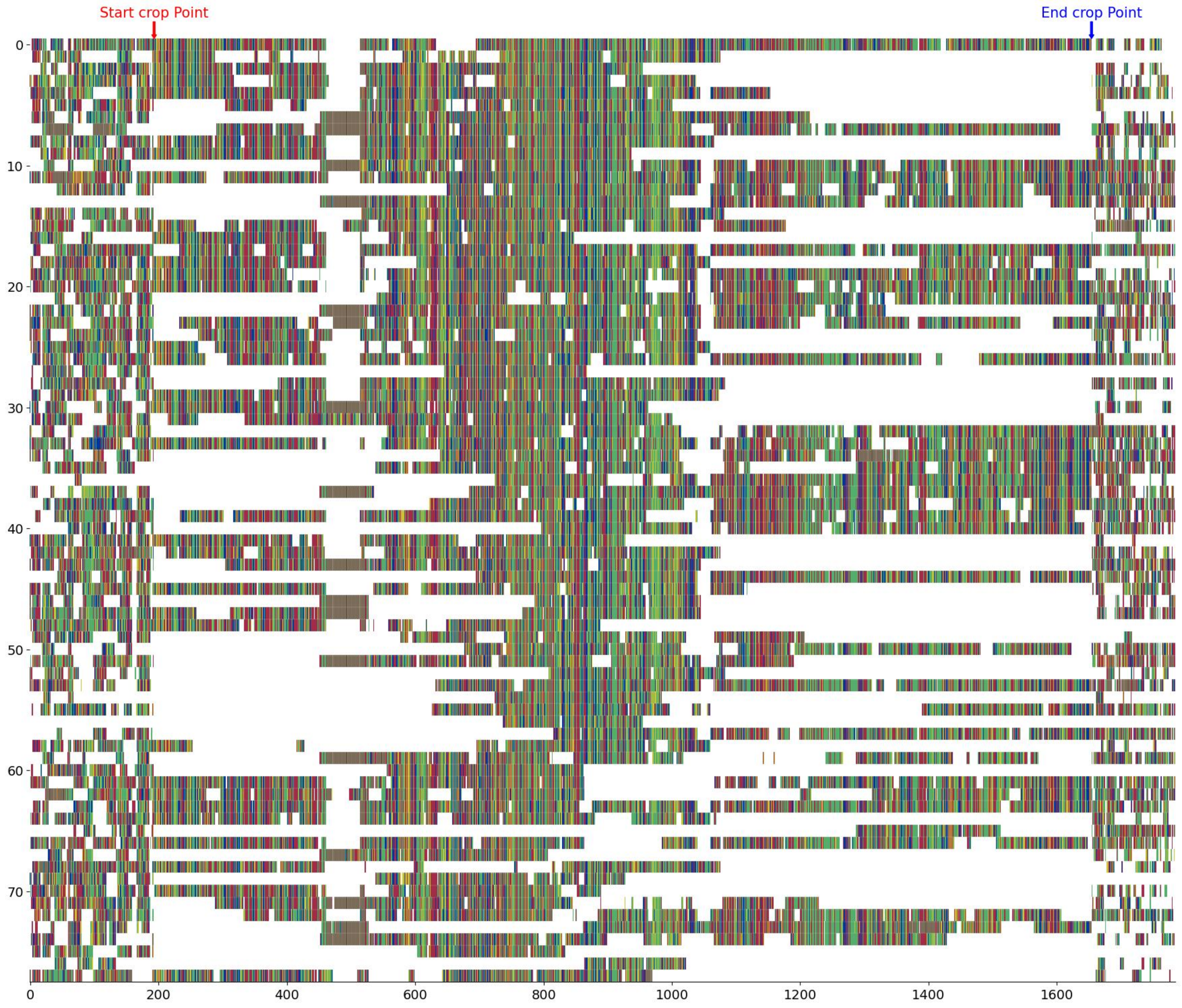




