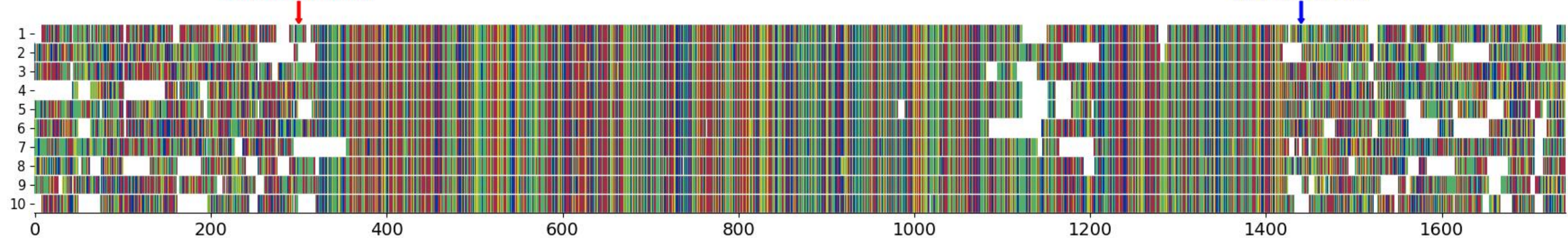


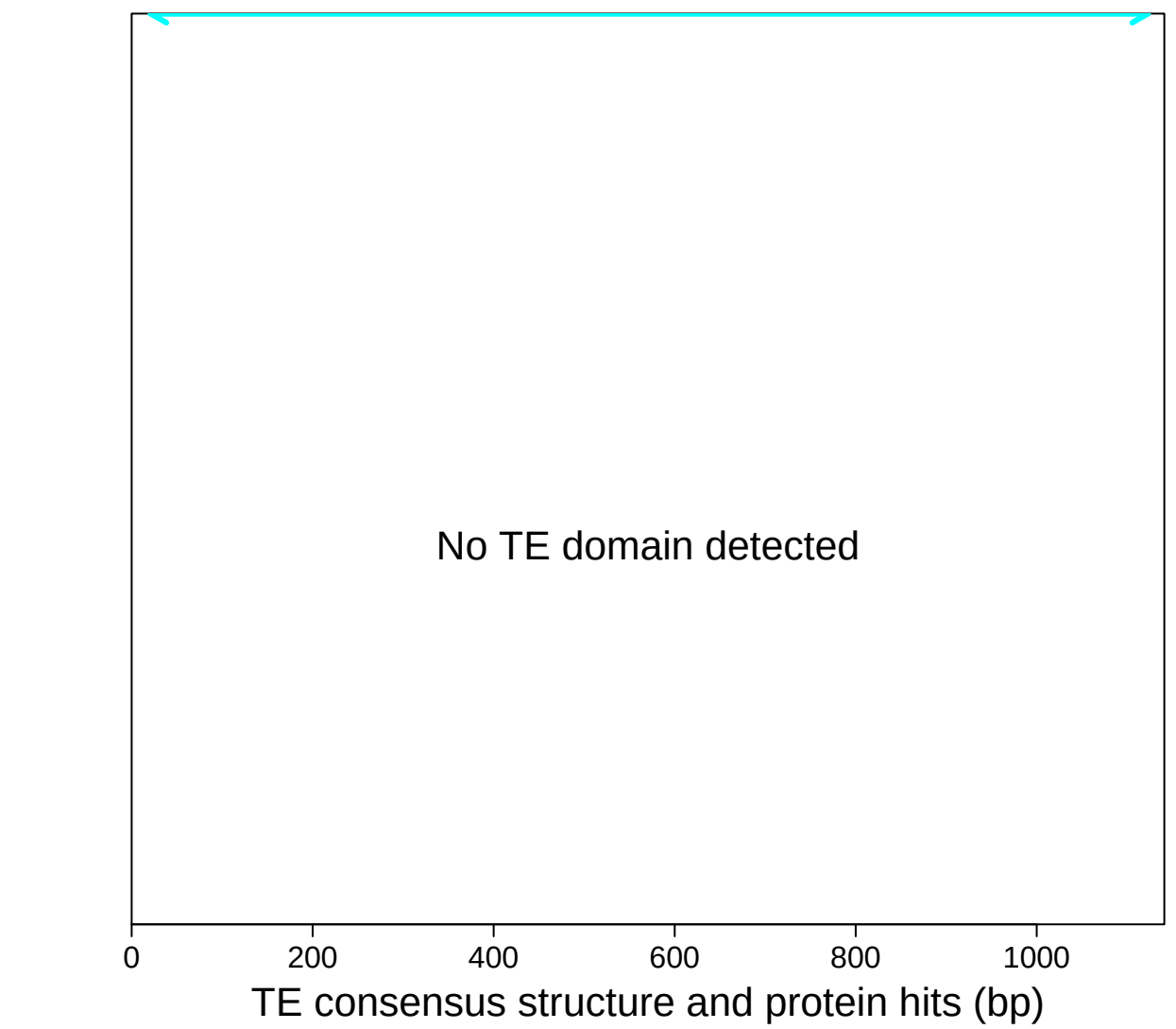
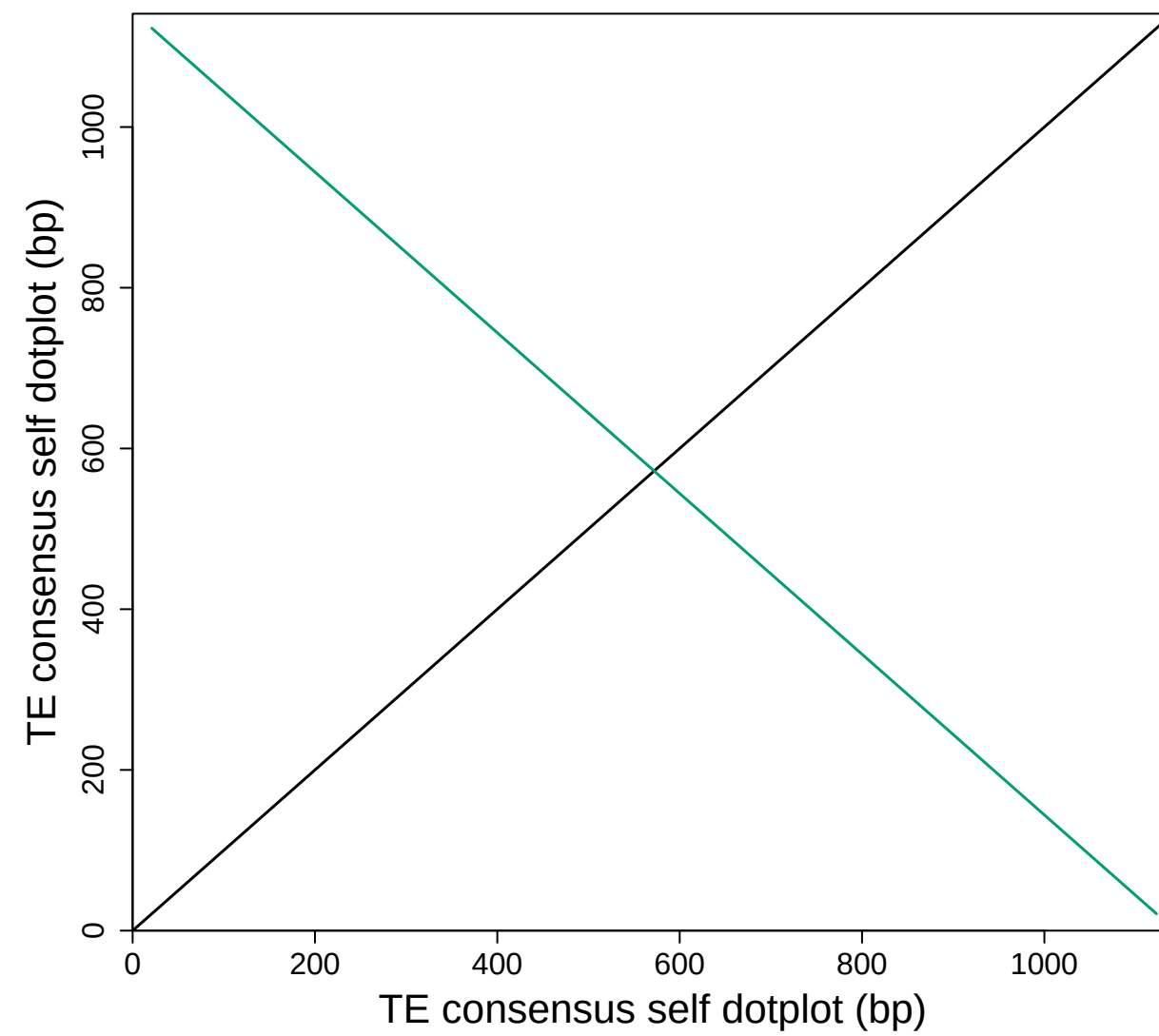
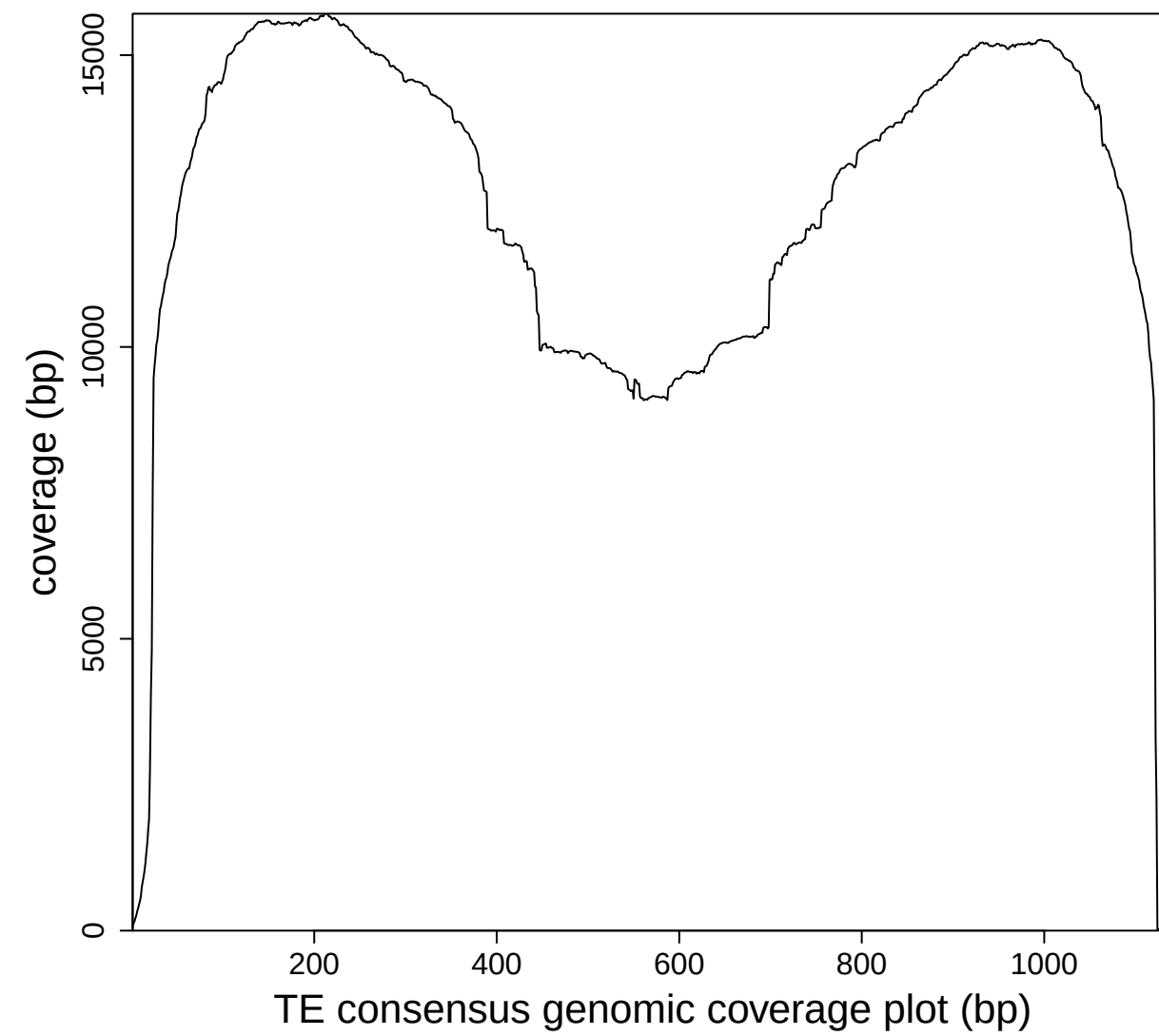
