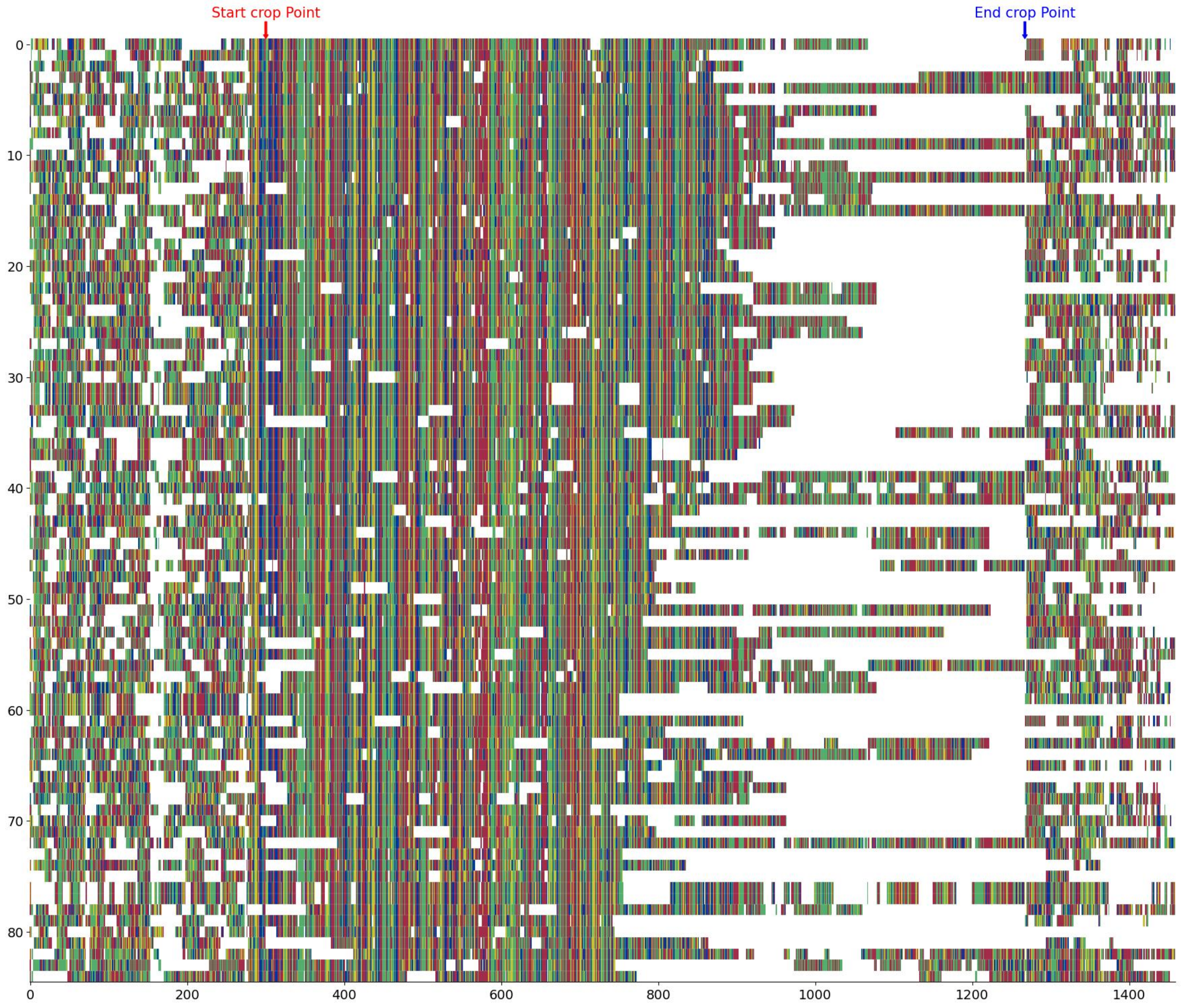




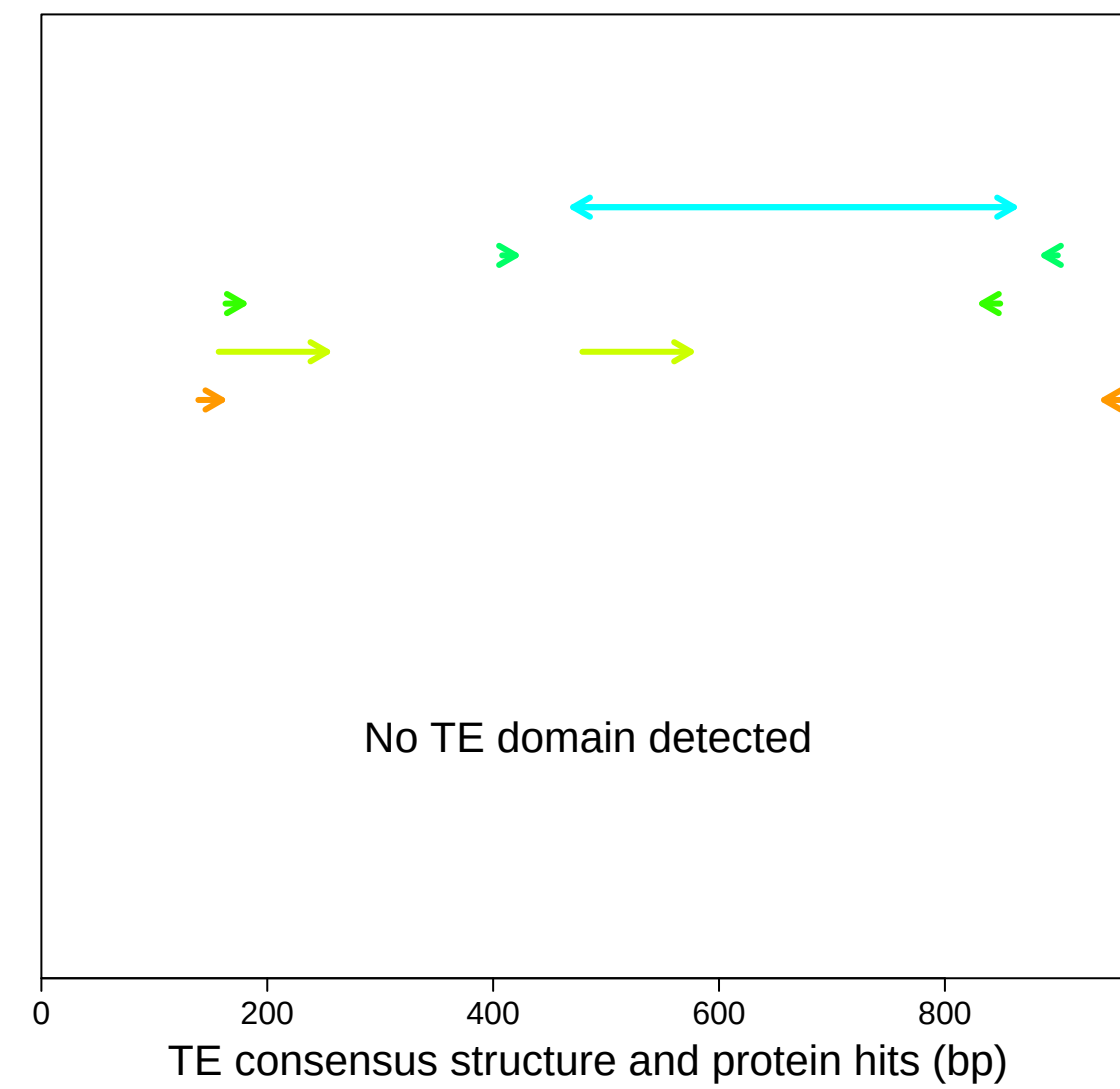
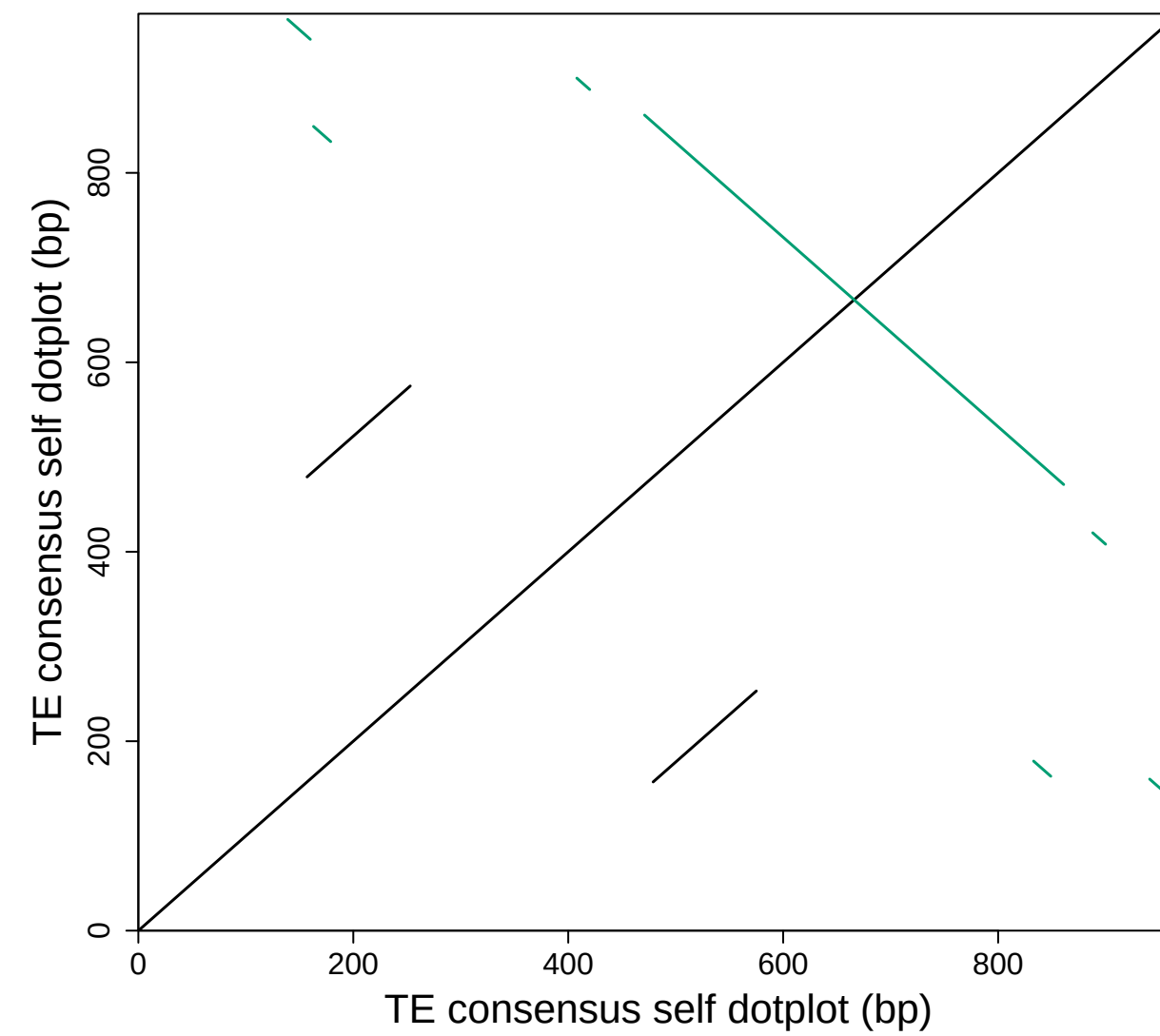
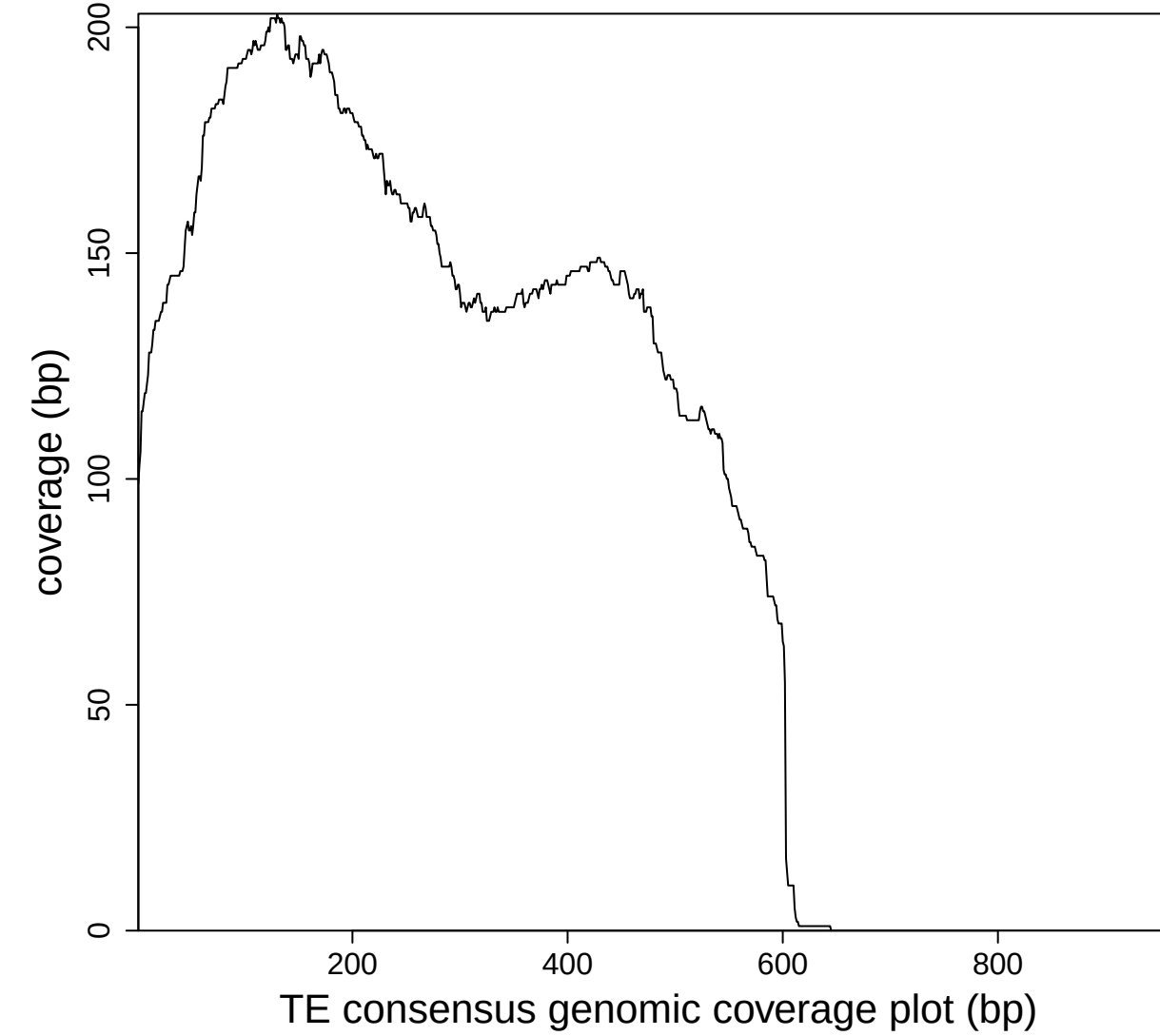
