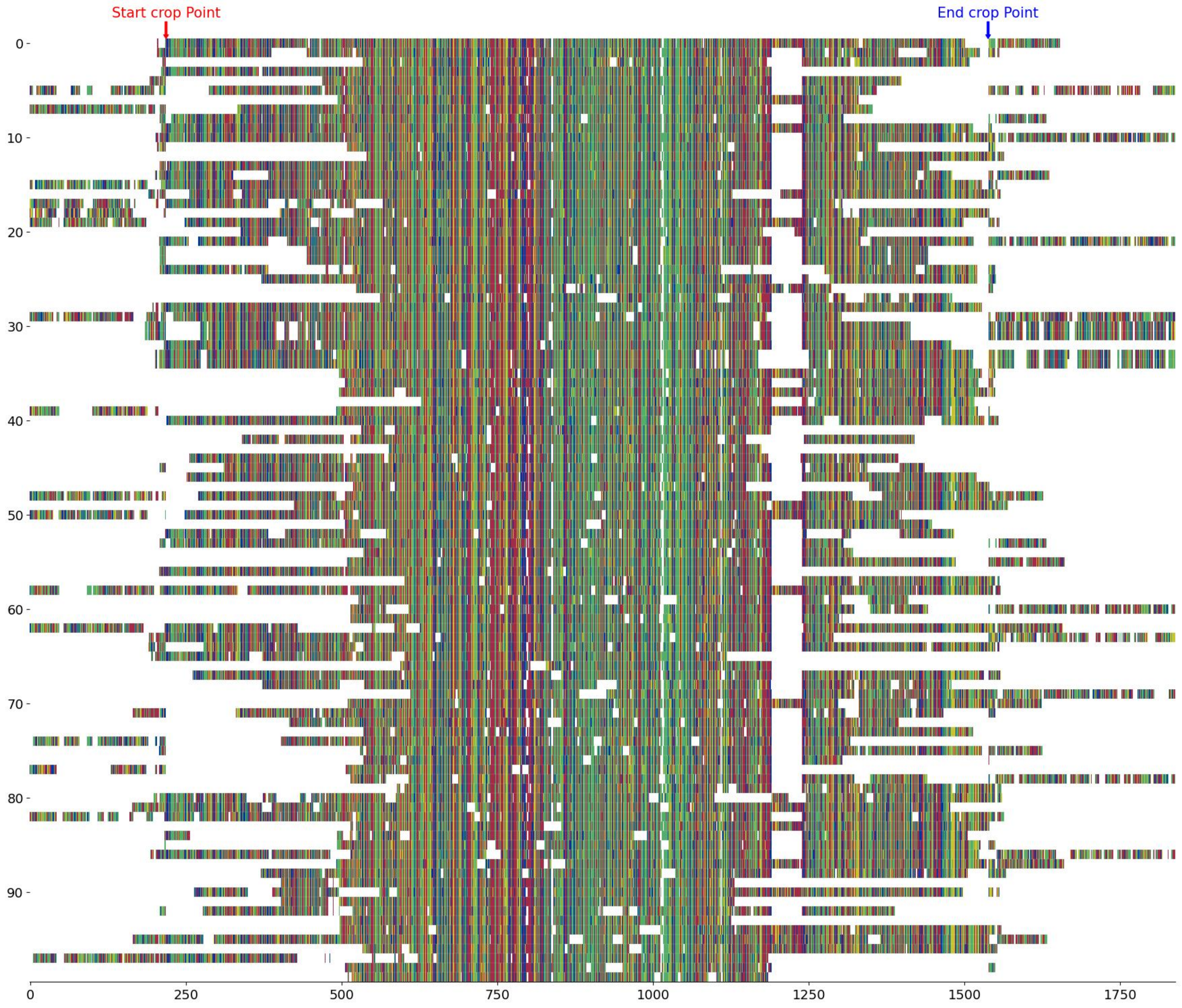




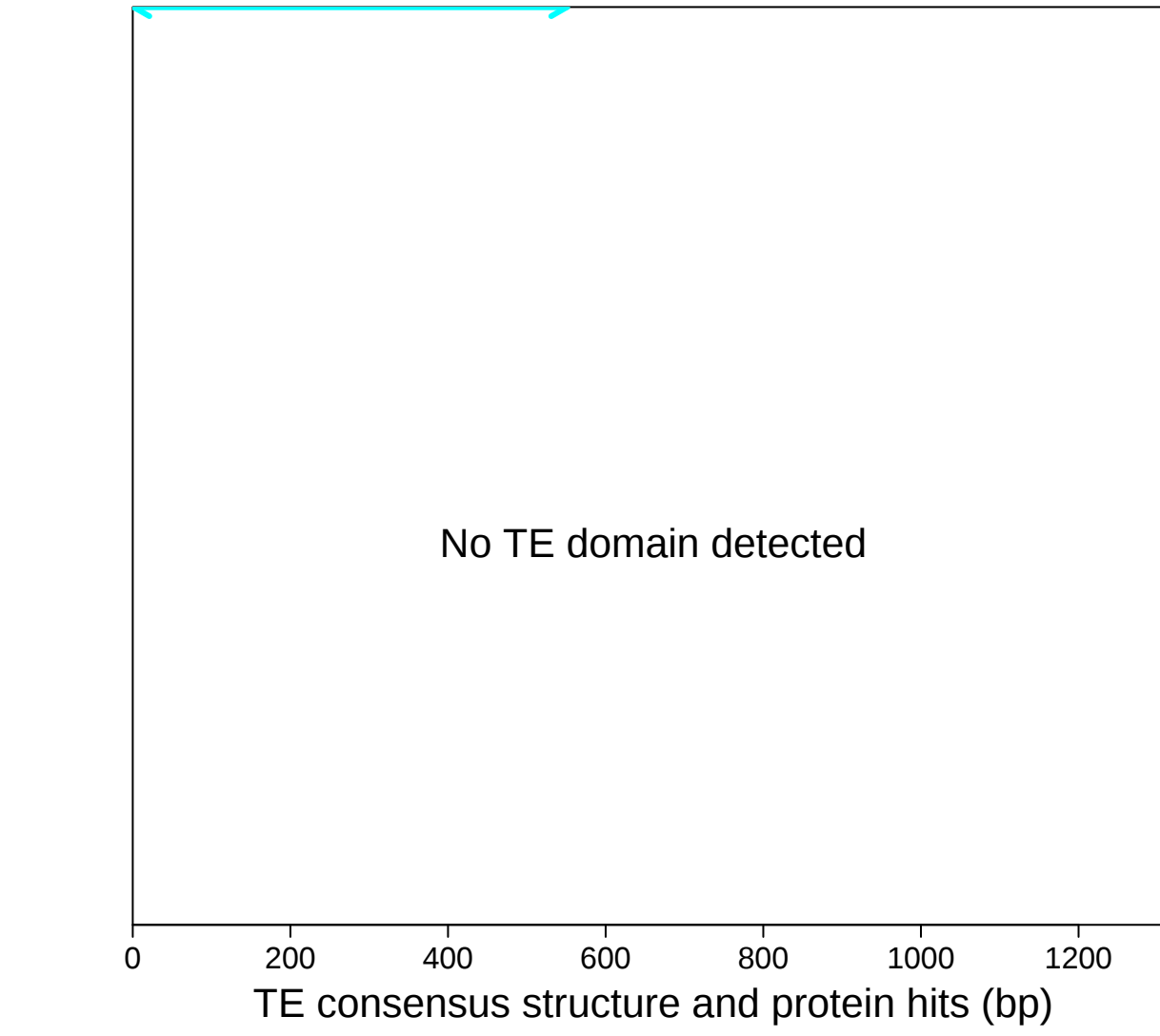
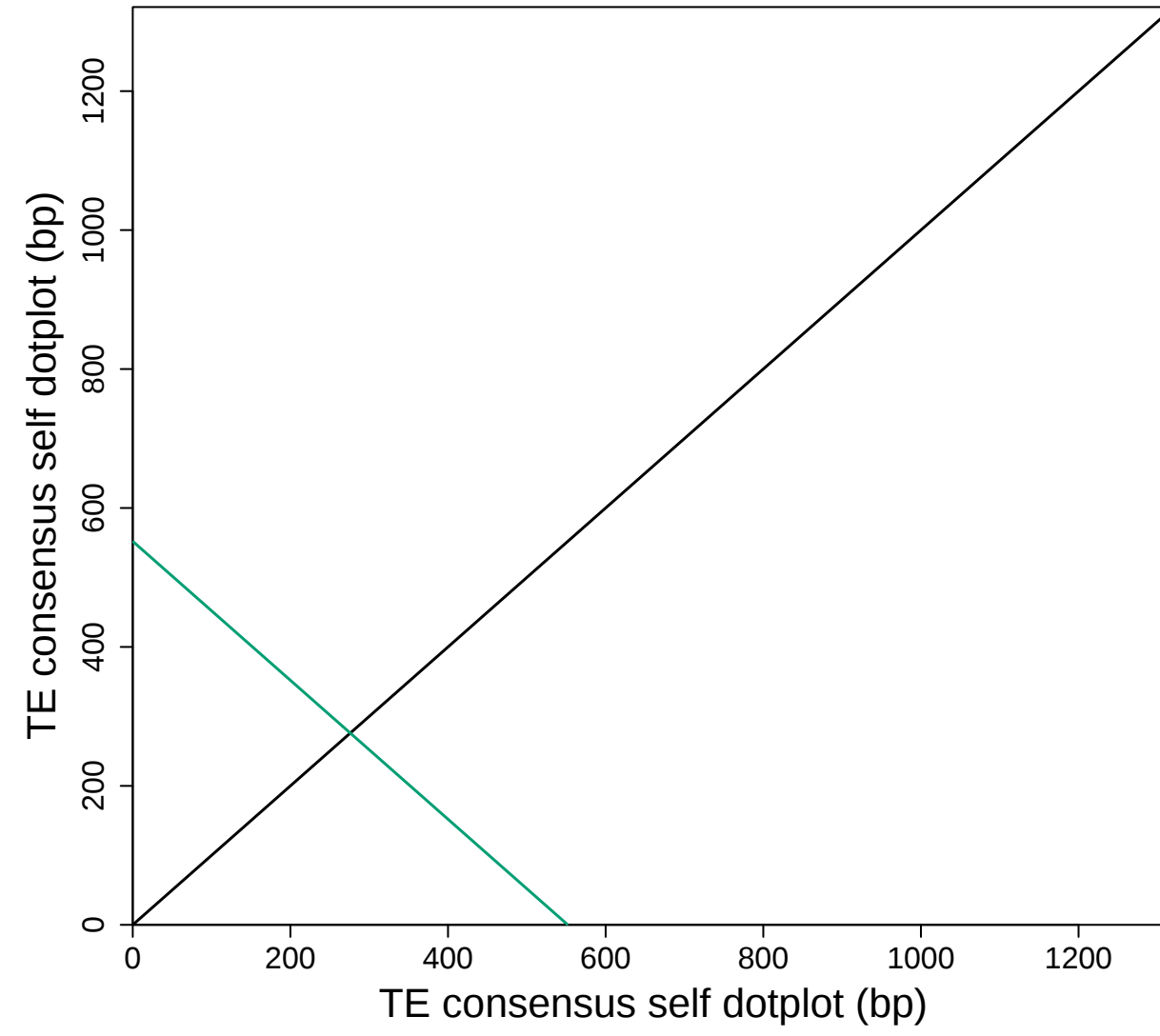