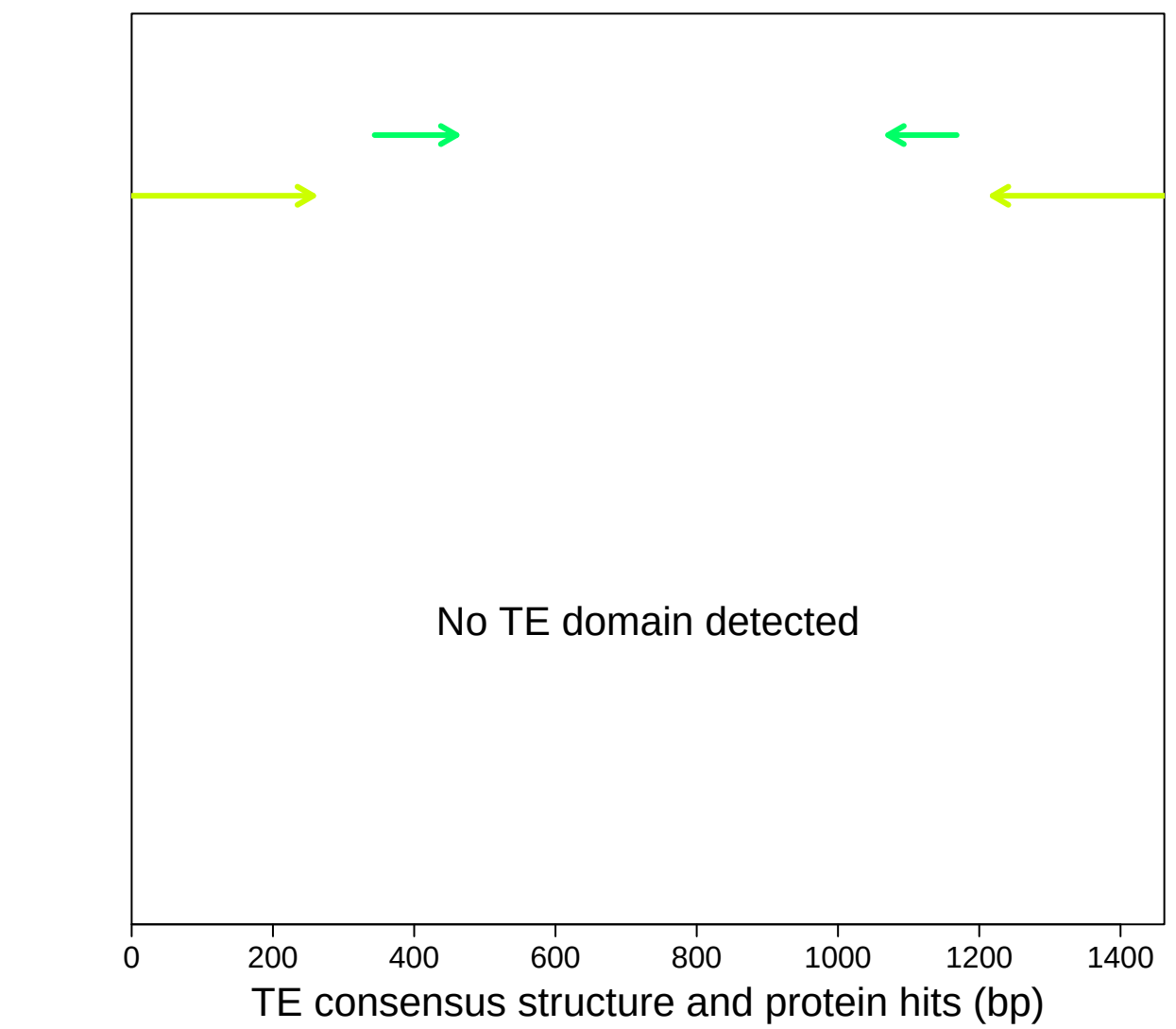