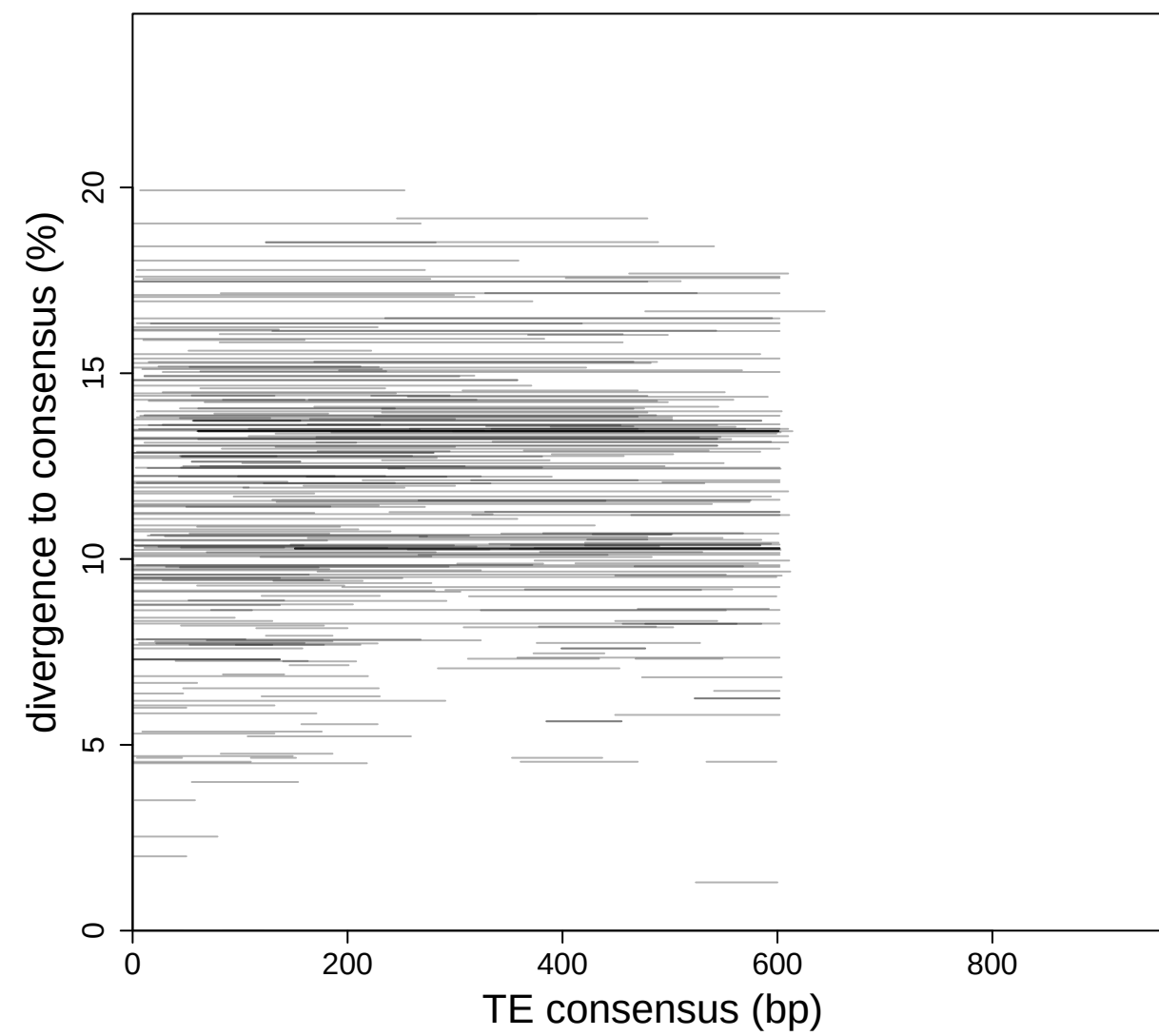
Start crop Point

End crop Point



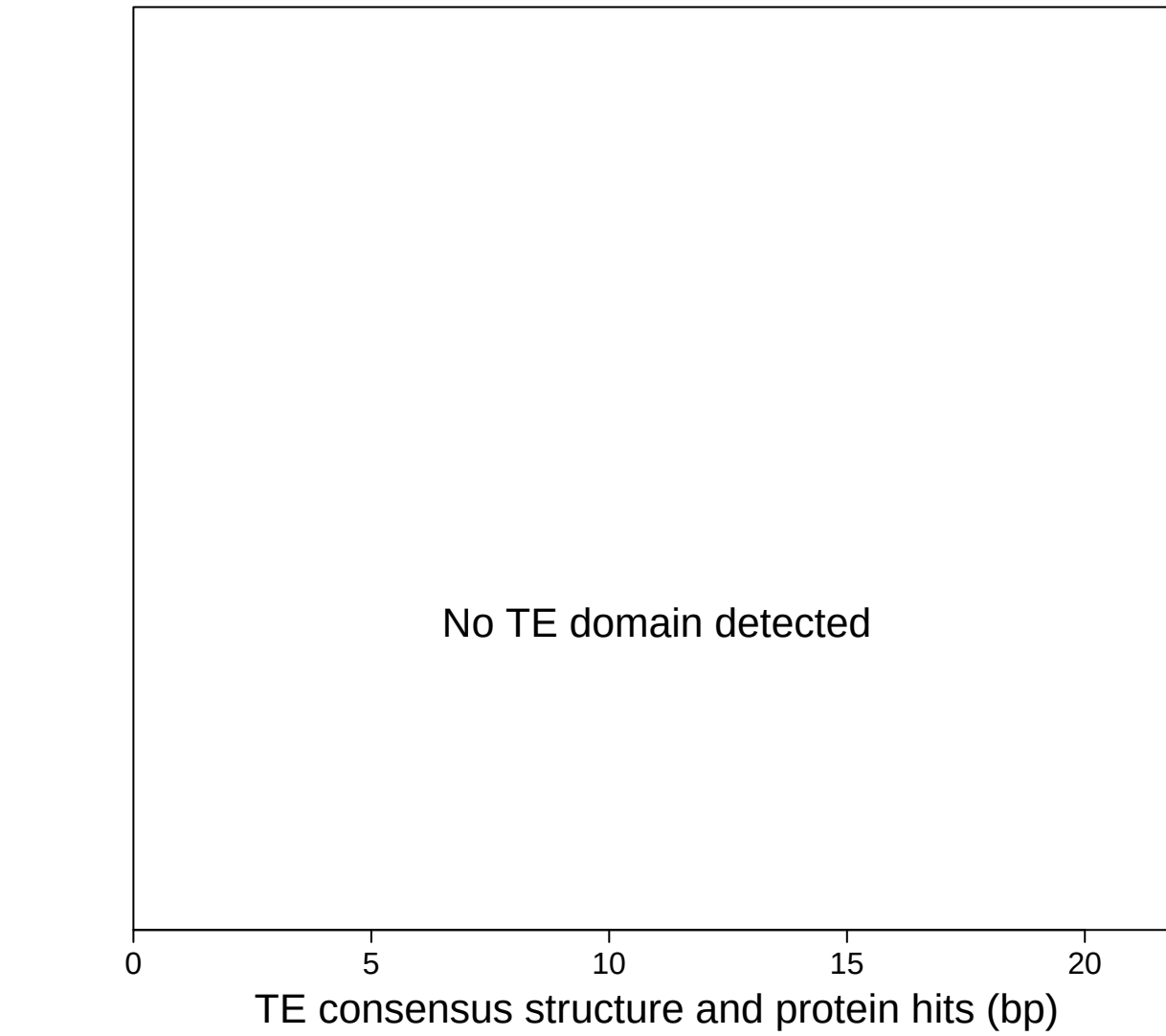