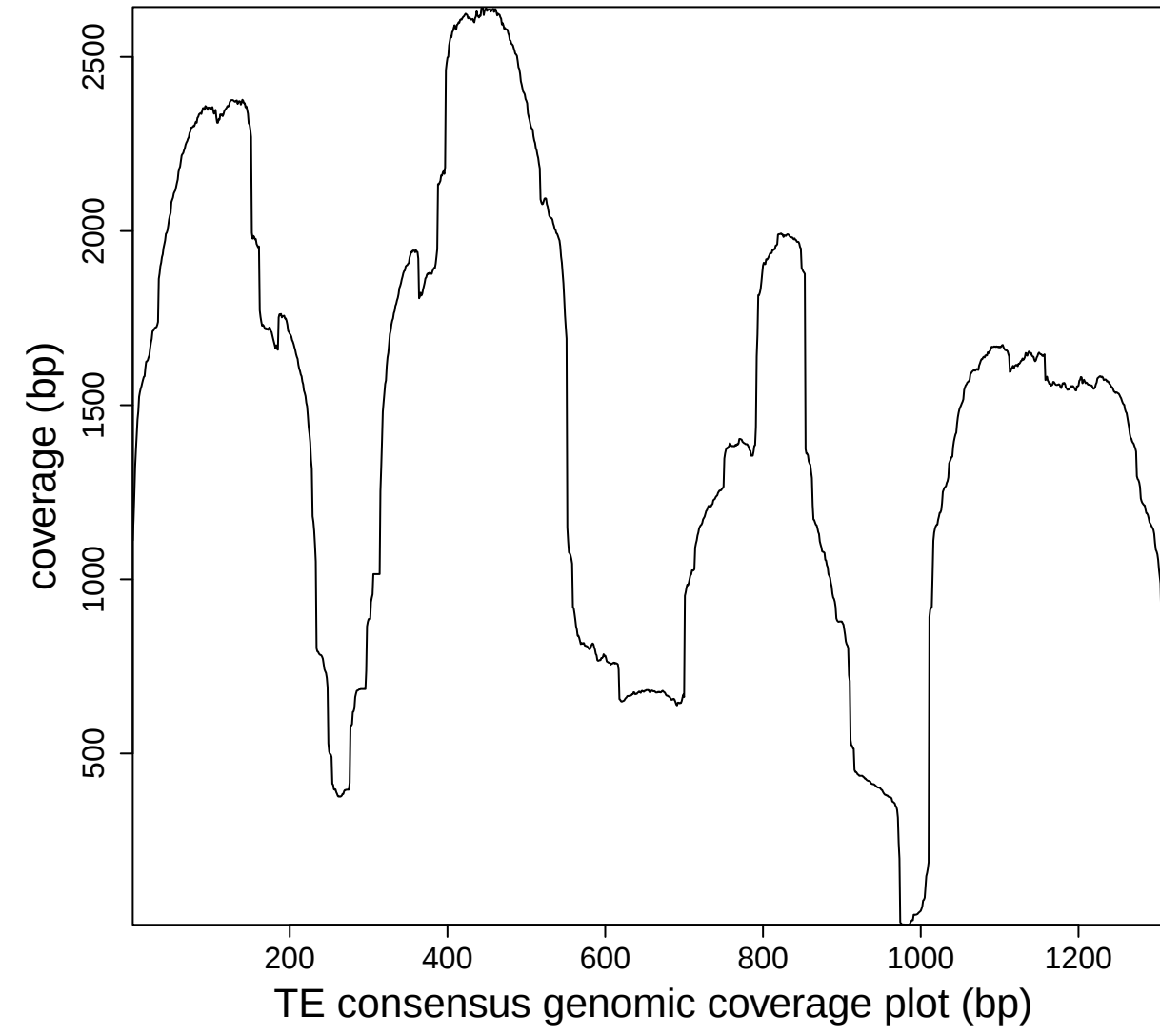
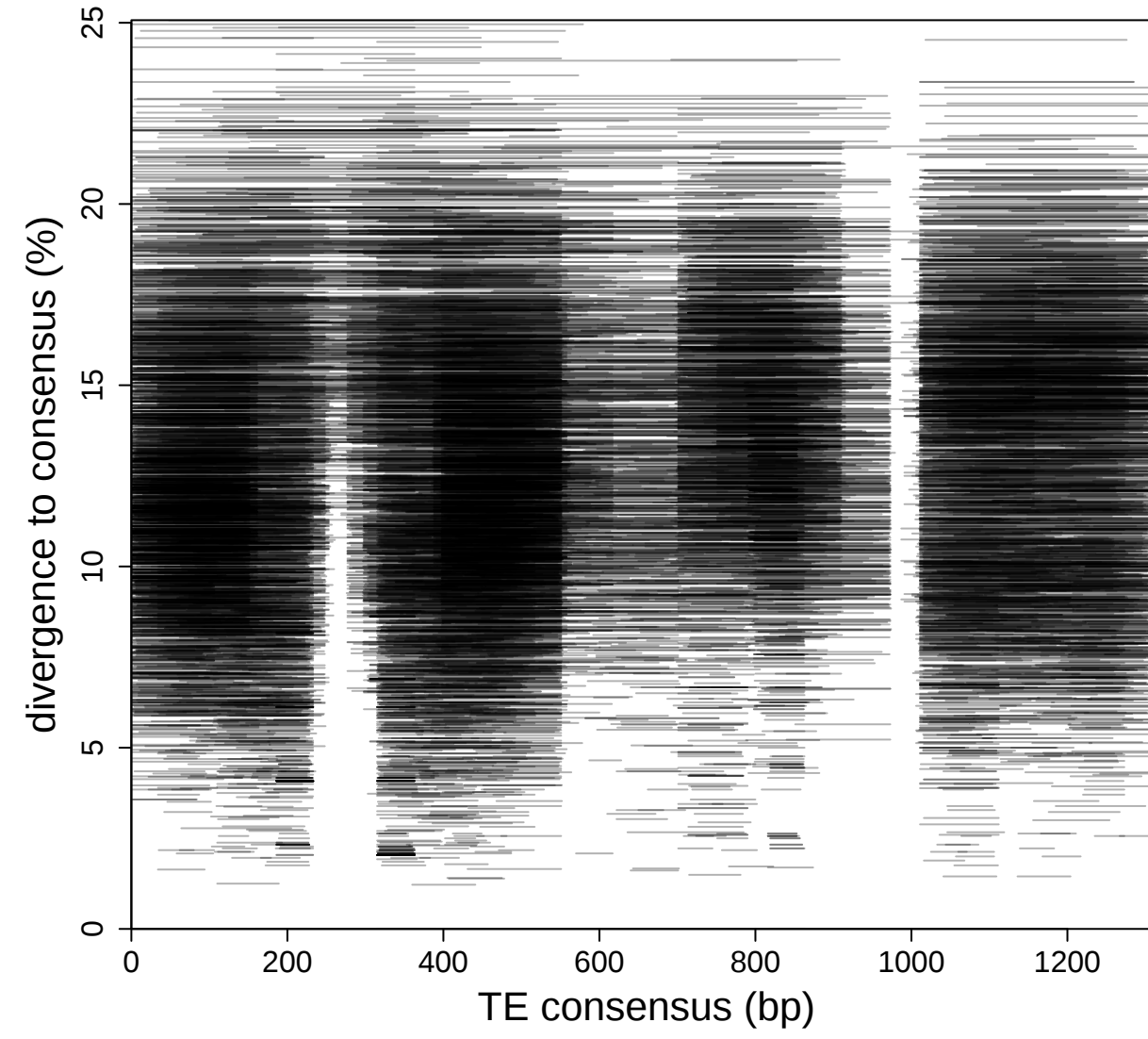
Start crop Point

End crop Point



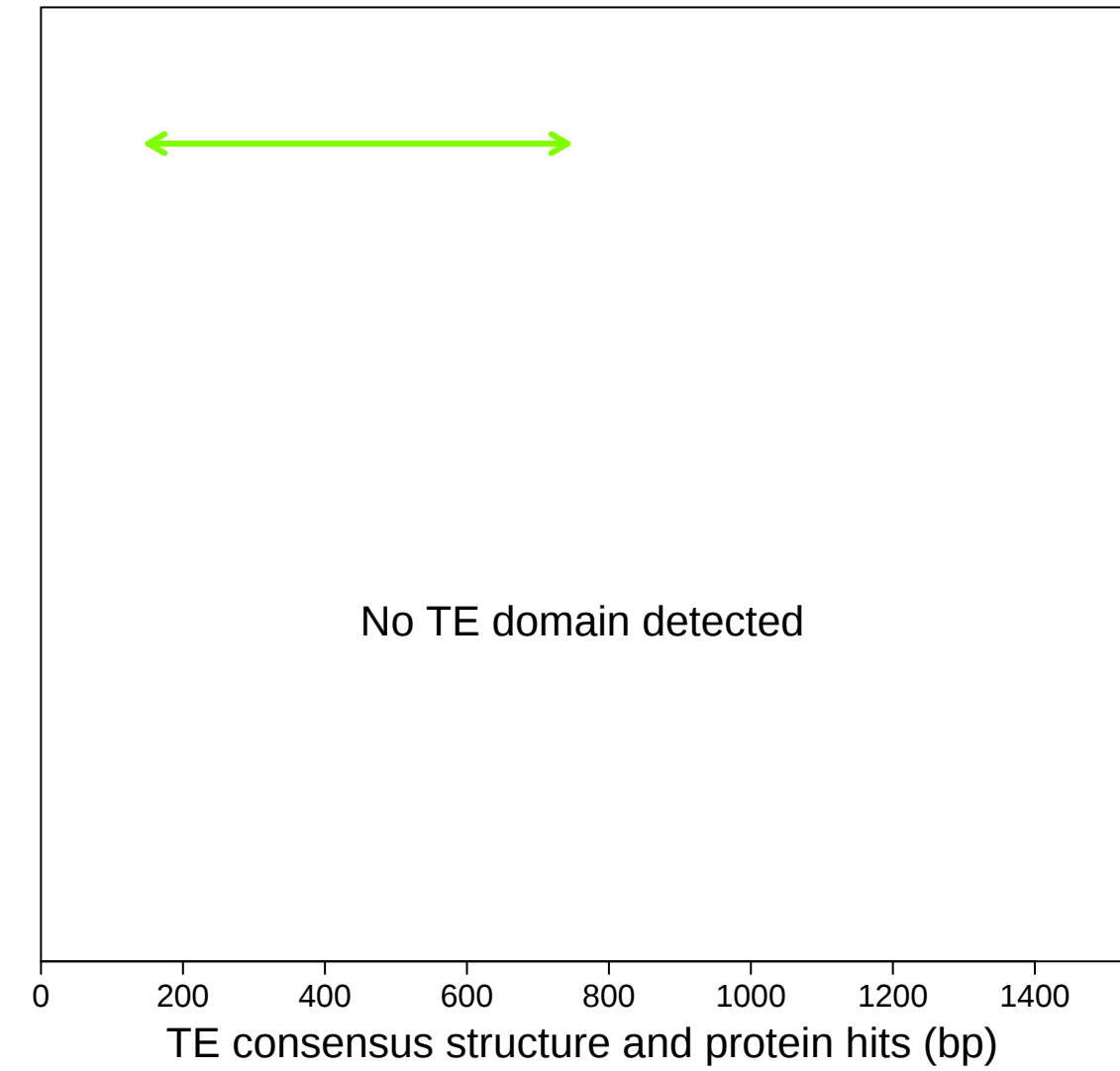
