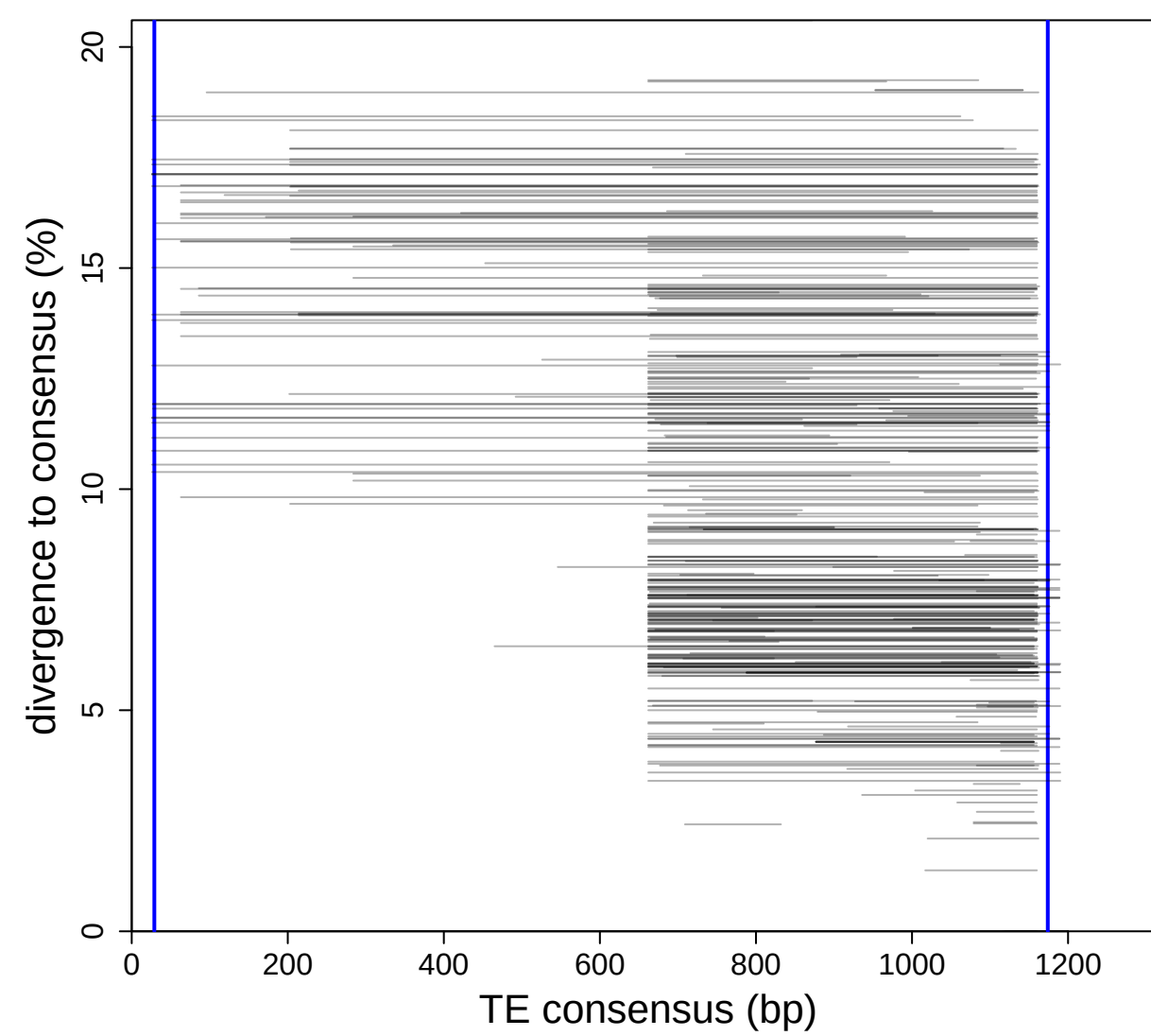
b.bed uf.bed g 3.bed fm 1.bed 0 0 bcln.fa aln.fa cl.fa gs.fa