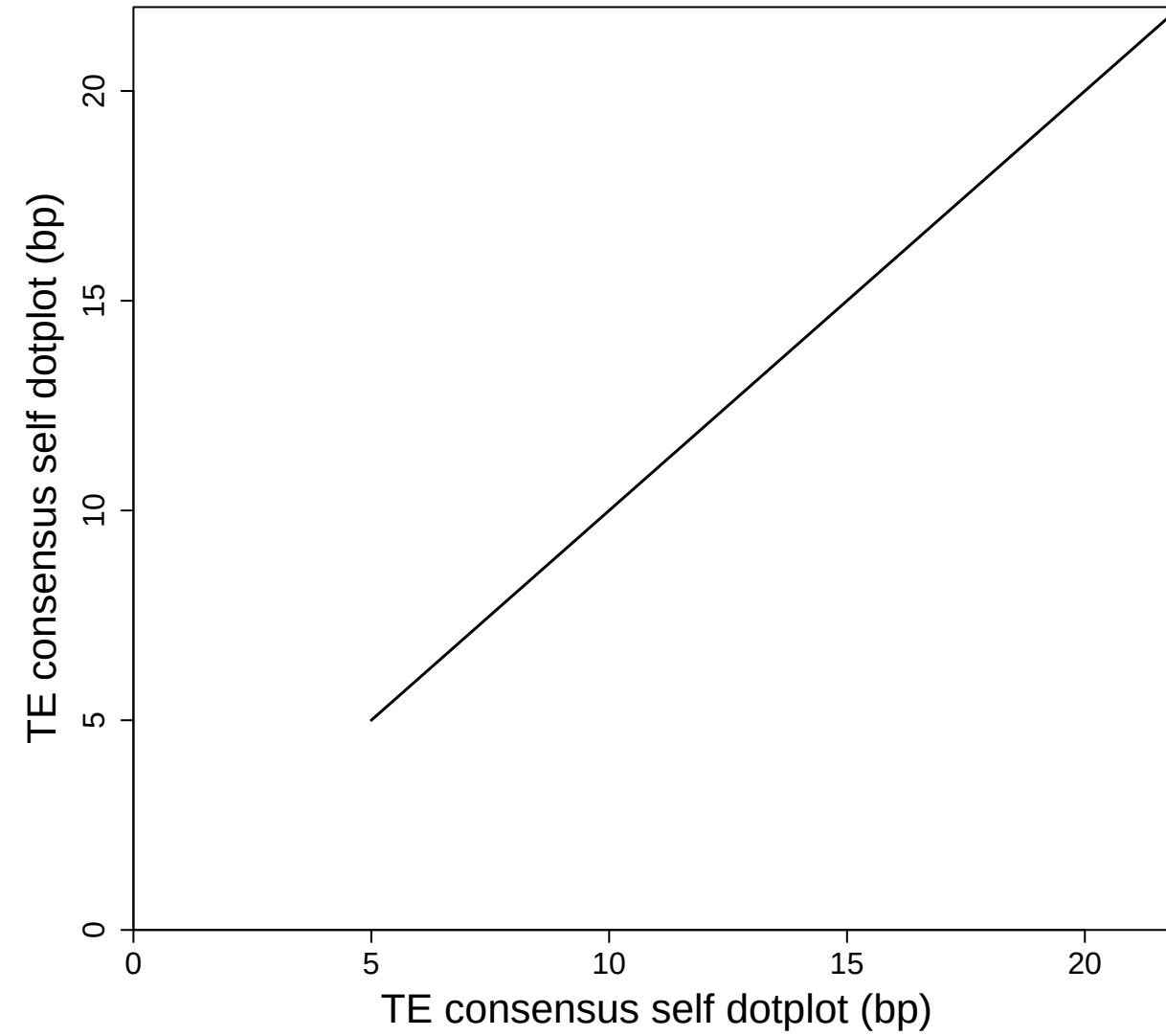
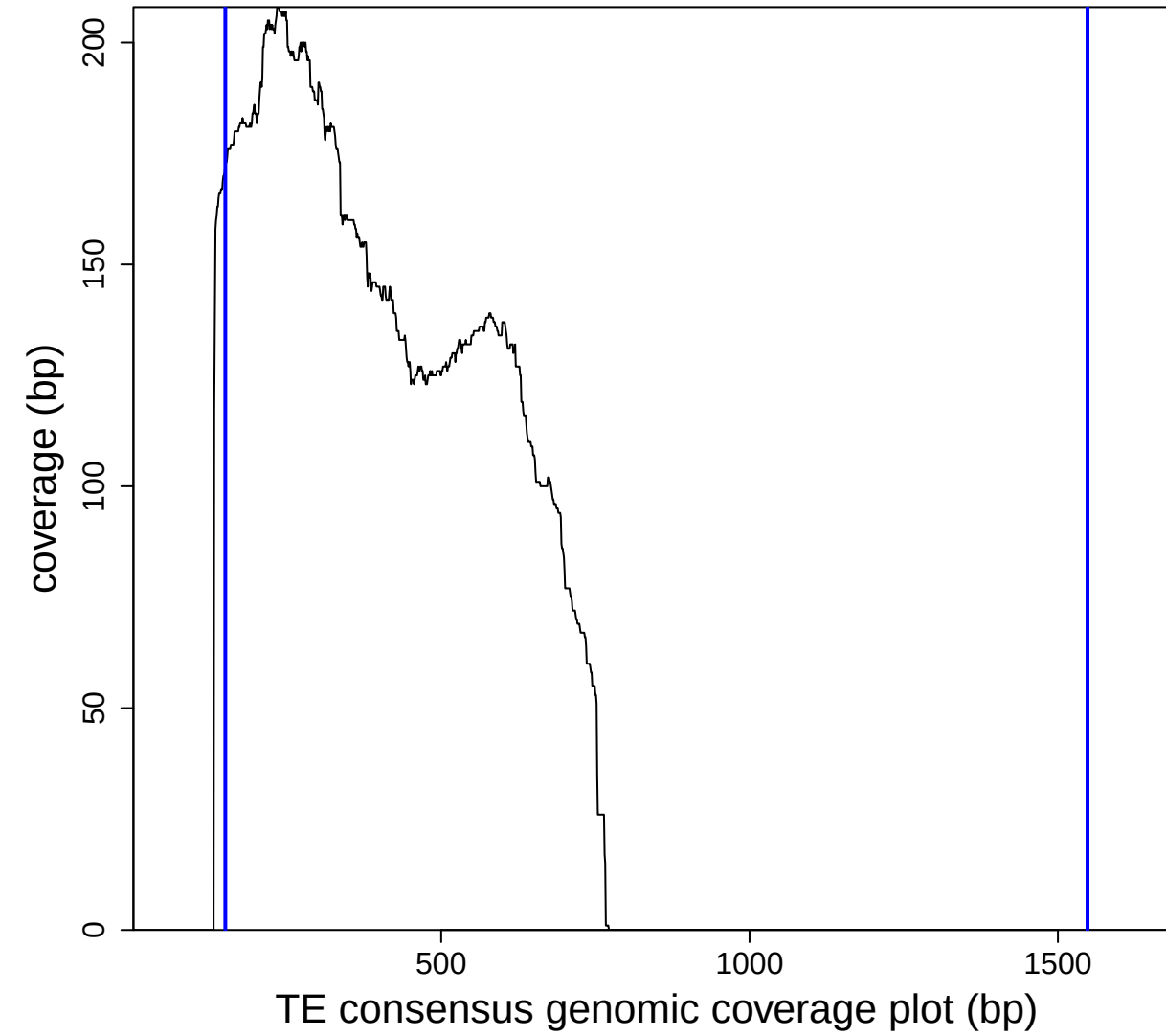
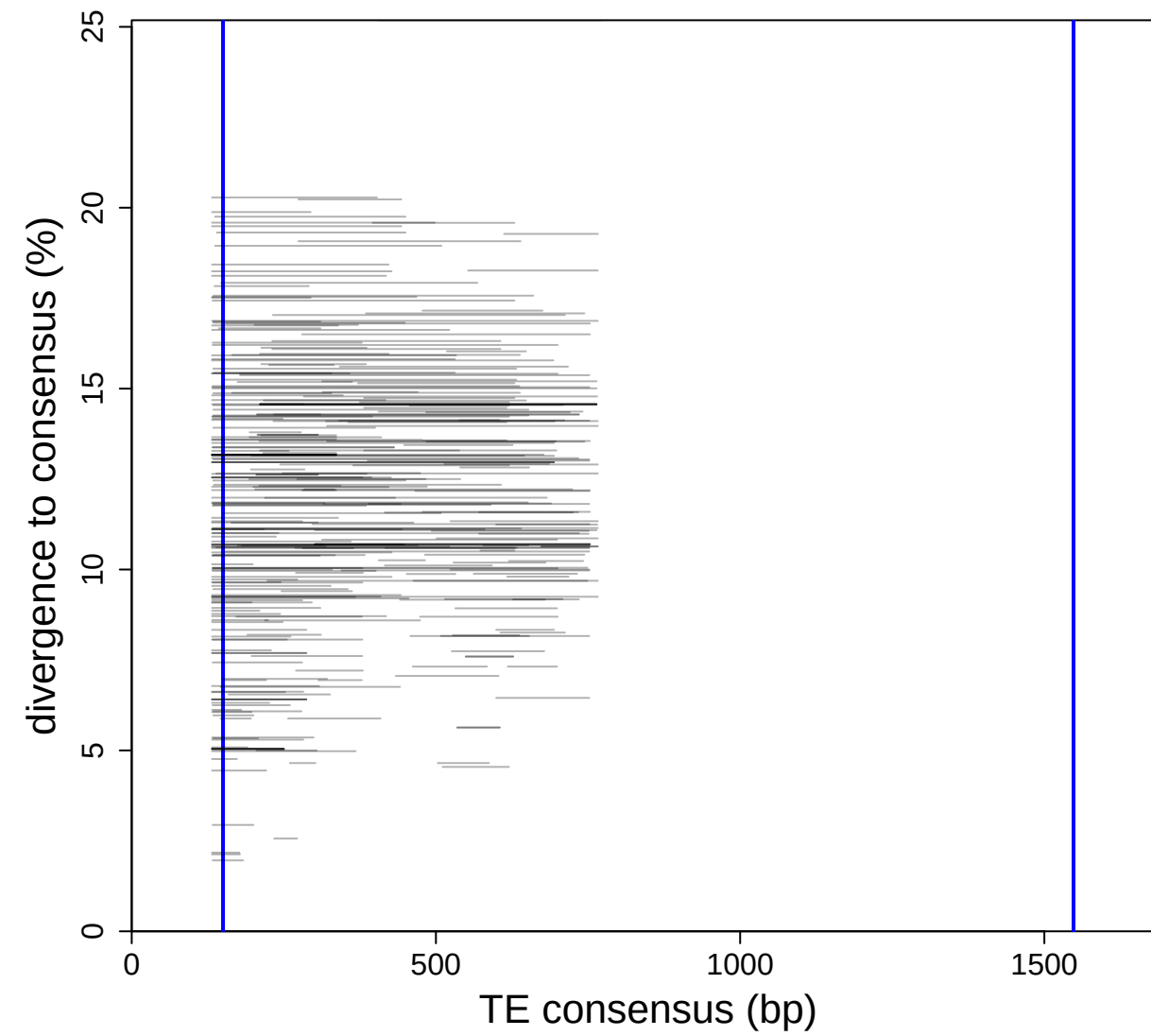
bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.fa_c