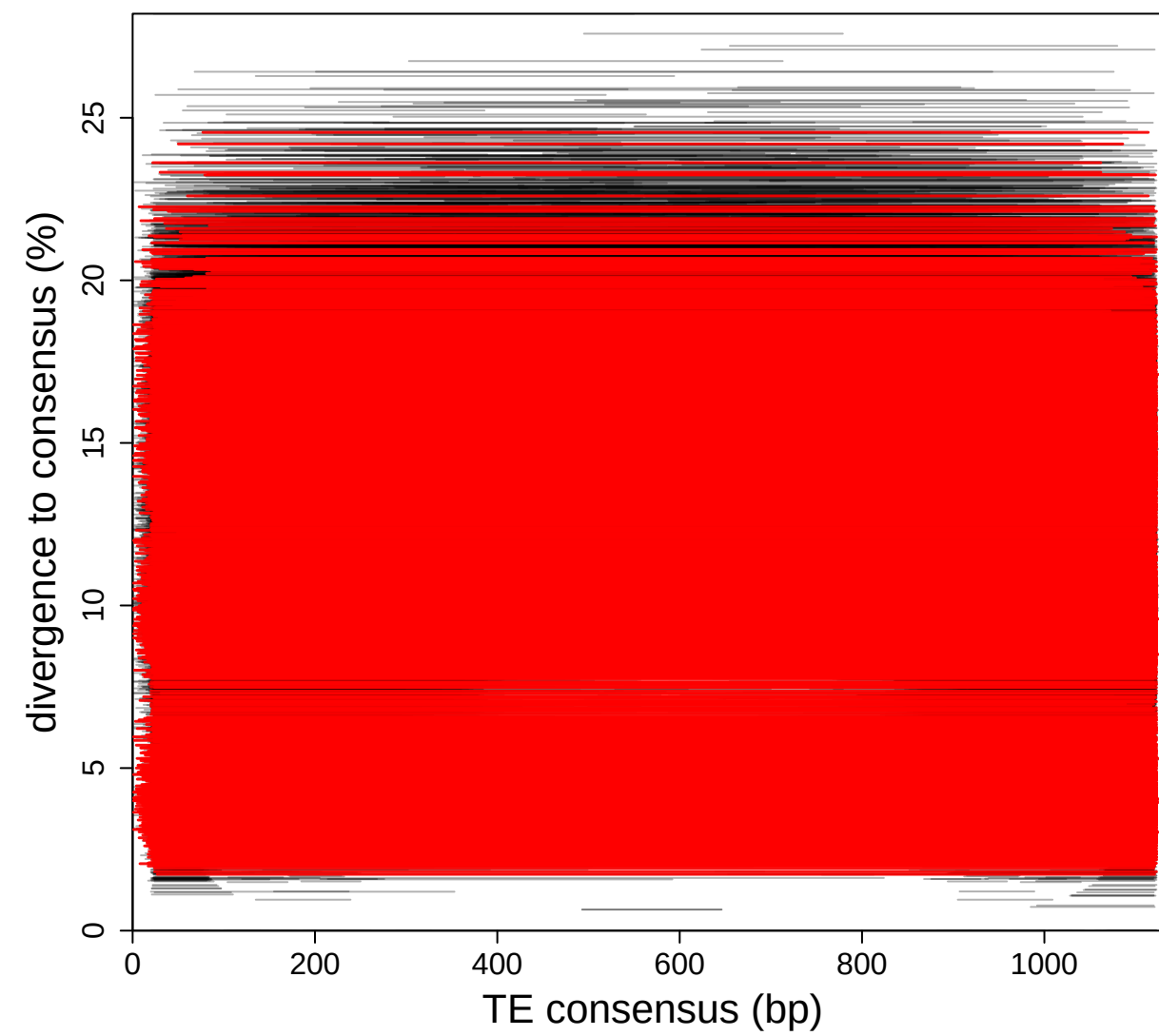
Start crop Point

End crop Point



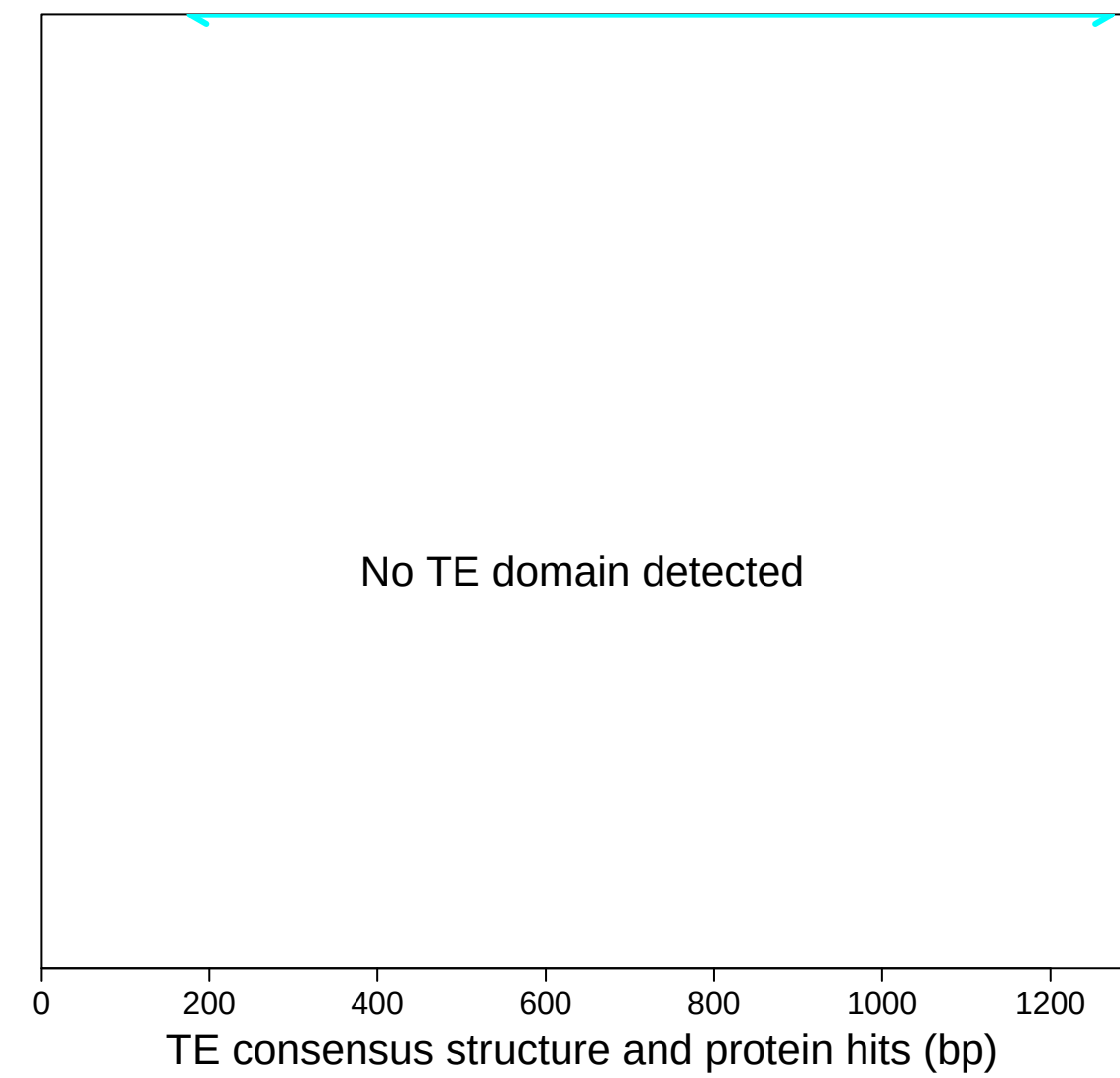