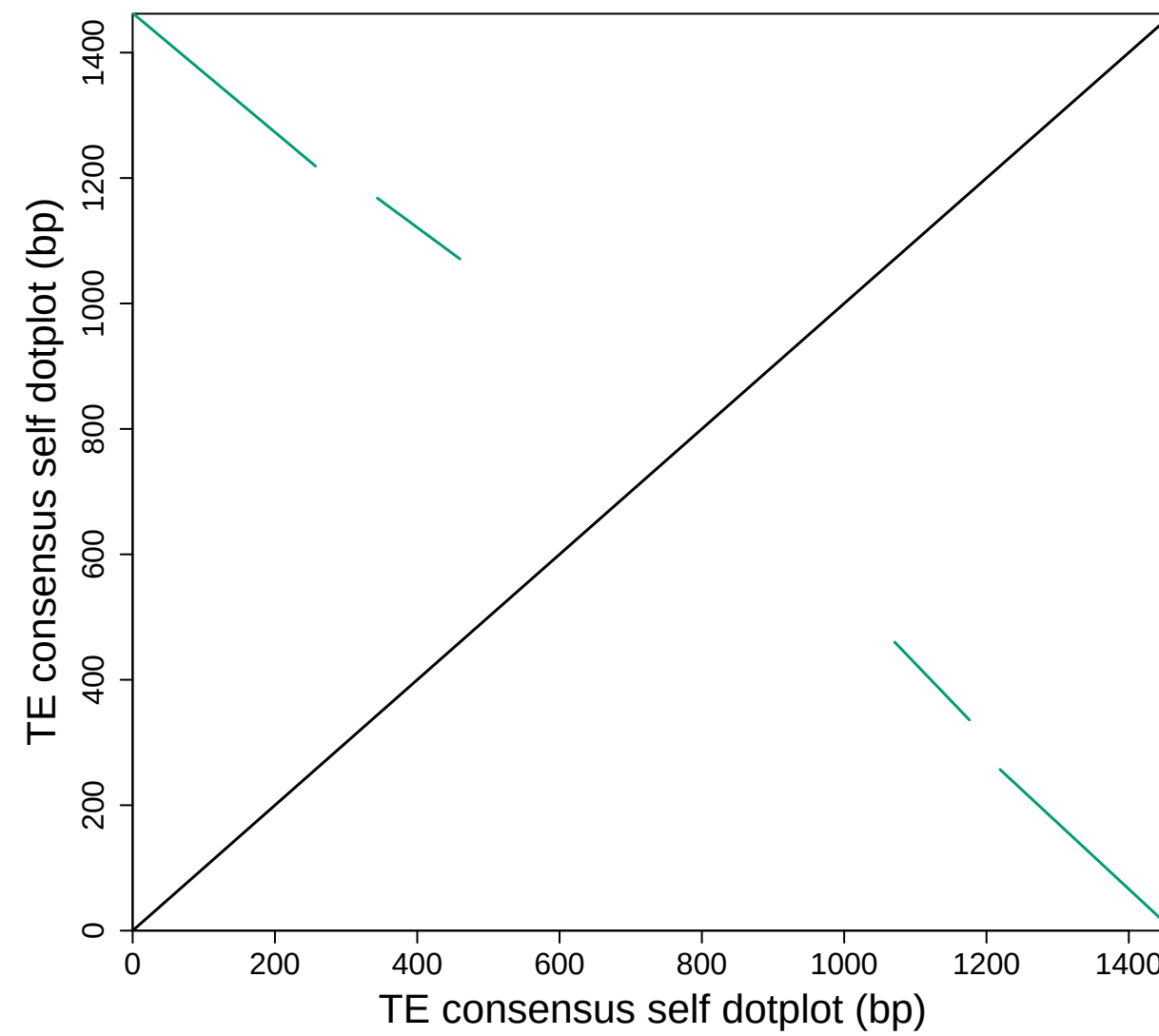