size: 968bp; fragments: 325; full length: 0 (>=871.2bp)

After TEtrimmer 968 bp



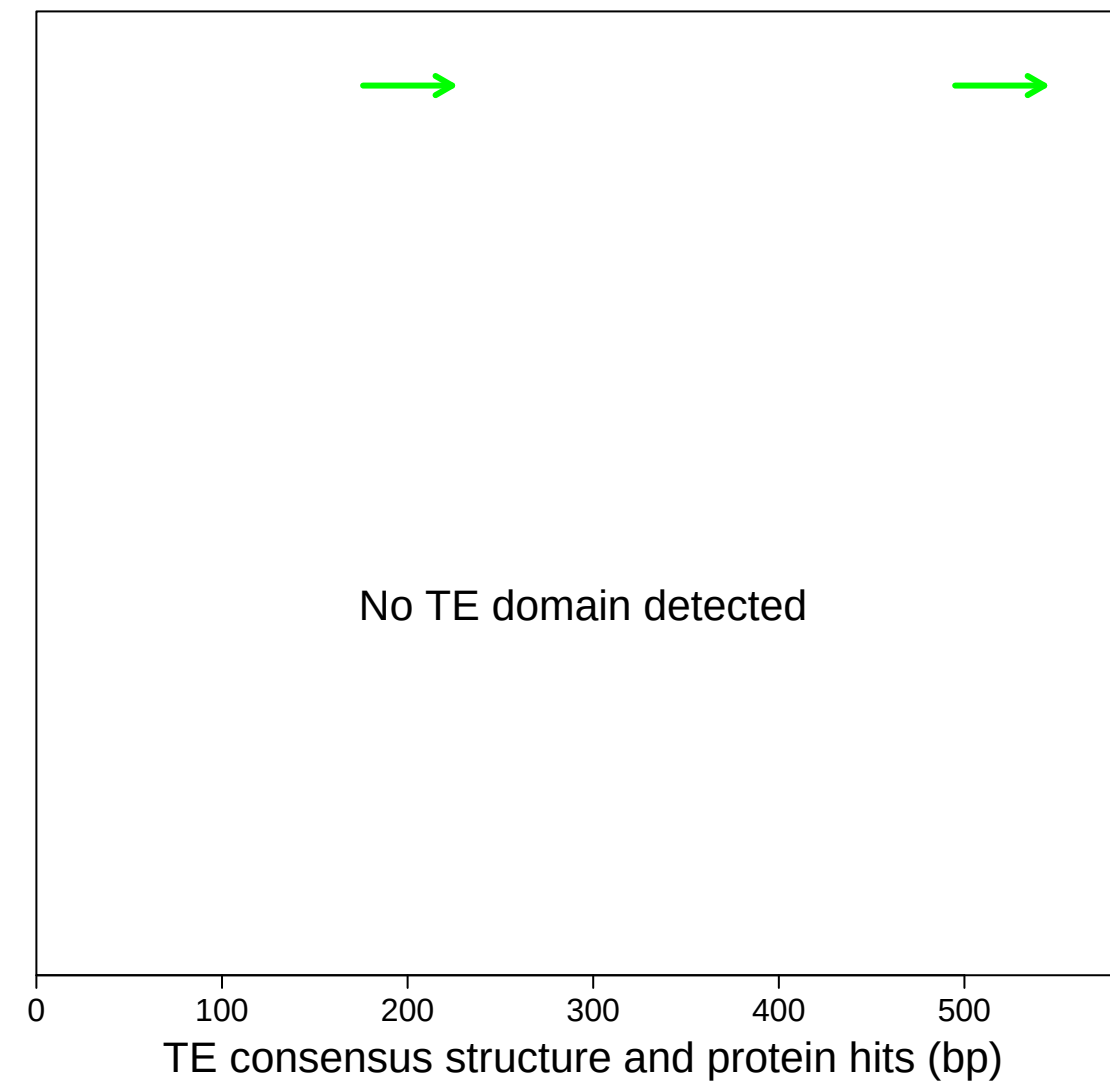