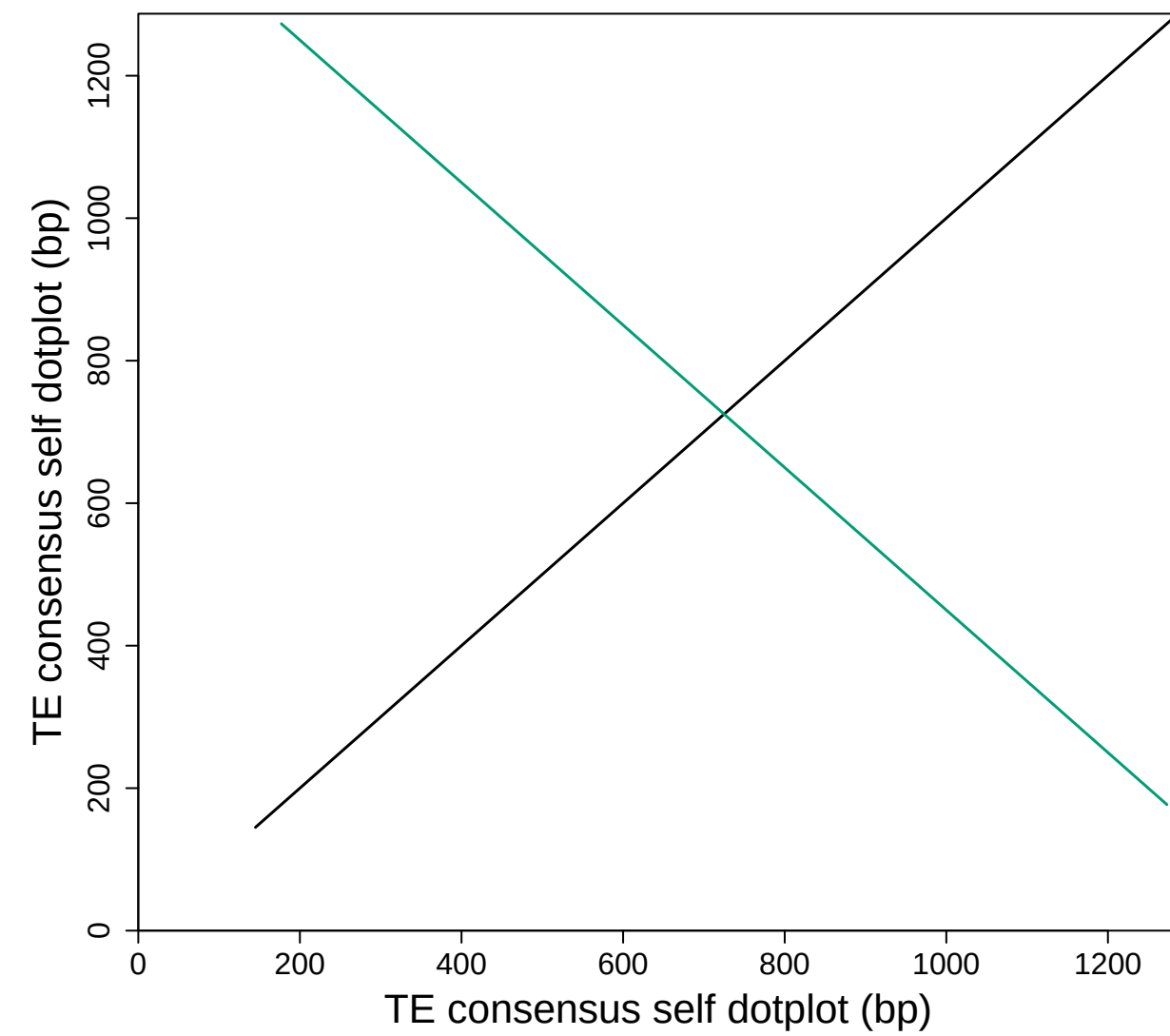
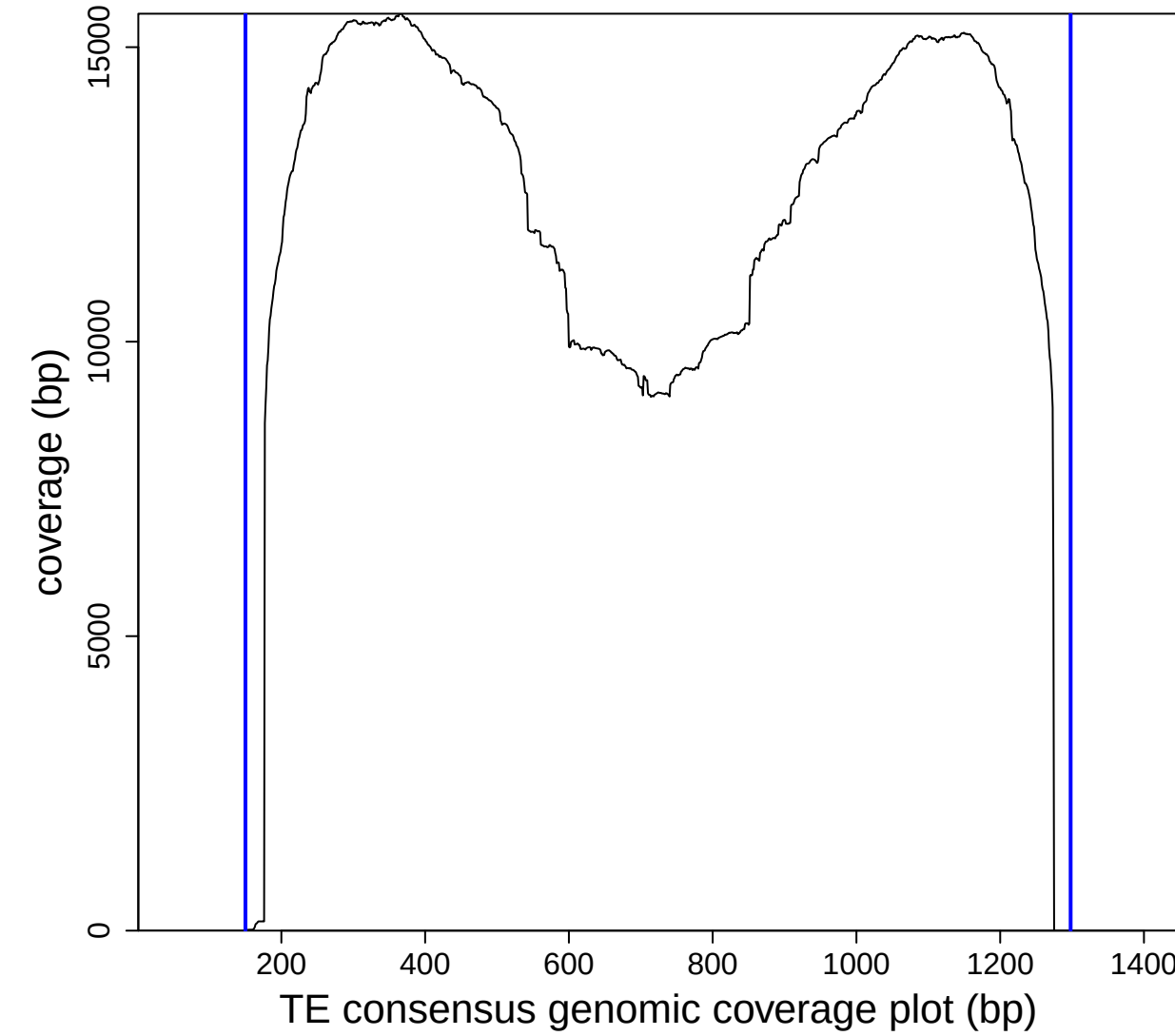
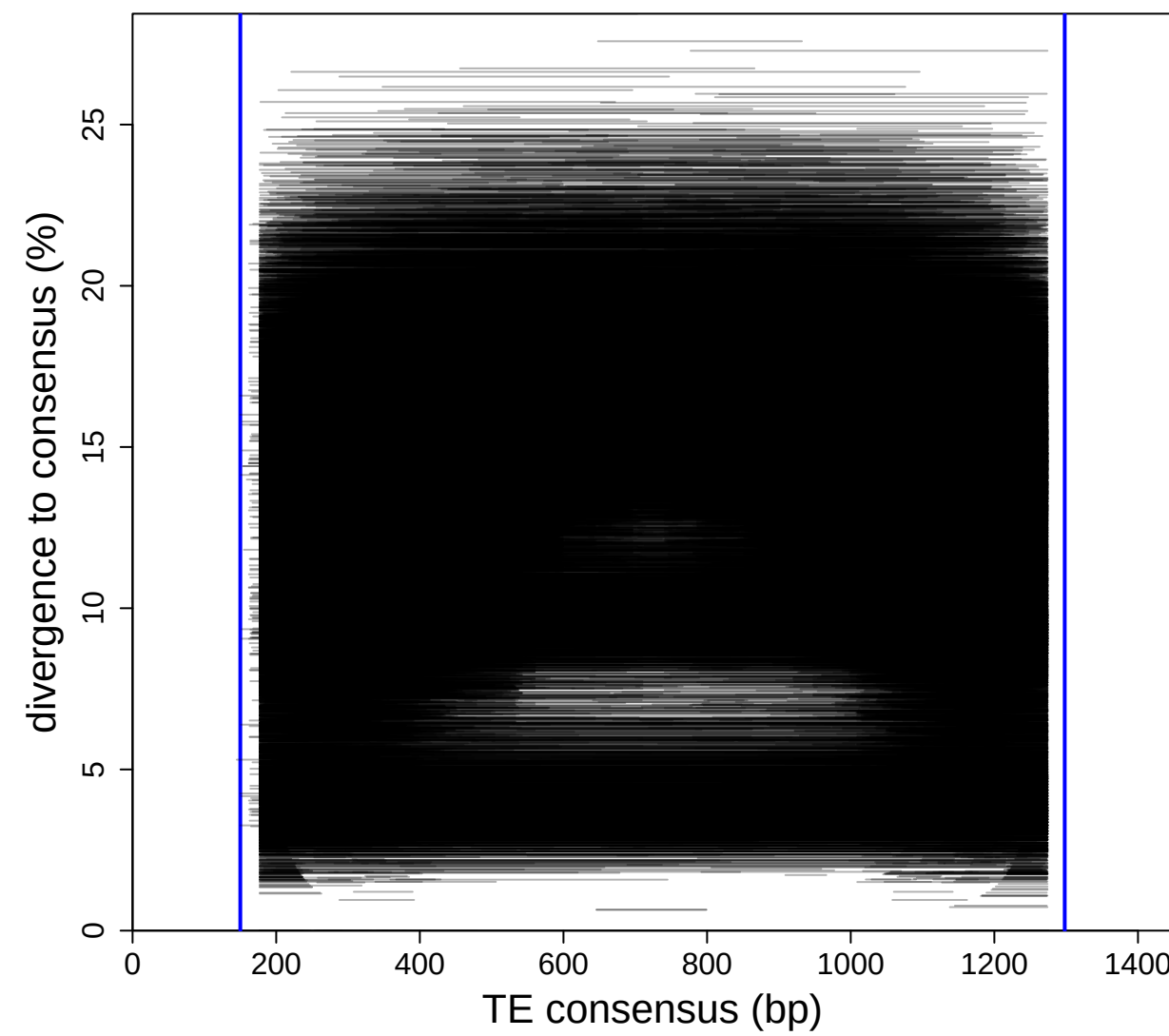
a.b.bed_uf.bed_g_3.