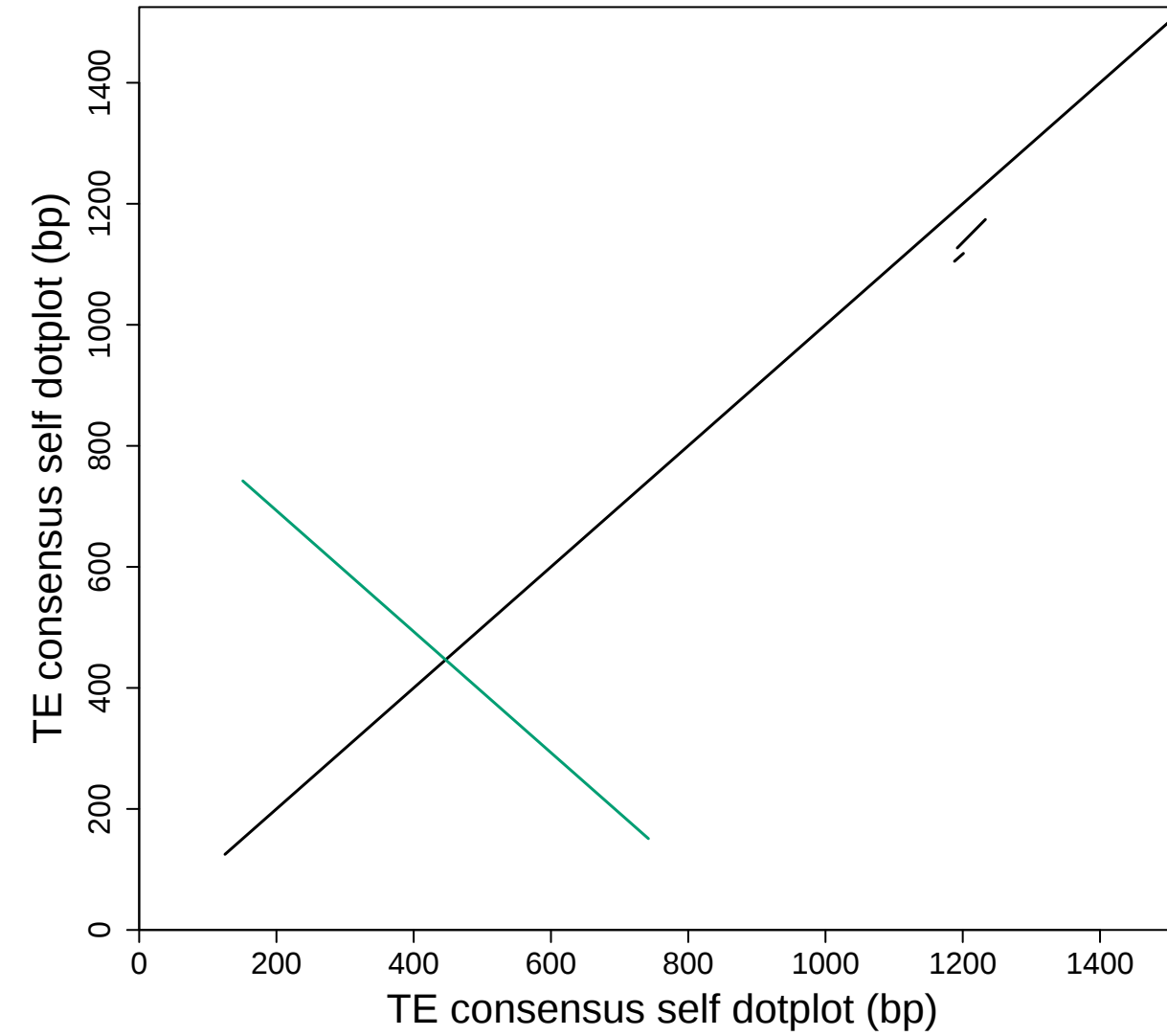
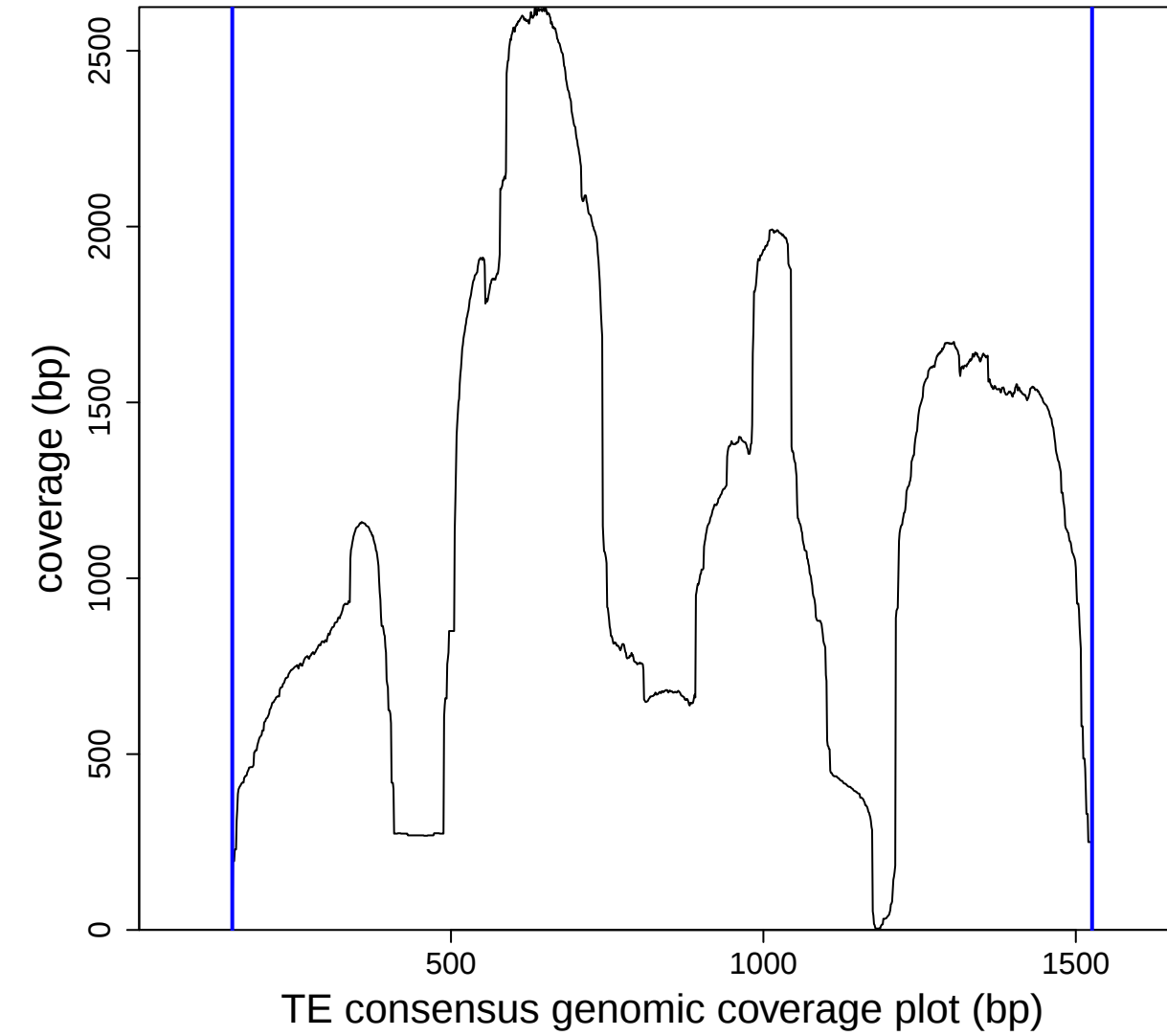
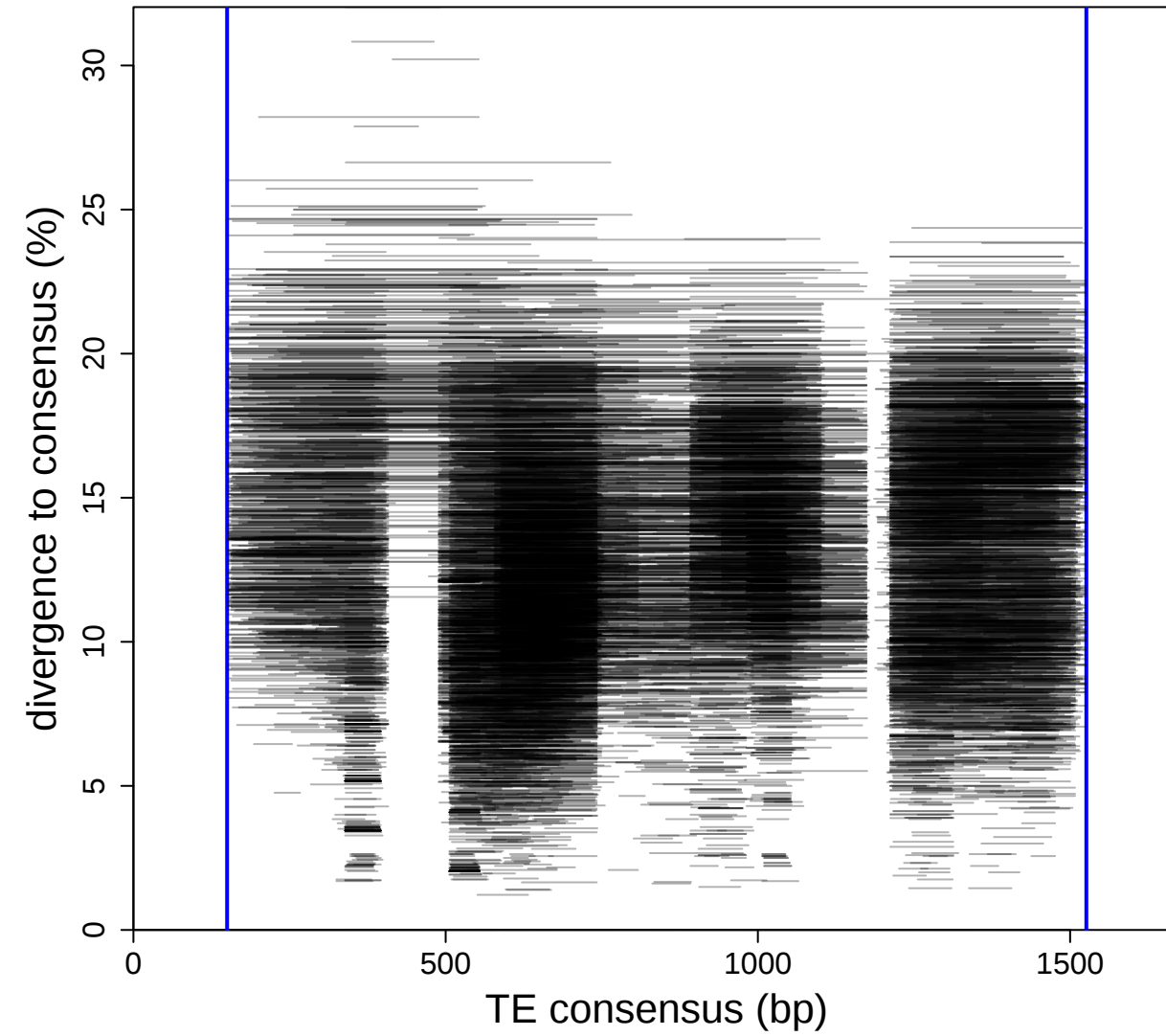
a.b.bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.f
size: 1321bp; fragments: 10789; full length: 0 (>=1188.9bp)

After TEtrimmer 1321 bp



9.fasta.b.bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.f
size: 1676bp; fragments: 8977; full length: 0 (>=1508.4bp)

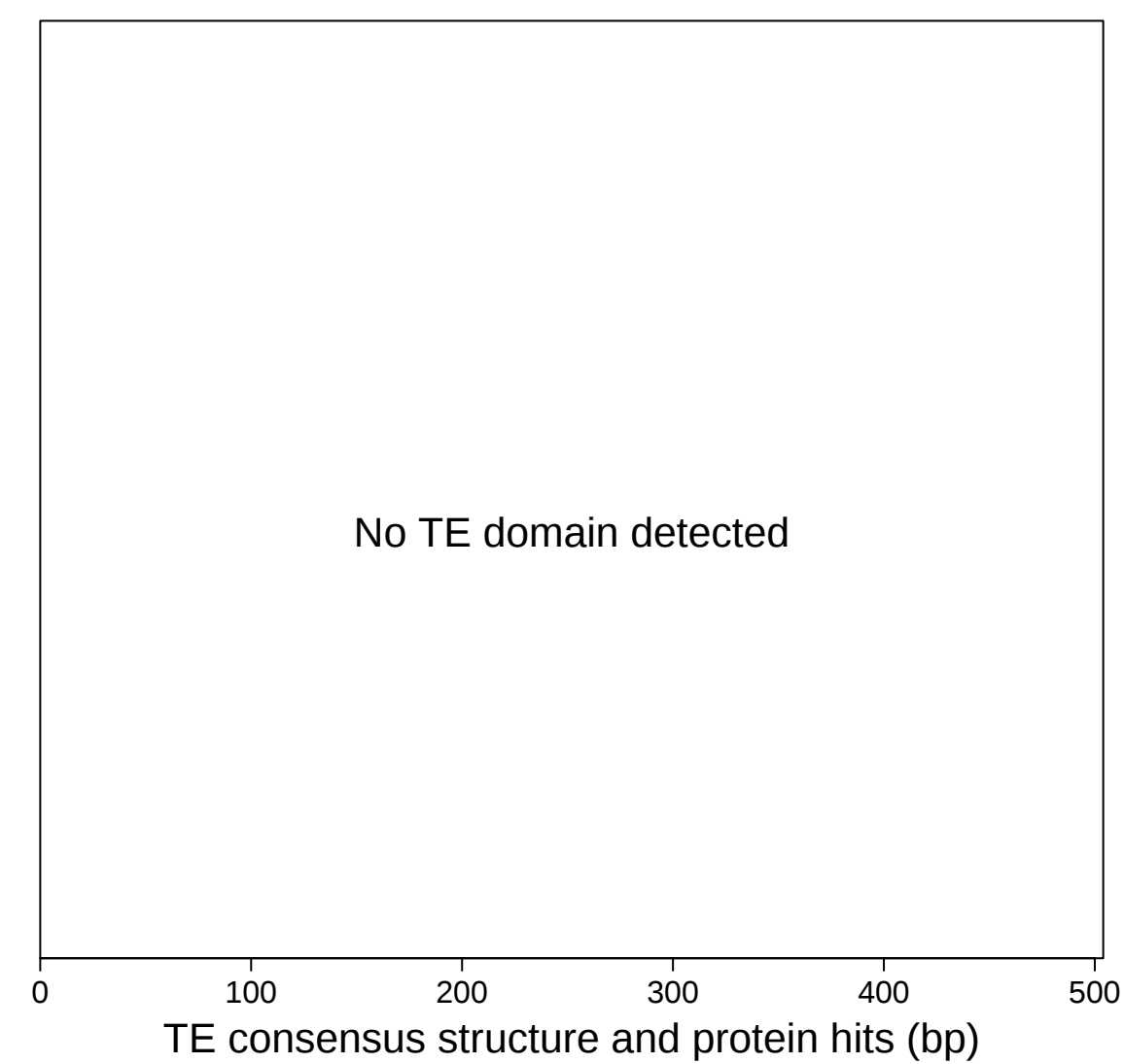