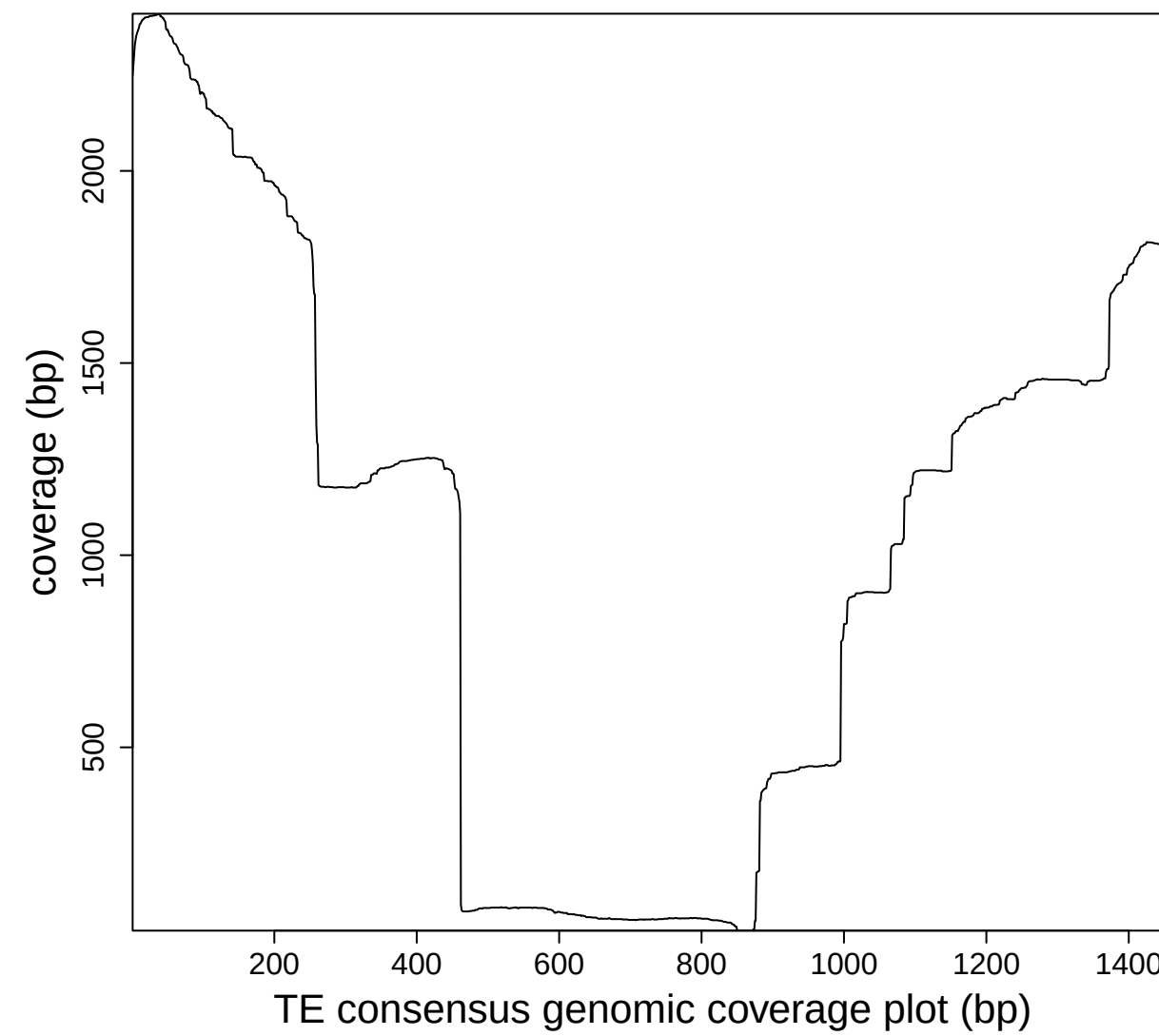