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.f
size: 1141bp; fragments: 31058; full length: 3976 (≥ 1026.9 bp)

After TEtrimmer 1141 bp



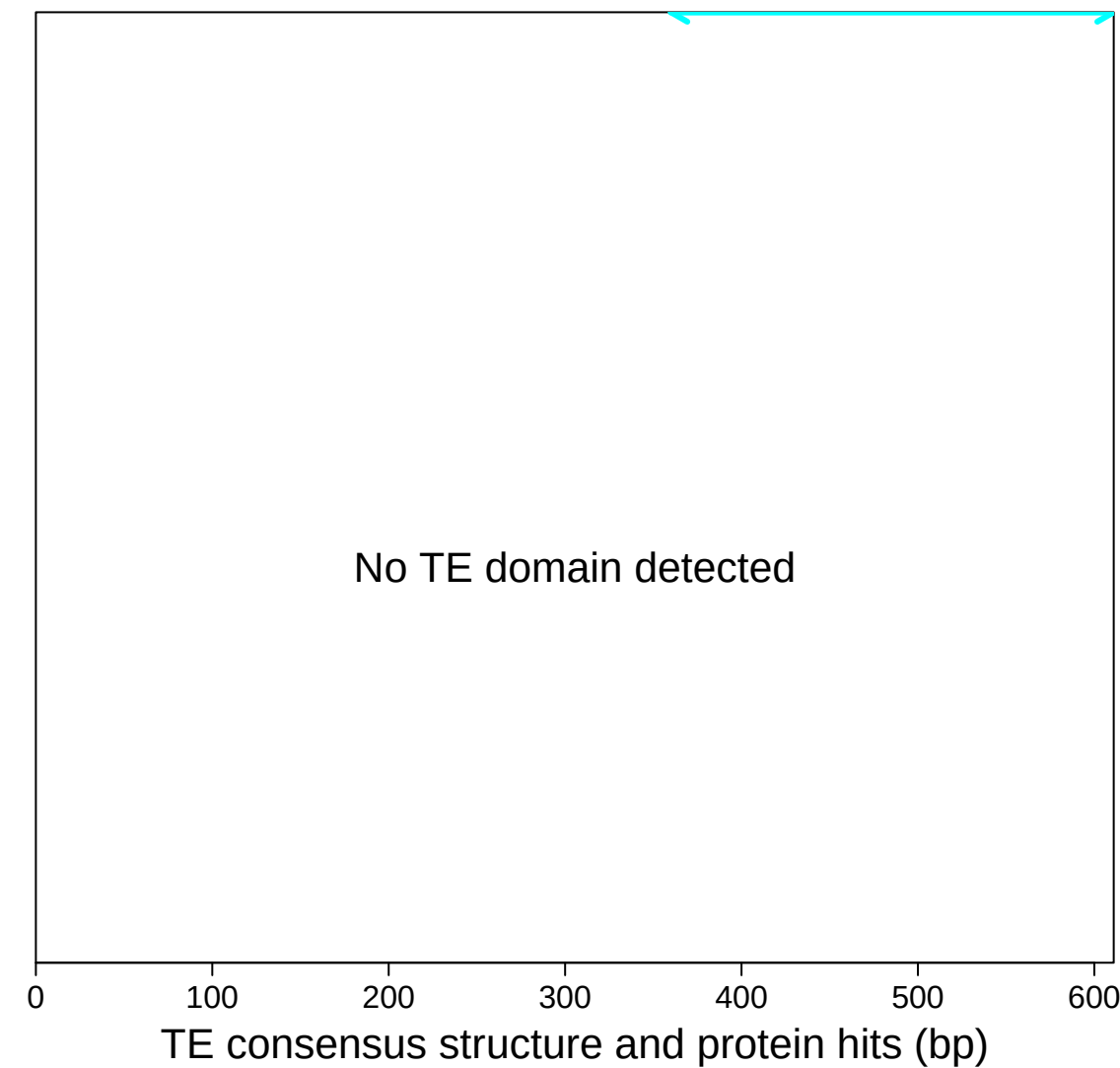