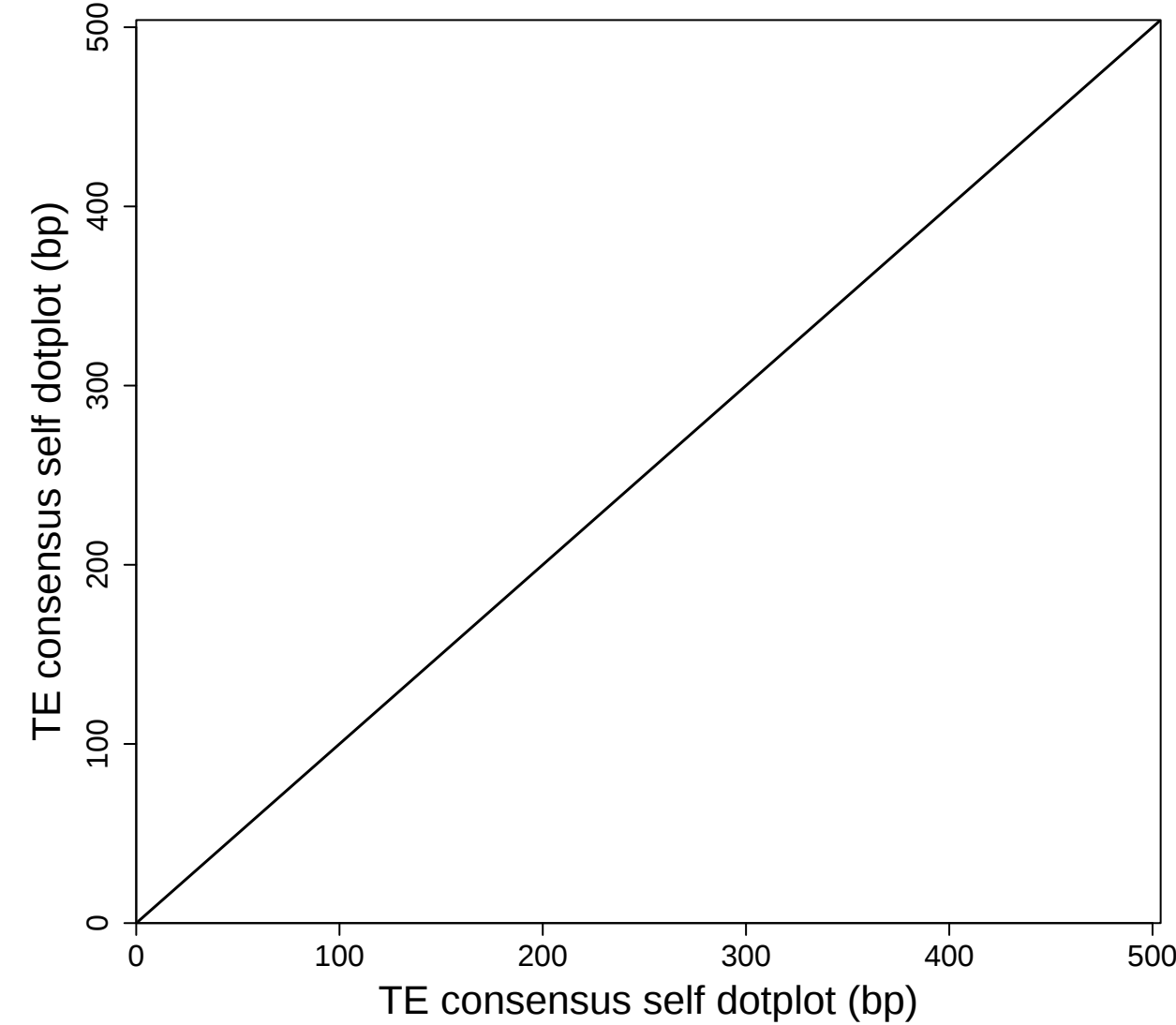
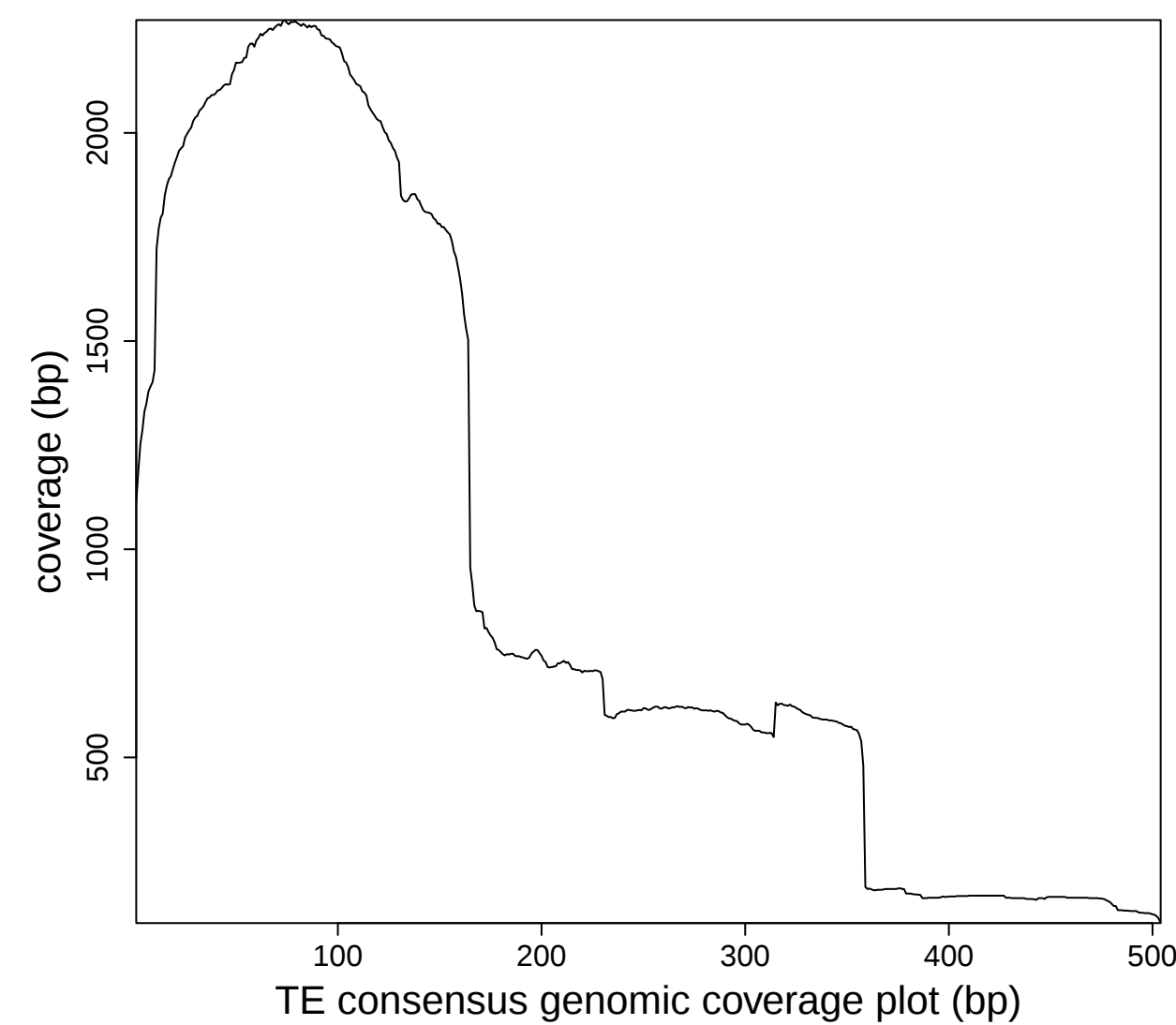
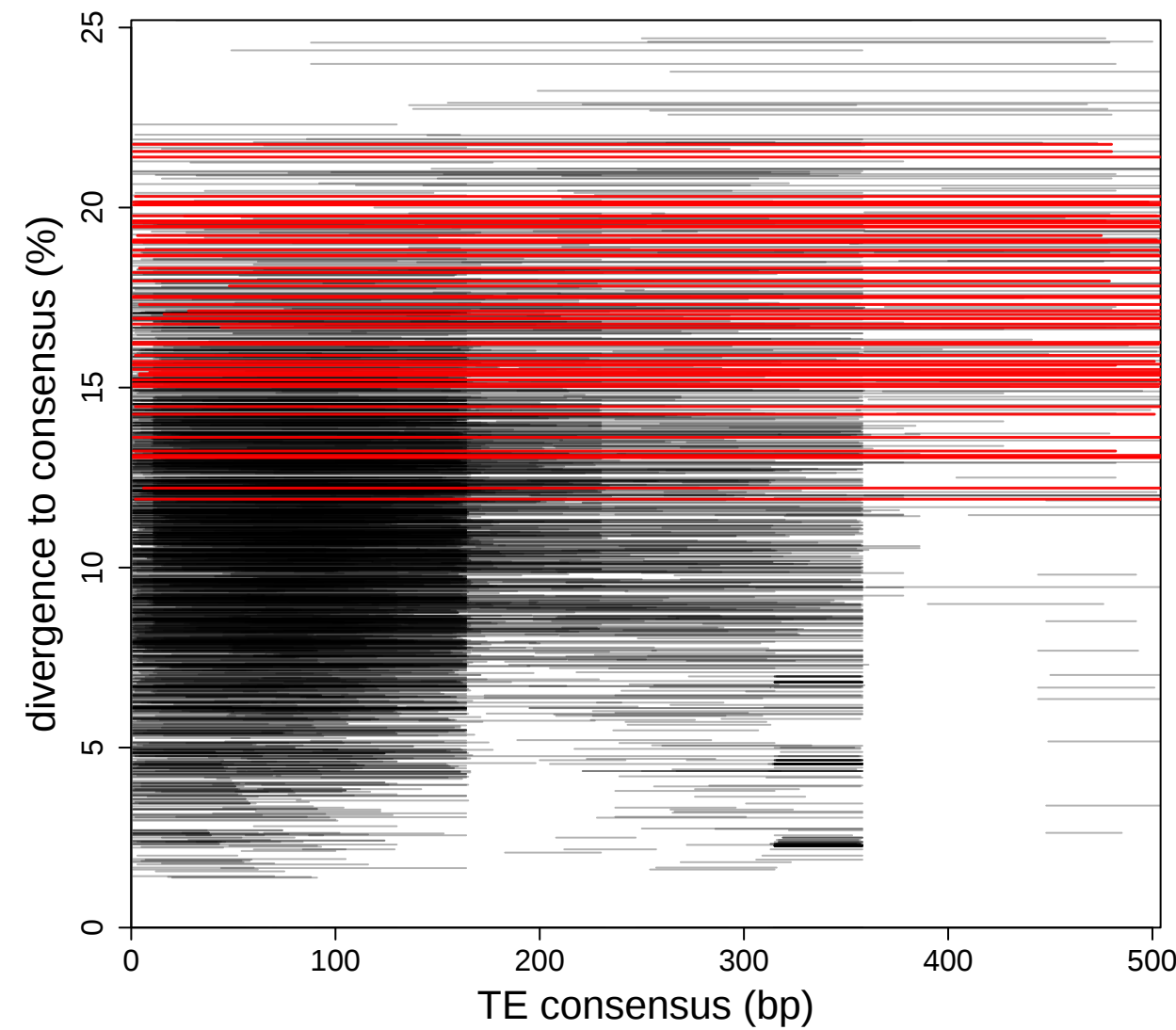