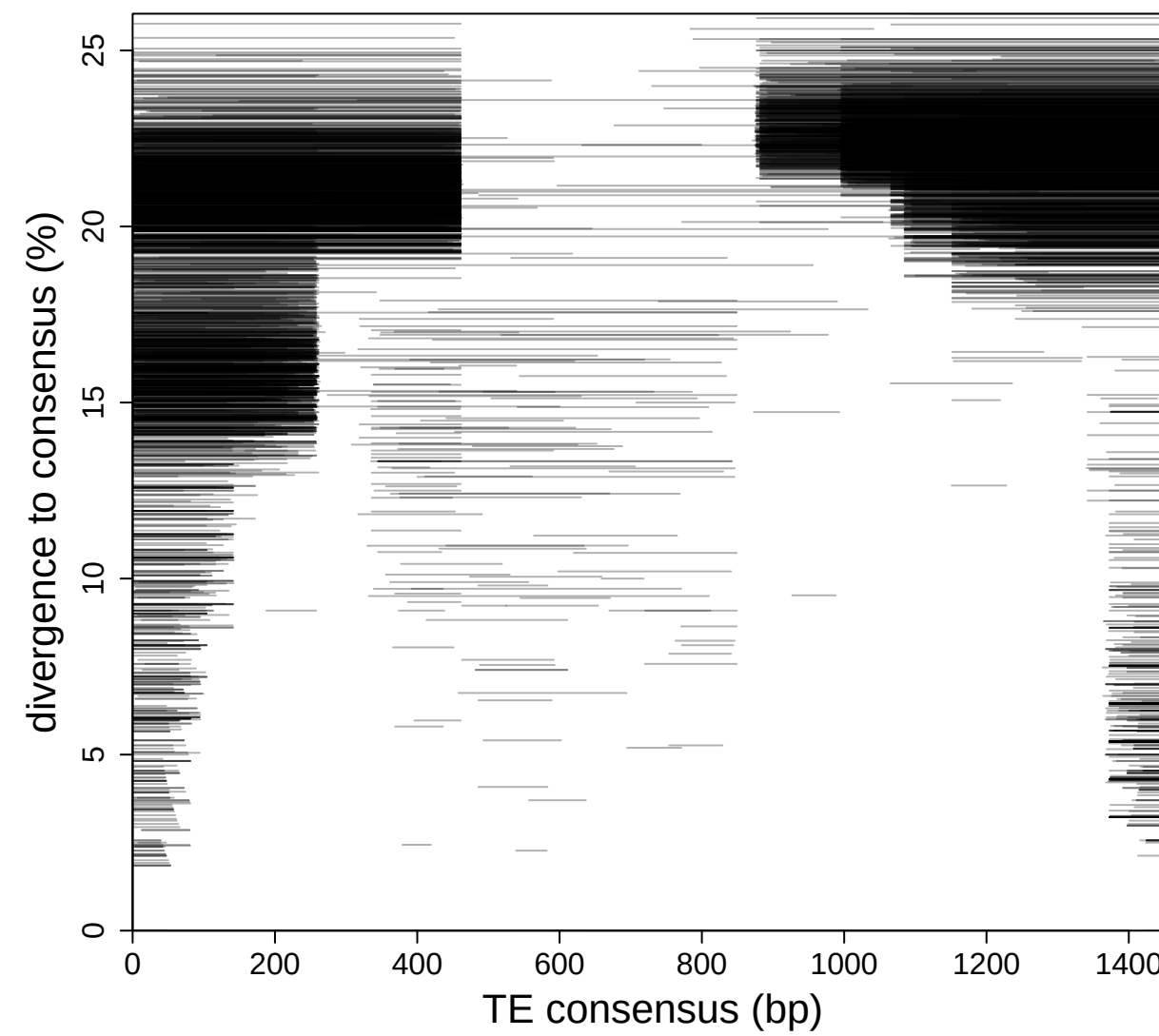
Start crop Point