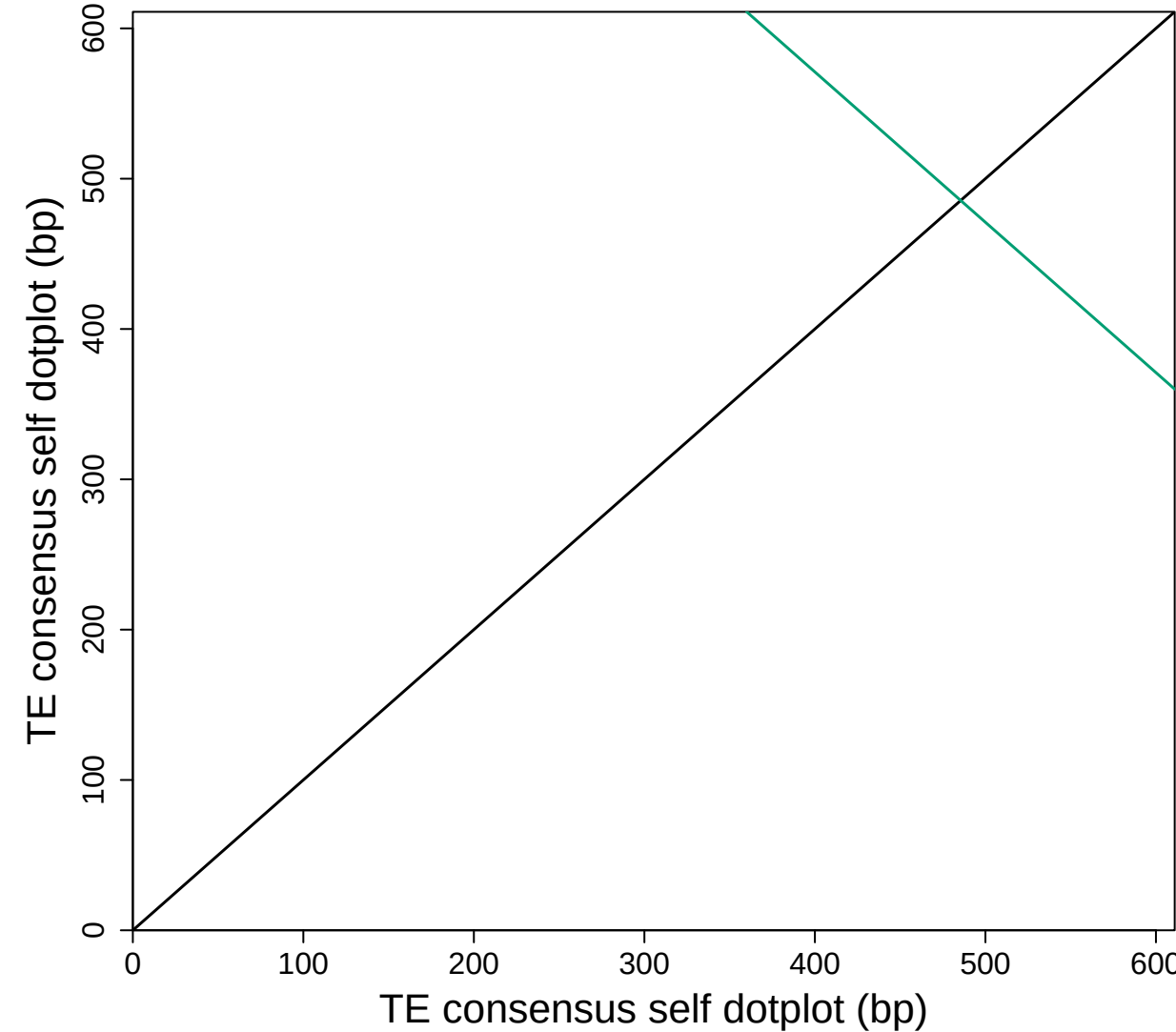
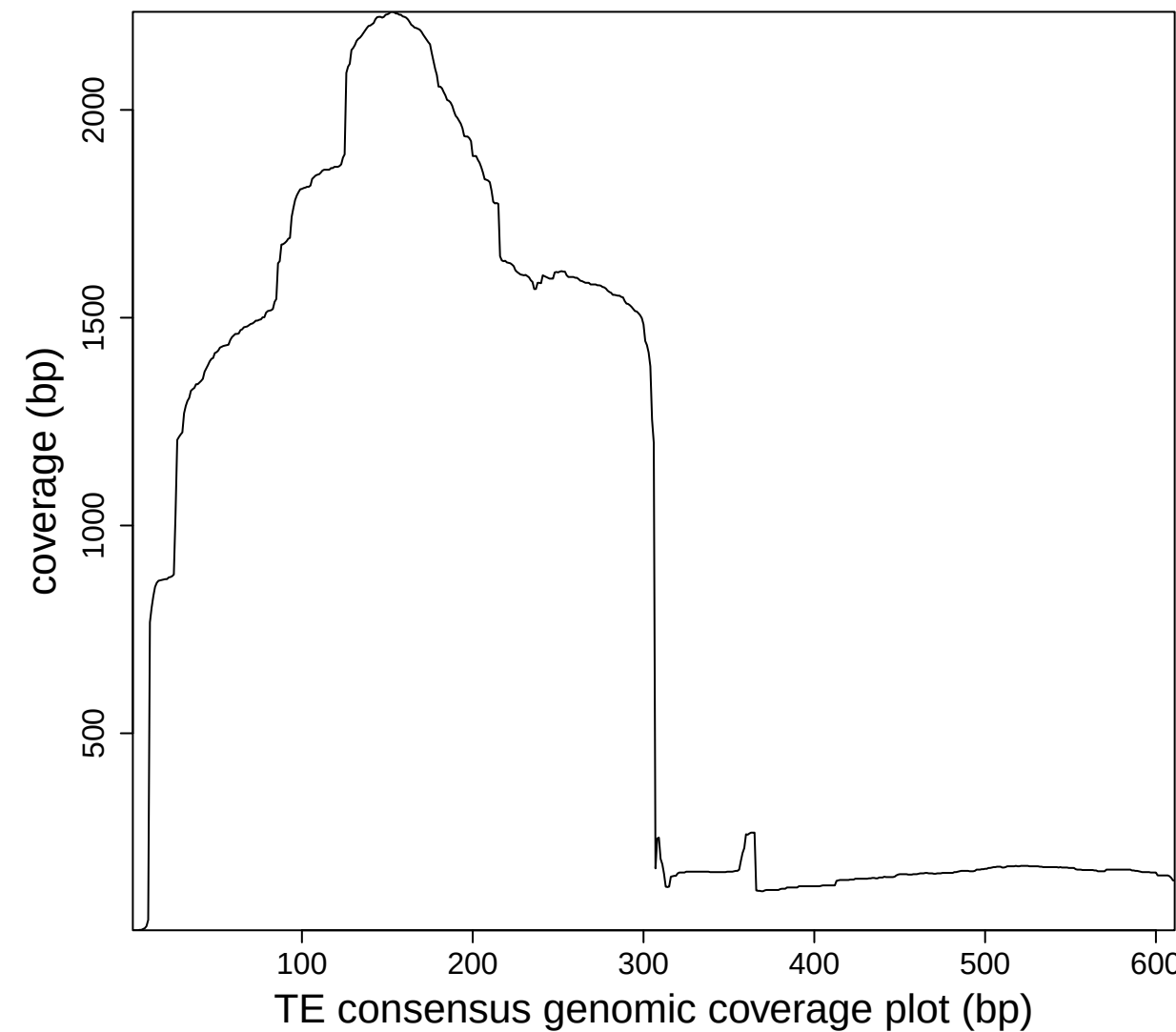
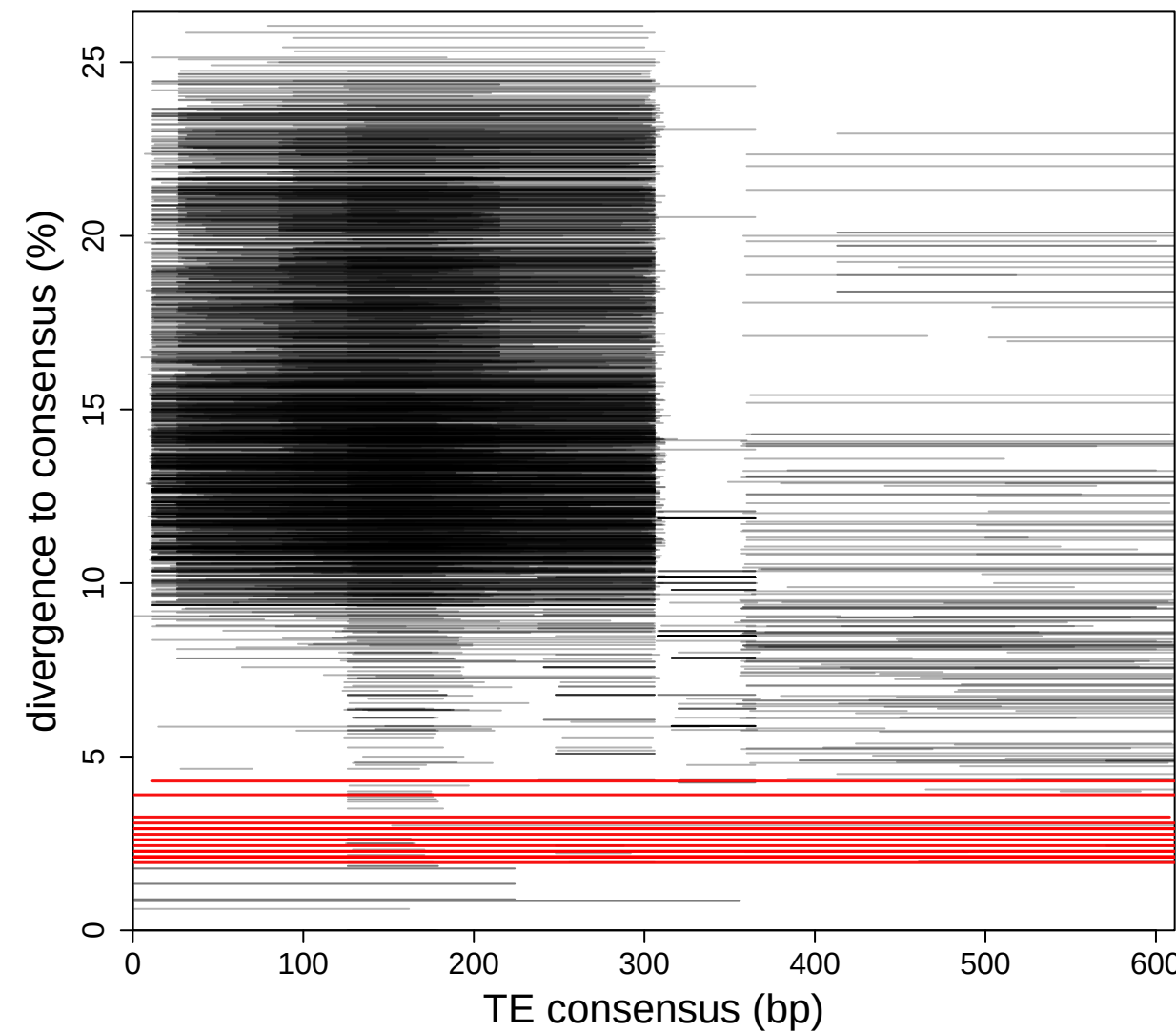