After TEtrimmer Extended plot Blue lines are boundaries



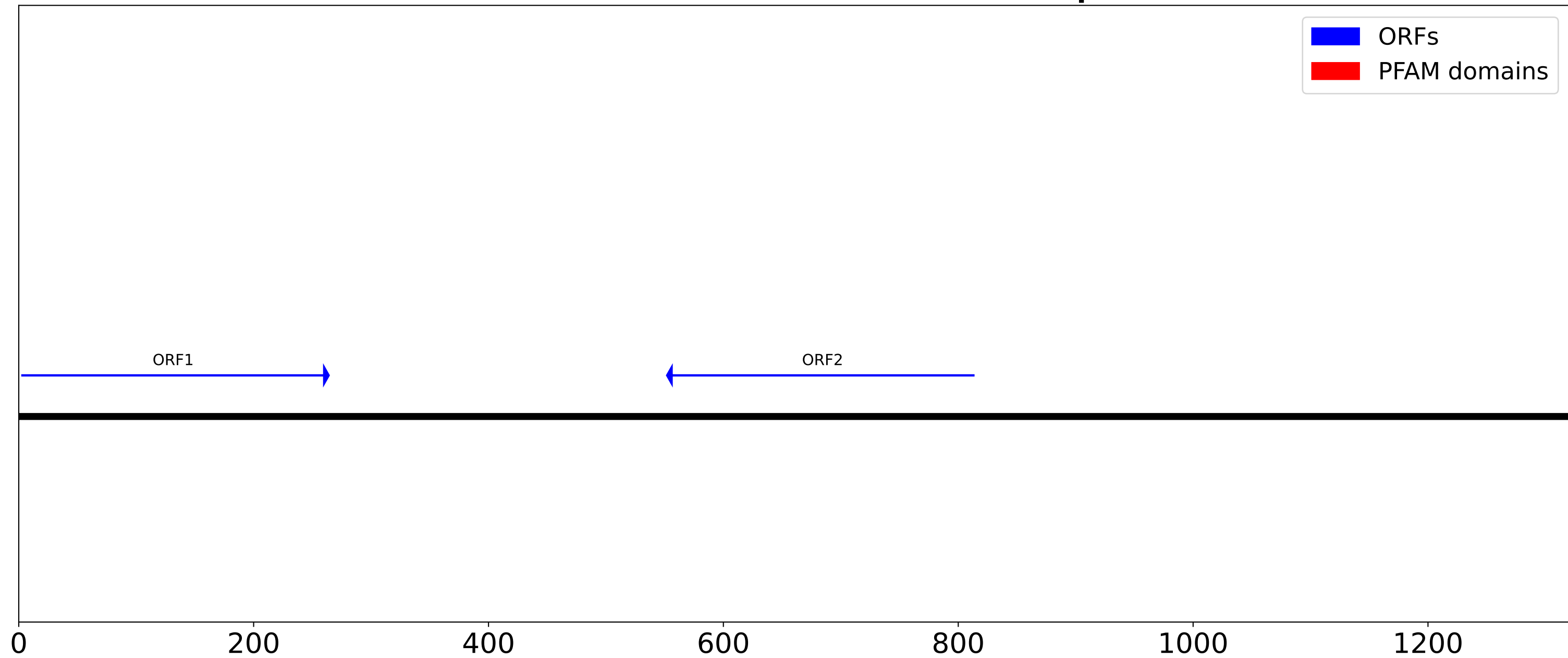
Before TEtrimmer 504 bp

TE: rnd_4_family_1229