End crop Point



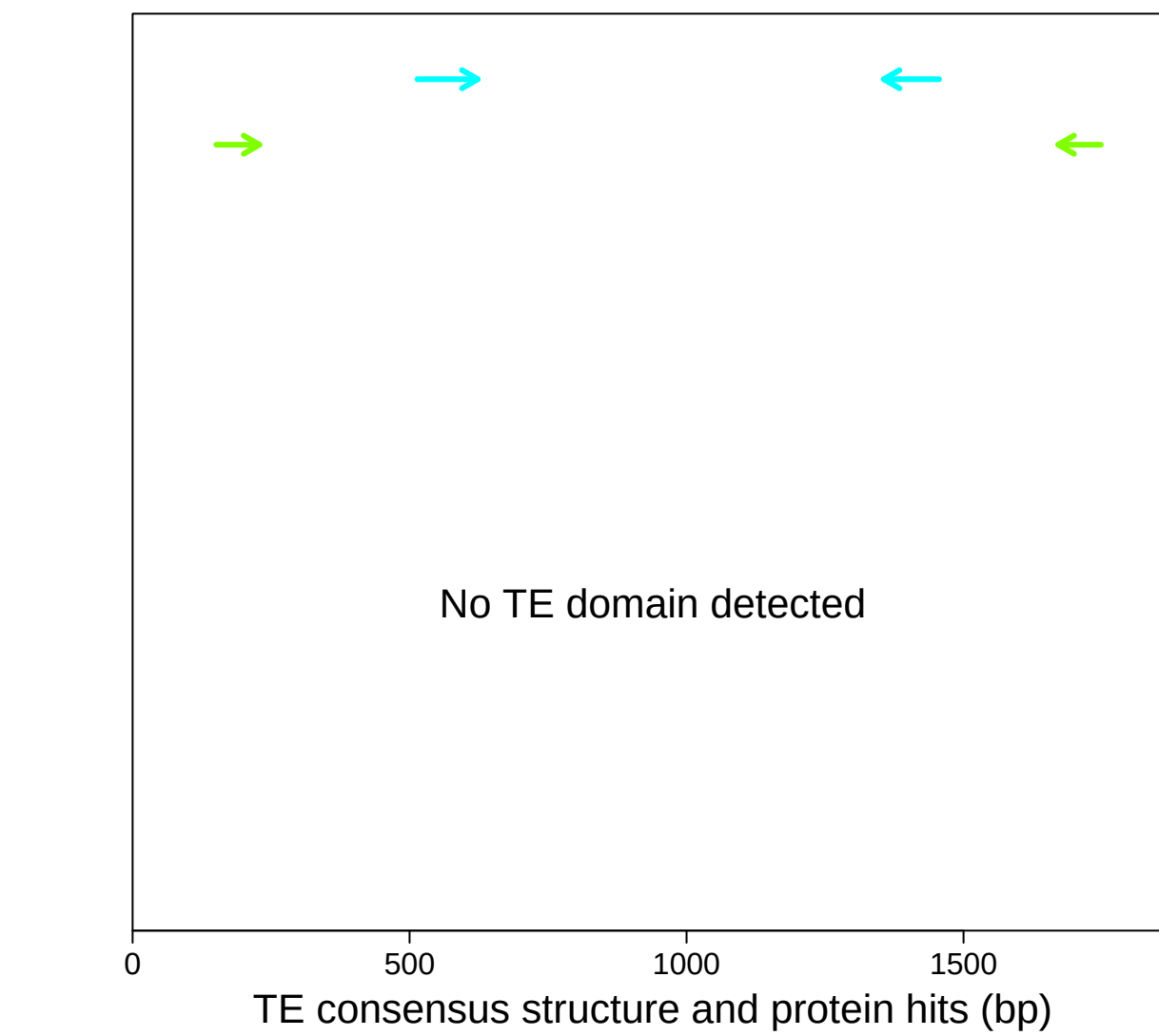