size: 1107bp; fragments: 374; full length: 57 (>=996.3bp)

After TEtrimmer 1107 bp



.fasta.b.bed_uf.bed_g_3.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa
size: 1324bp; fragments: 356; full length: 0 (>=1191.6bp)

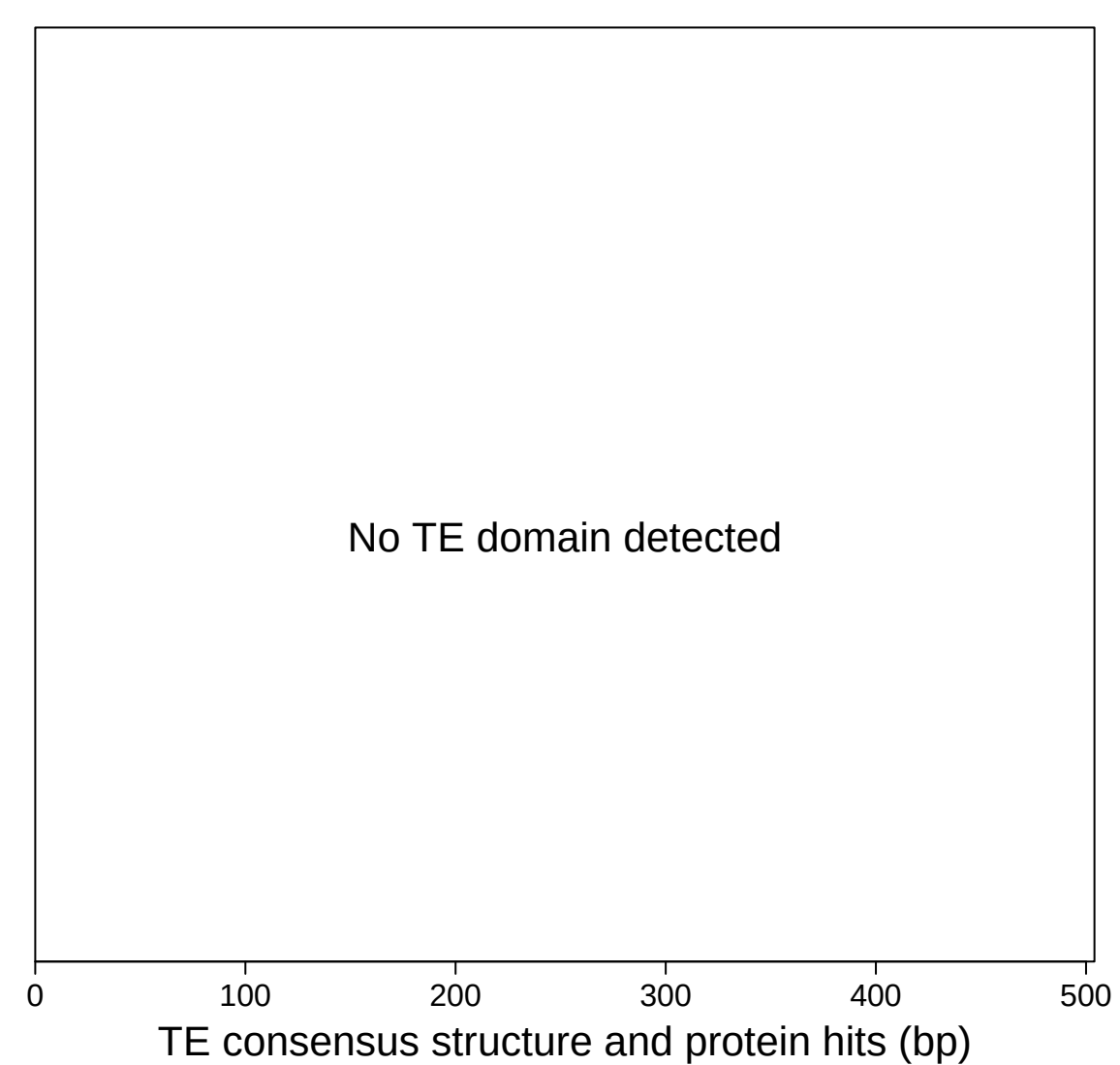
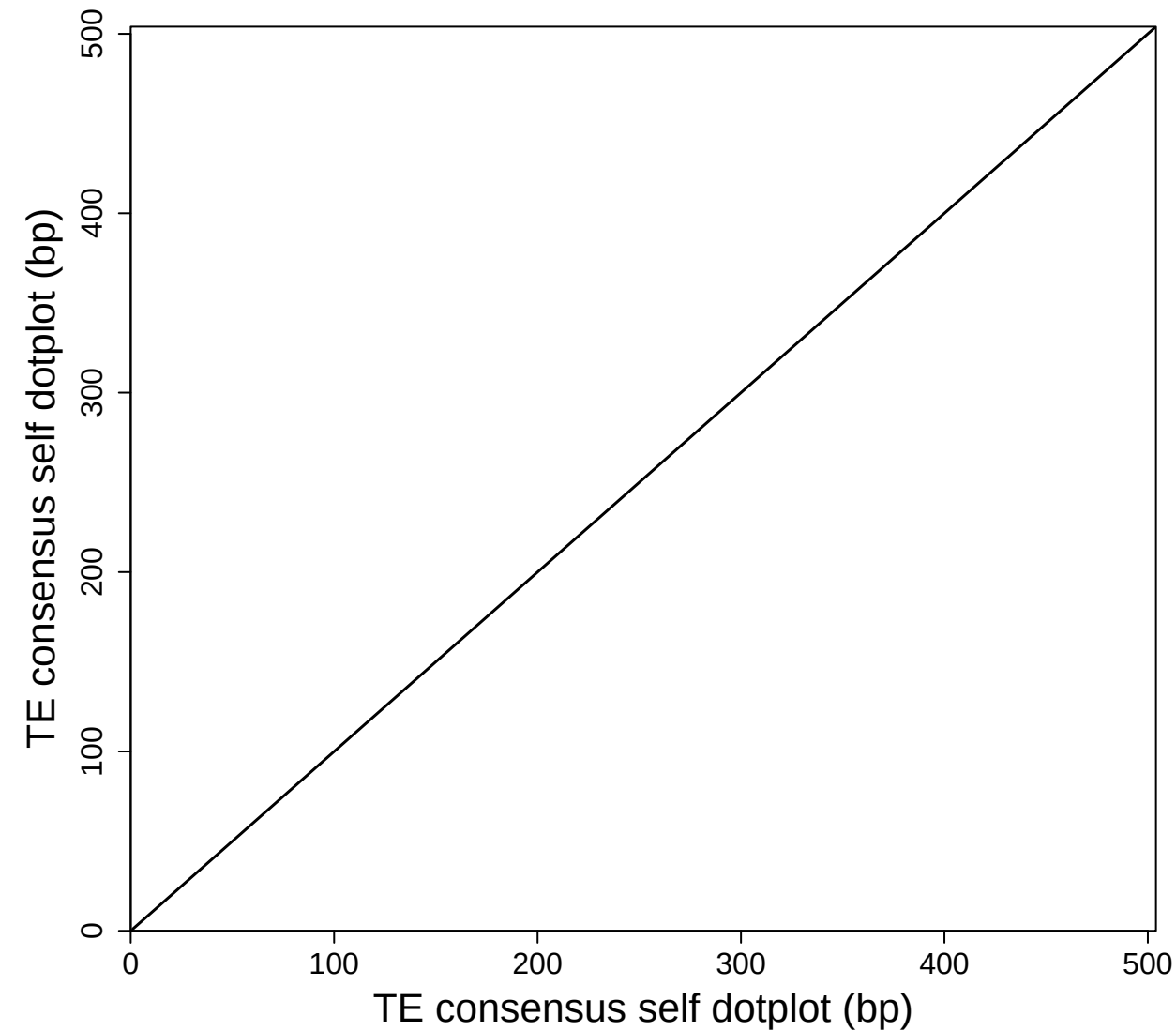
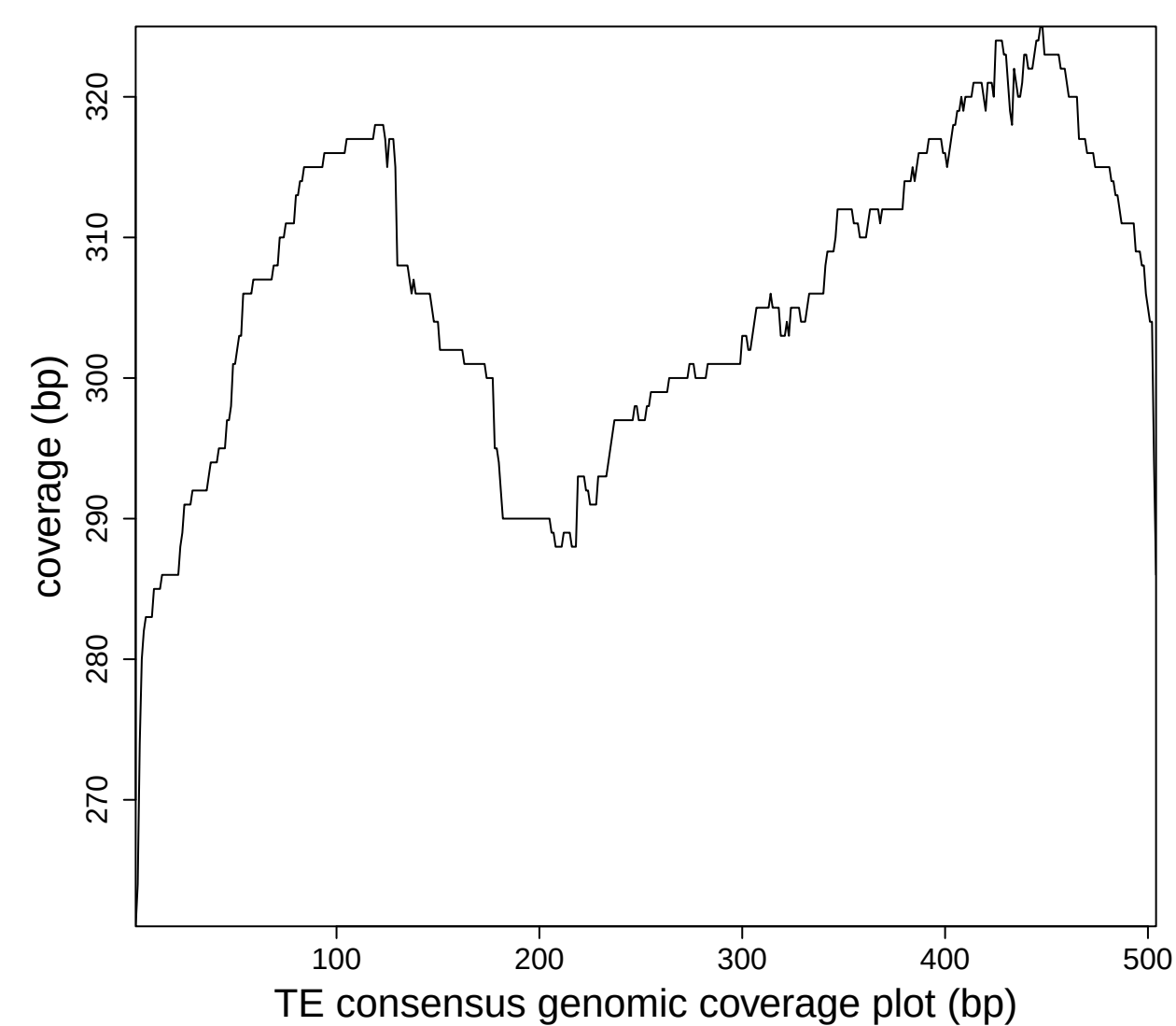