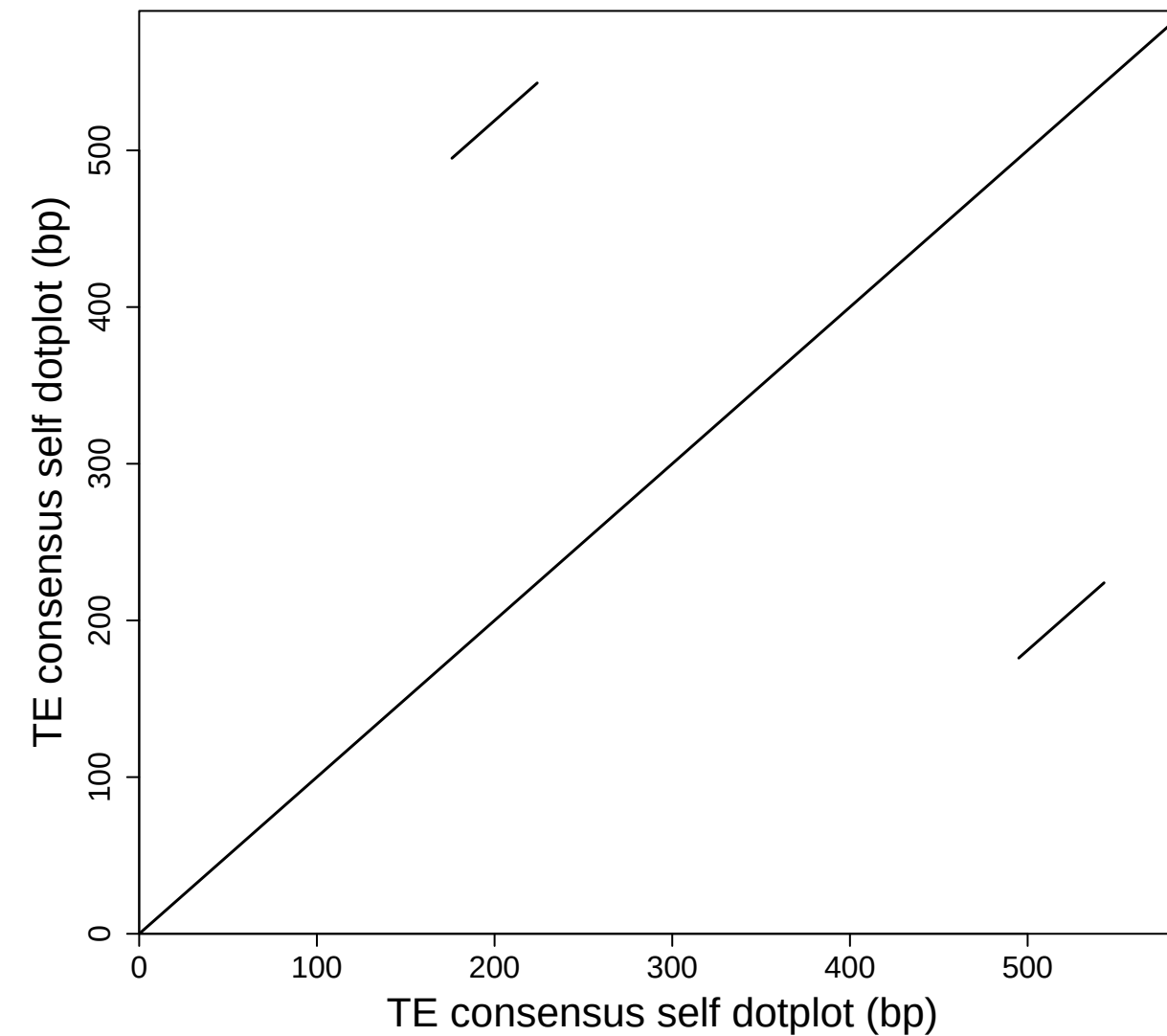
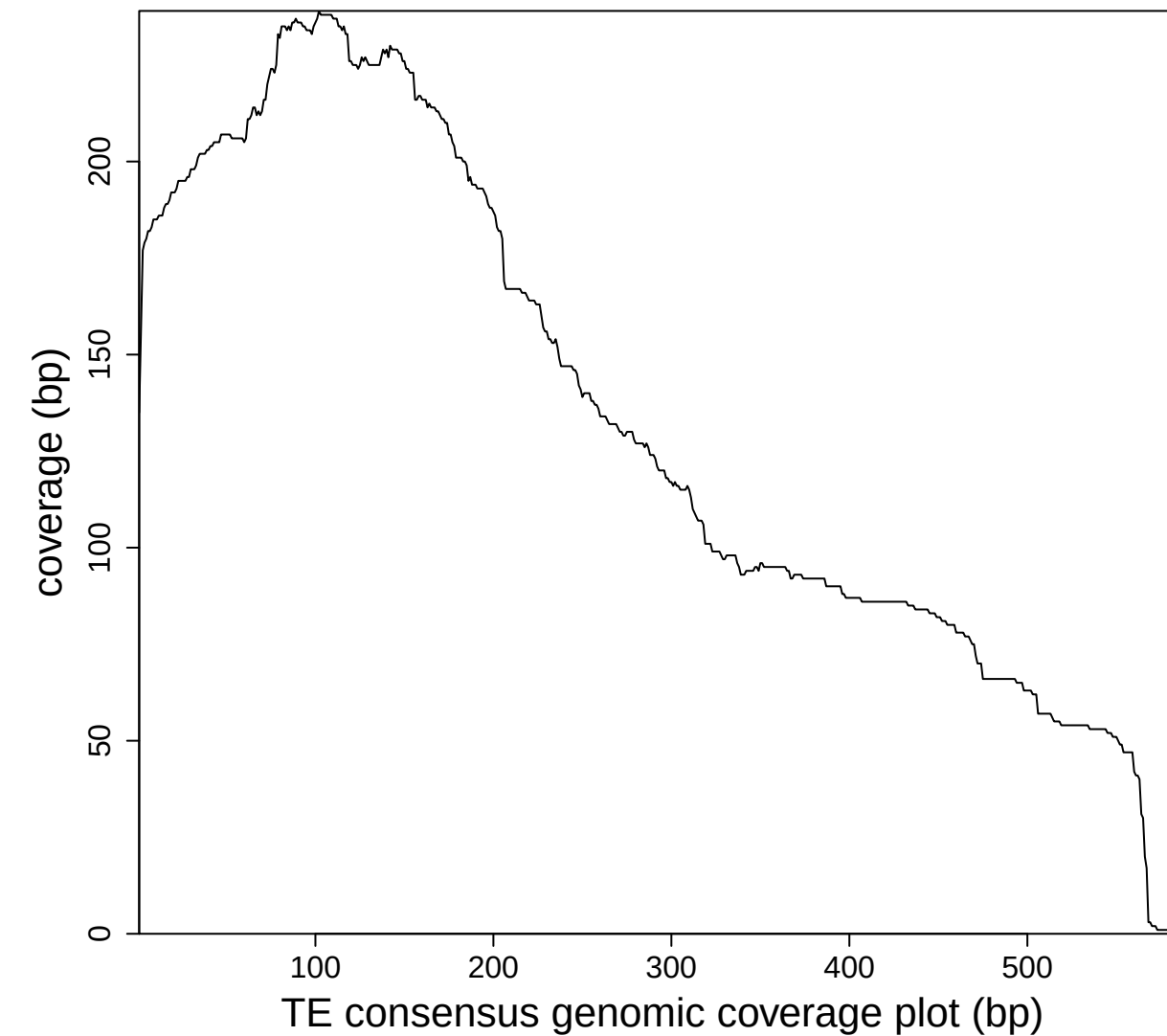
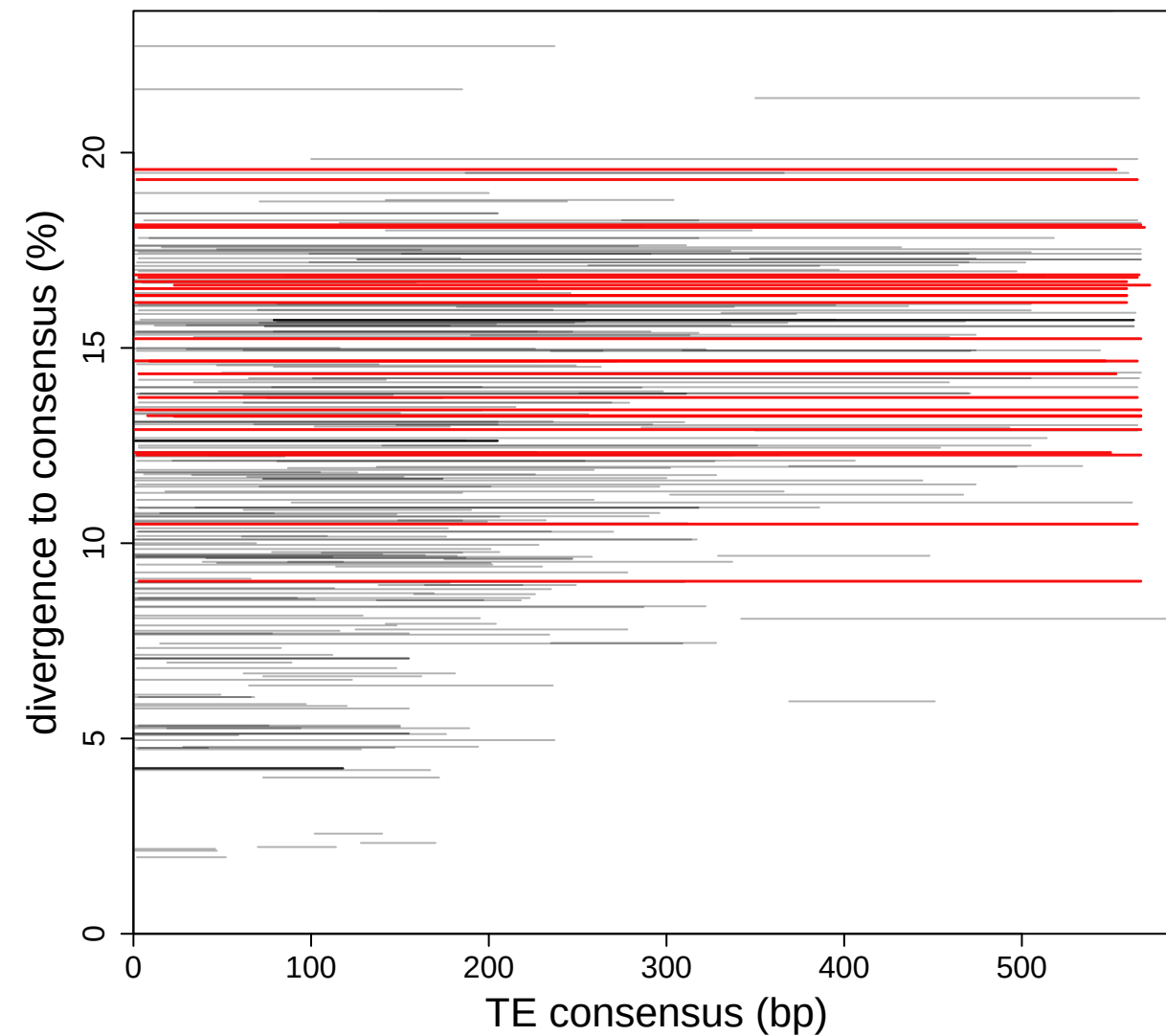
ly_1834.fasta.b.bed uf.bed g_1.bed fm_1.bed_0_0 bcln.fa aln.f