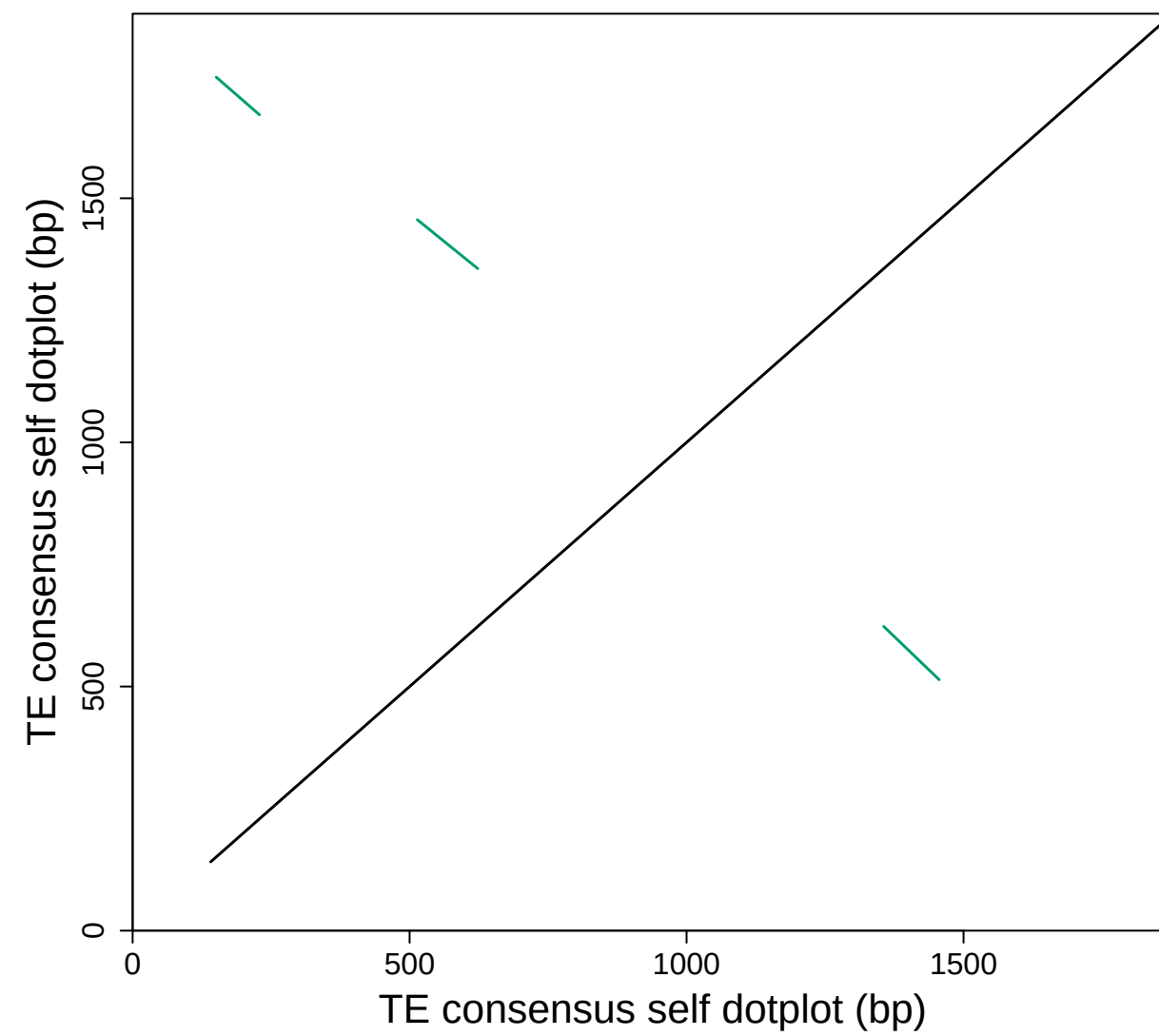