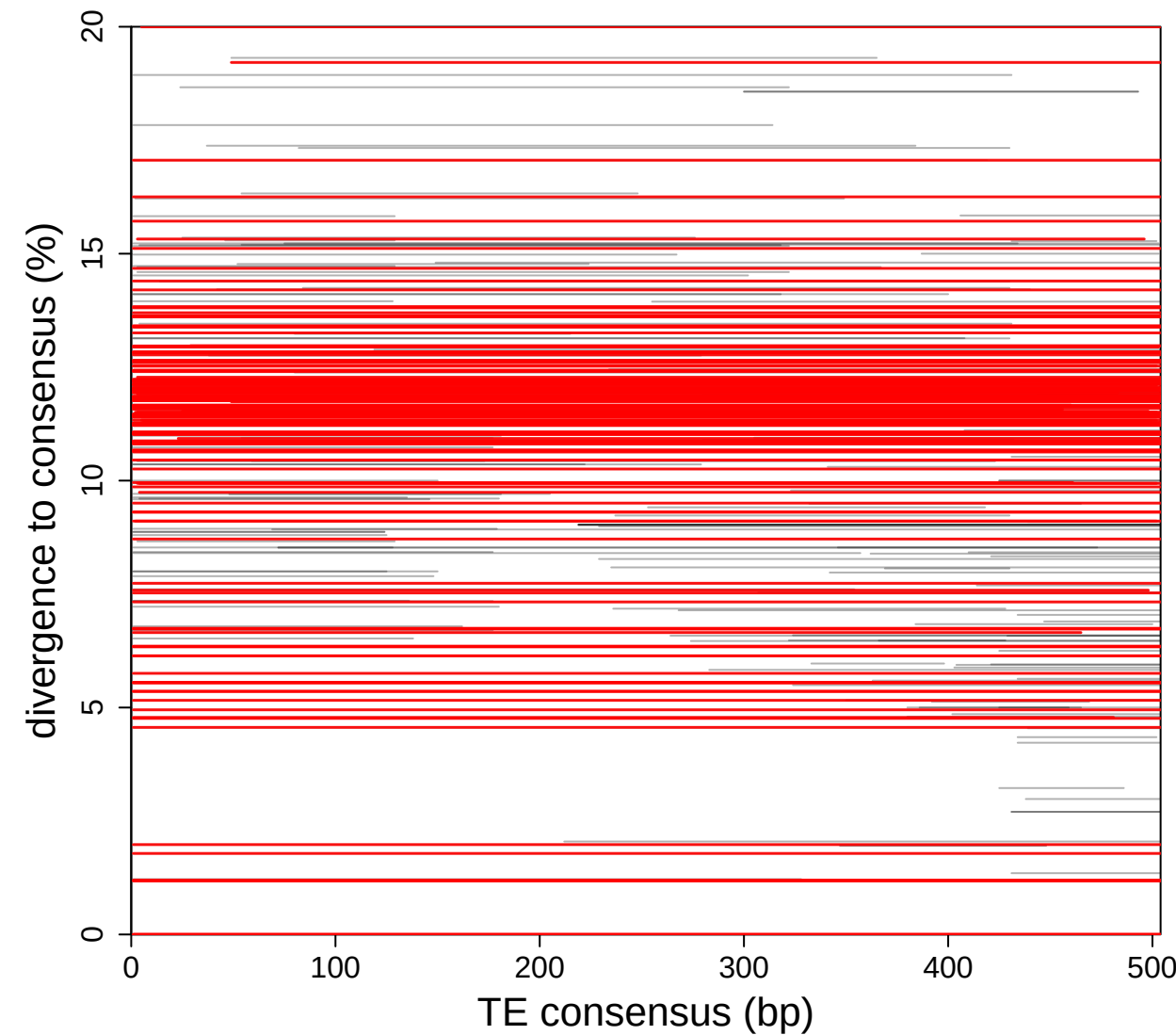
After TEtrimmer Extended plot Blue lines are boundaries



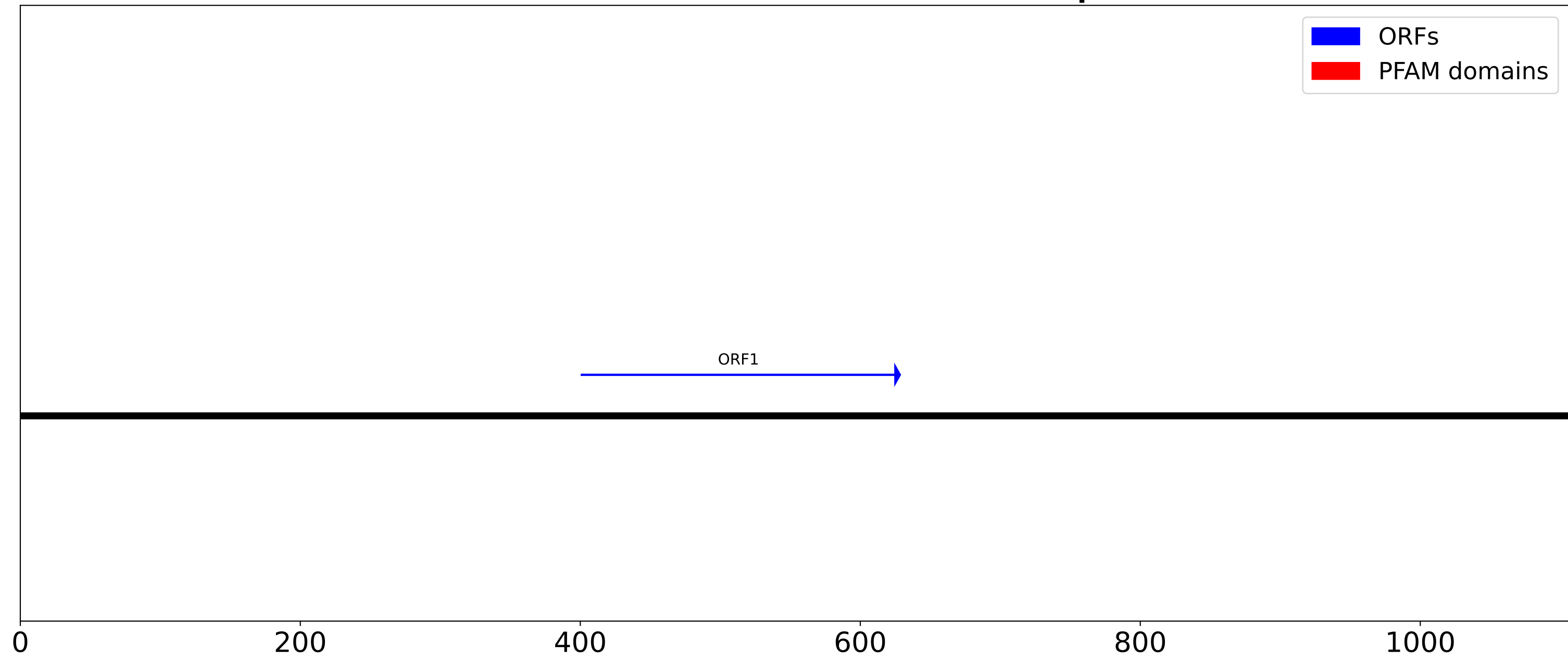