size: 1698bp; fragments: 331; full length: 0 (>=1528.2bp)

After TEtrimmer Extended plot Blue lines are boundaries

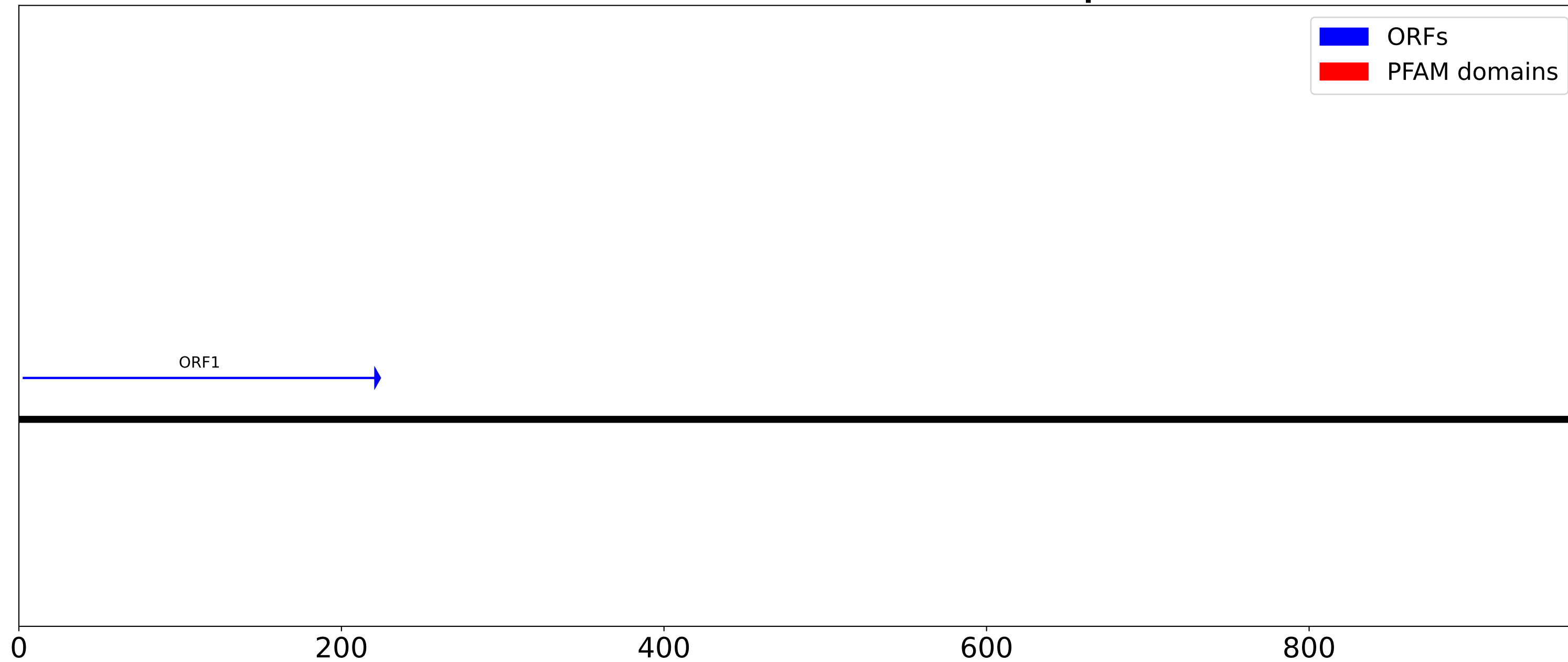