size: 504bp; fragments: 3016; full length: 59 (≥ 453.6 bp)



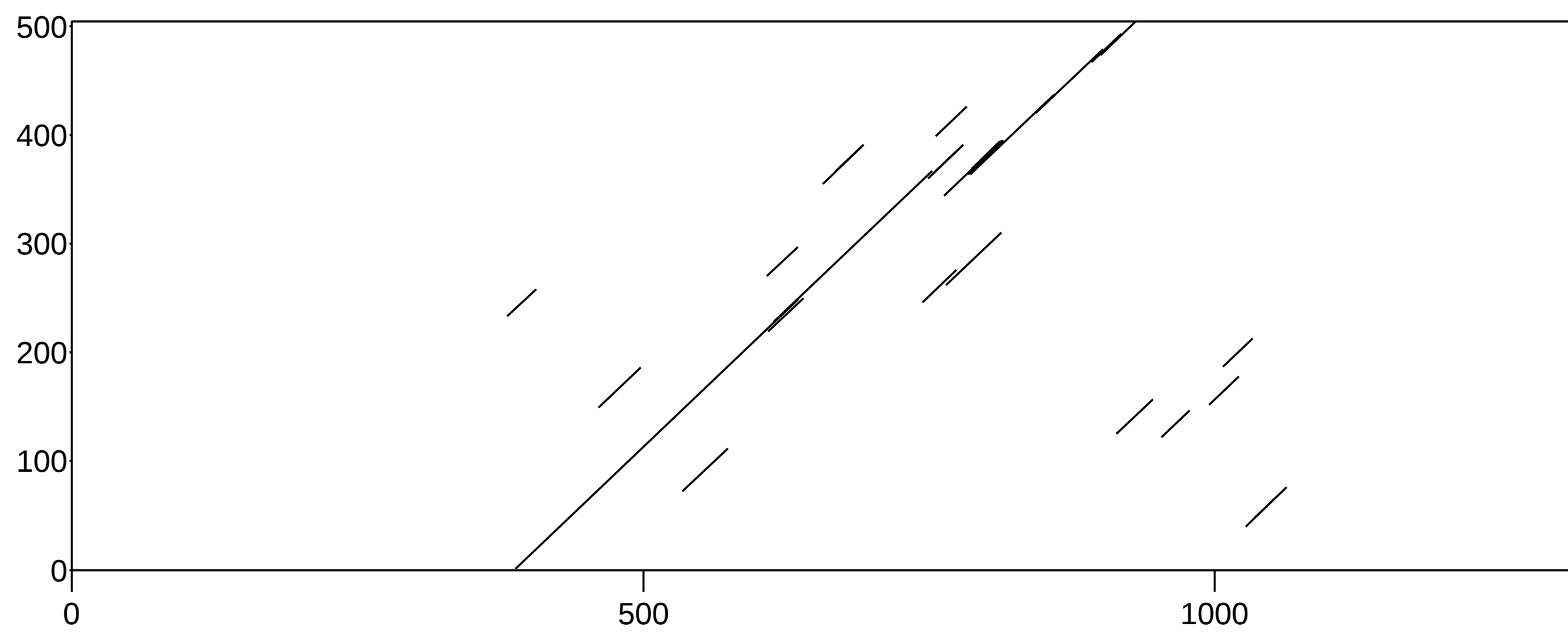
After TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)