Before TEtrimmer 504 bp

TE: ltr_1_family_196

size: 504bp; fragments: 406; full length: 235 (≥ 453.6 bp)



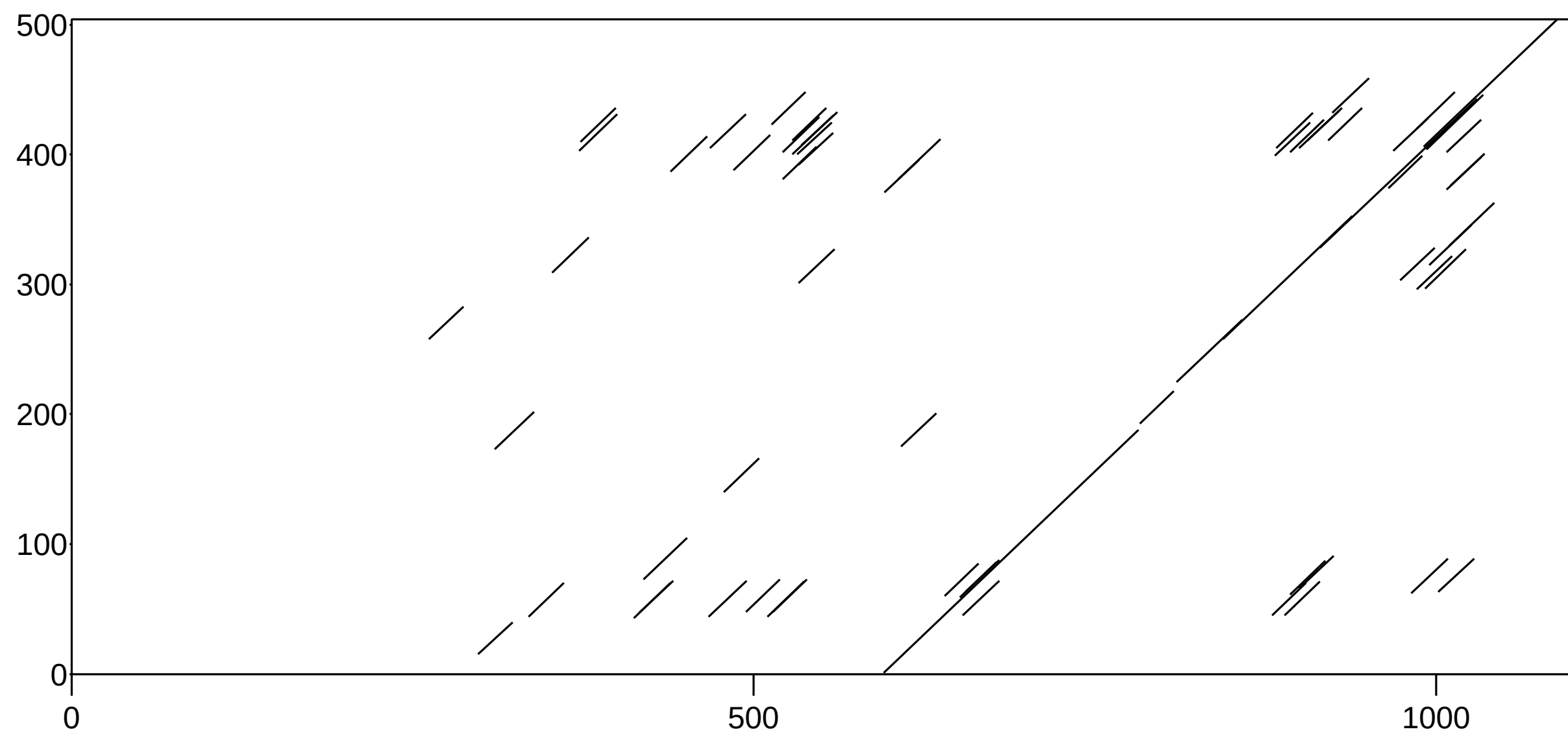
After TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)