TE: rnd_4_family_1834
size: 589bp; fragments: 298; full length: 25 (≥ 530.1 bp)

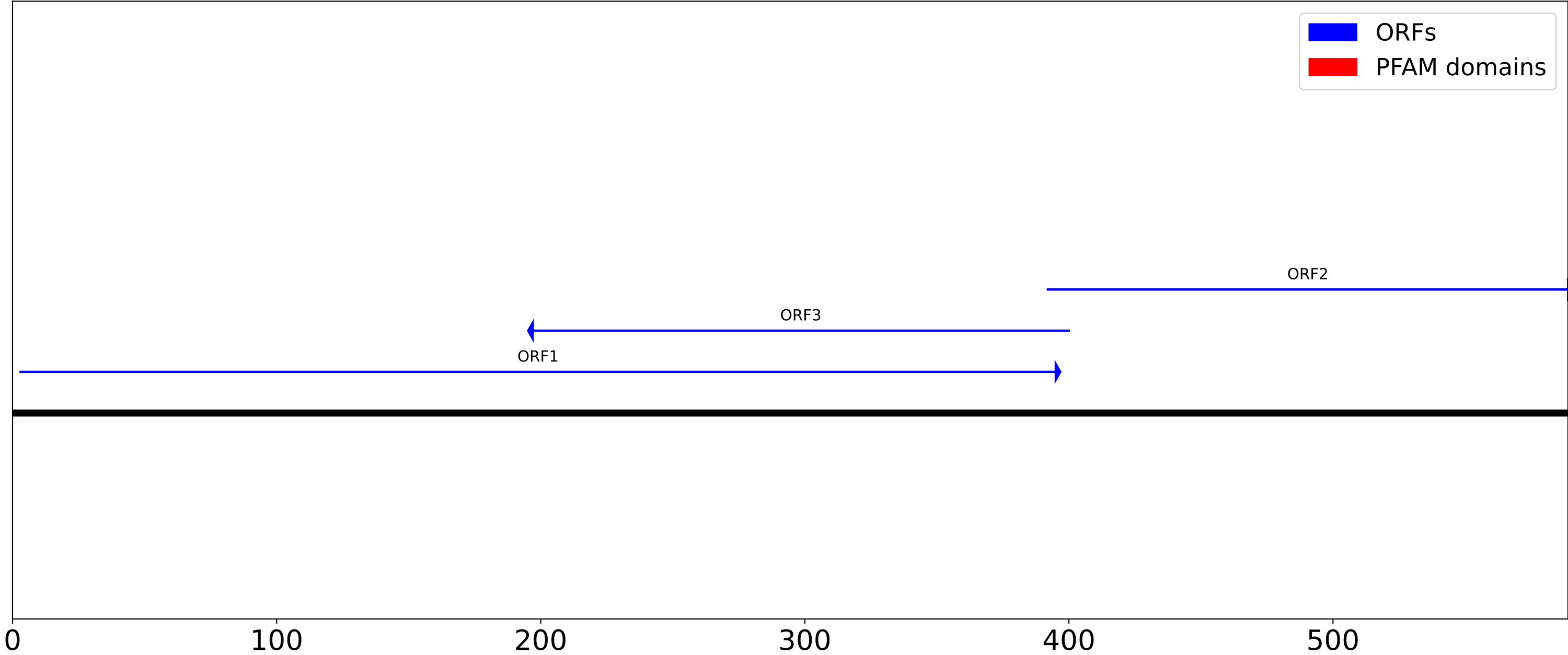