5.fasta.b.bed_uf.bed_g_3.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.f
size: 1448bp; fragments: 30820; full length: 0 (>=1303.2bp)

After TEtrimmer Extended plot Blue lines are boundaries

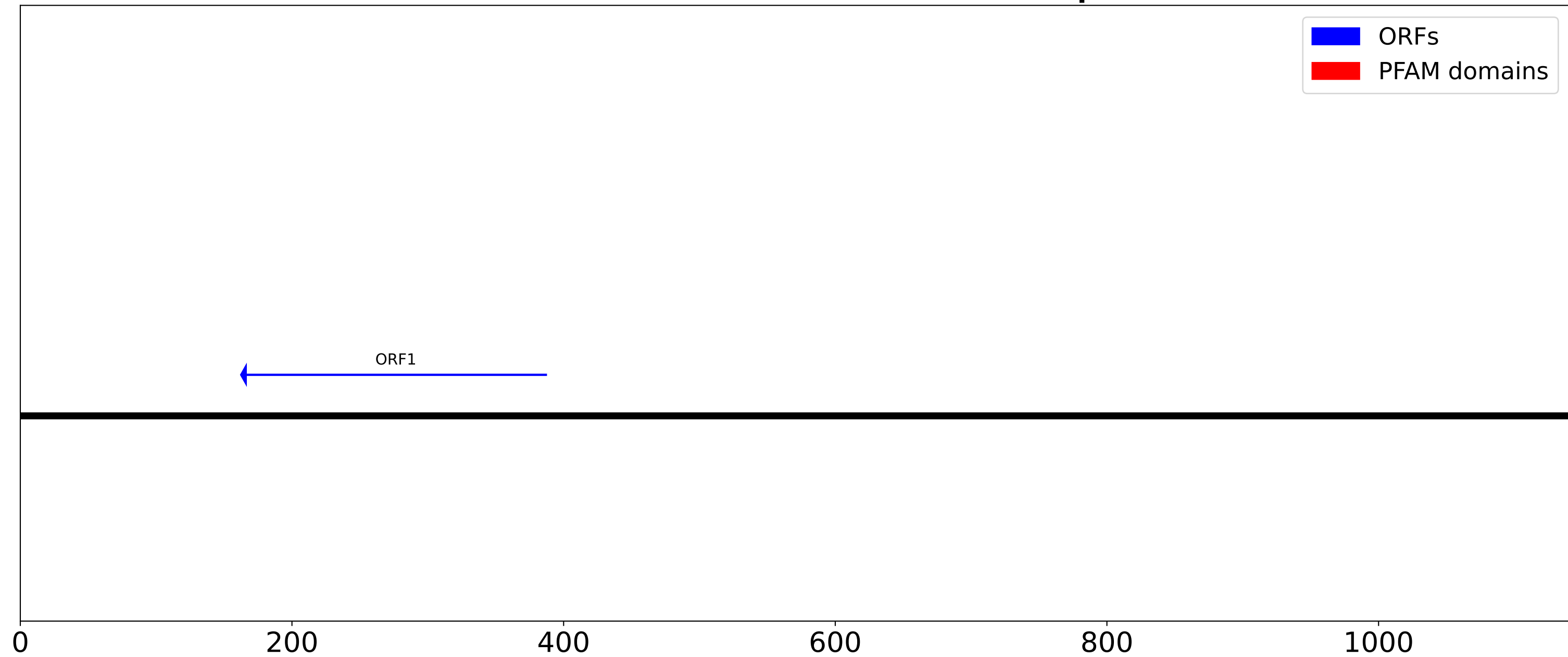


TE: rnd_5_family_2045
size: 611bp; fragments: 2650; full length: 17 (≥ 549.9 bp)