Before TEtrimmer 589 bp



After TEtrimmer ORF and PFAM domain plot



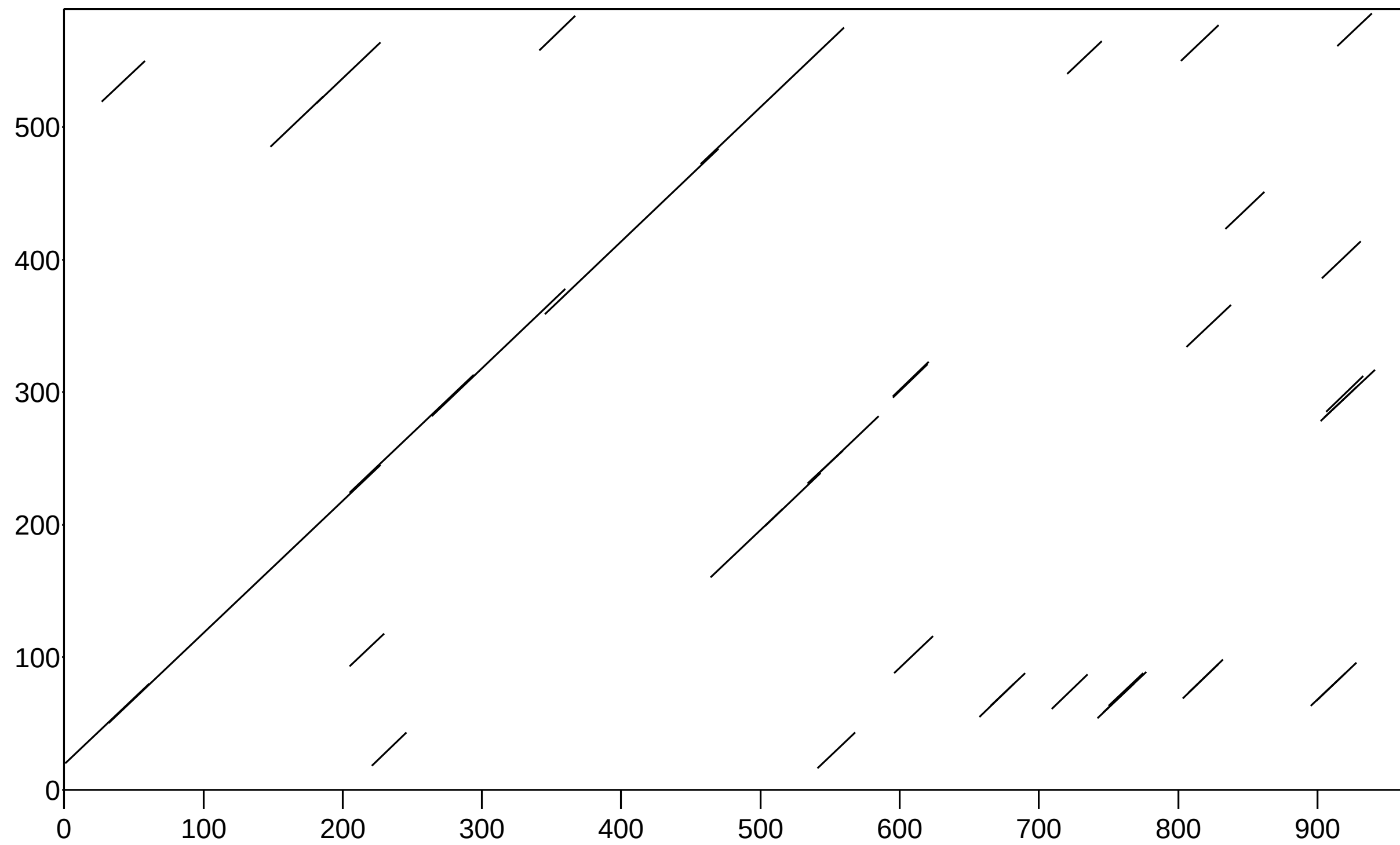
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)