Before TEtrimmer 611 bp



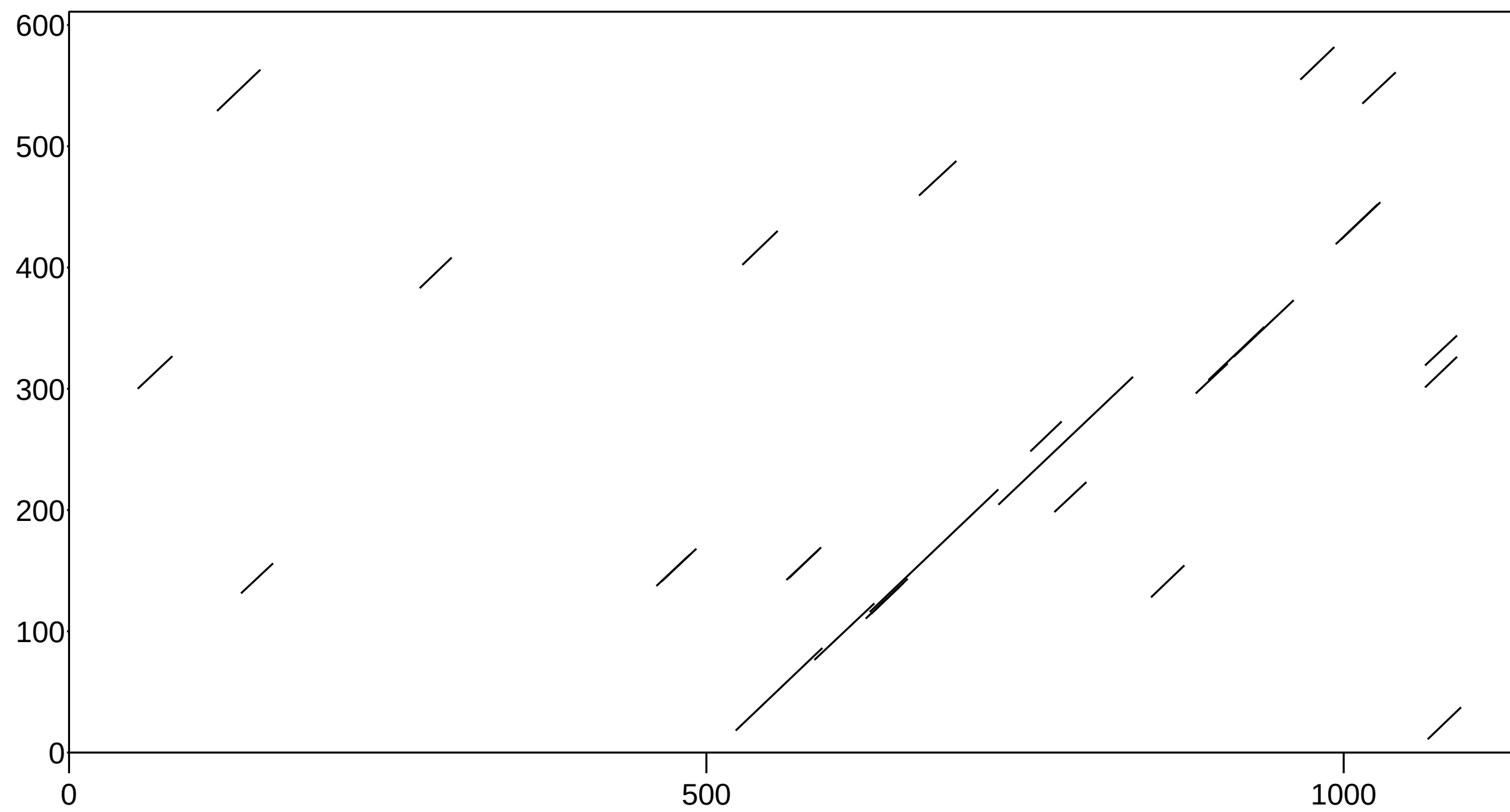
After TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)