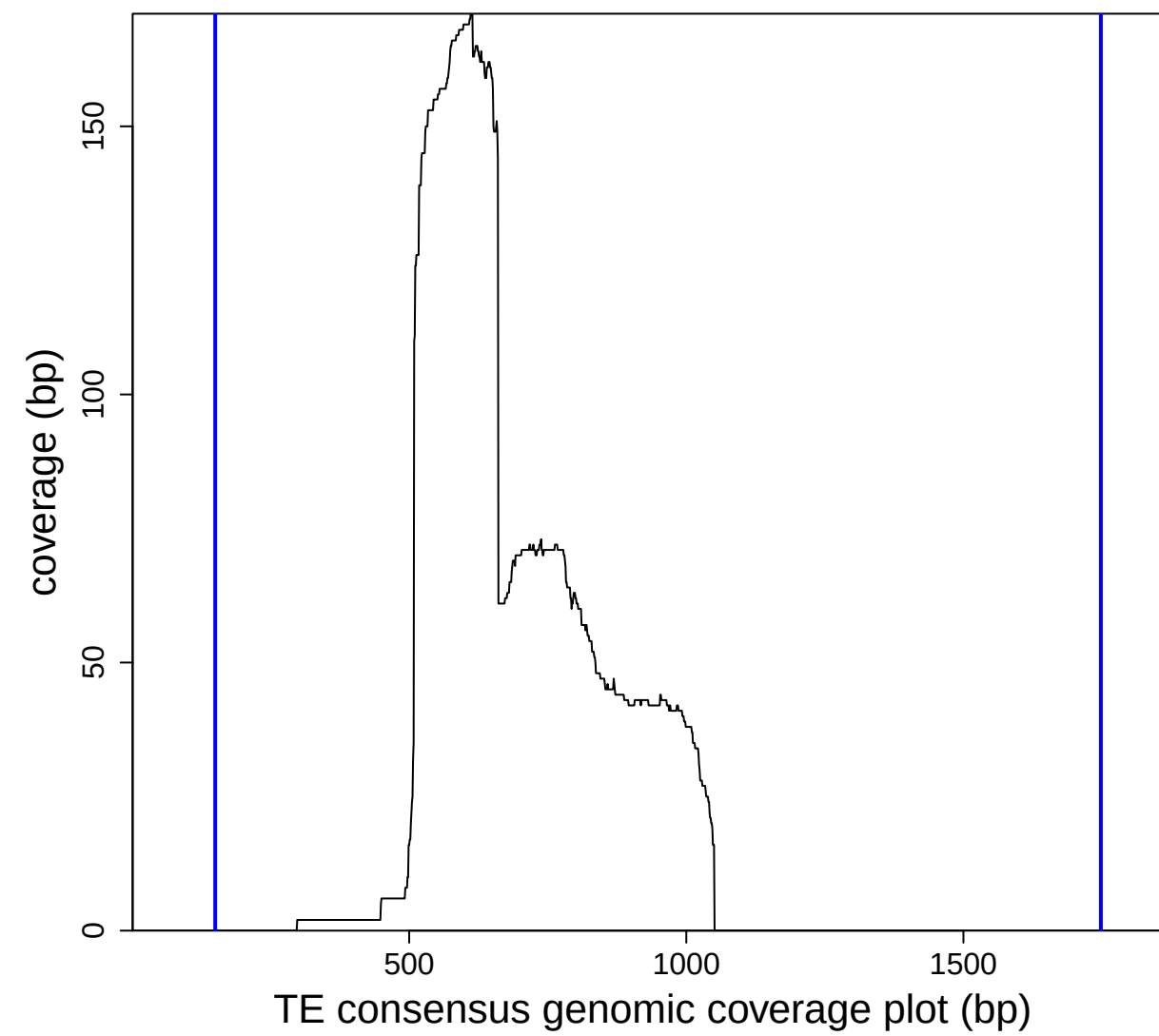
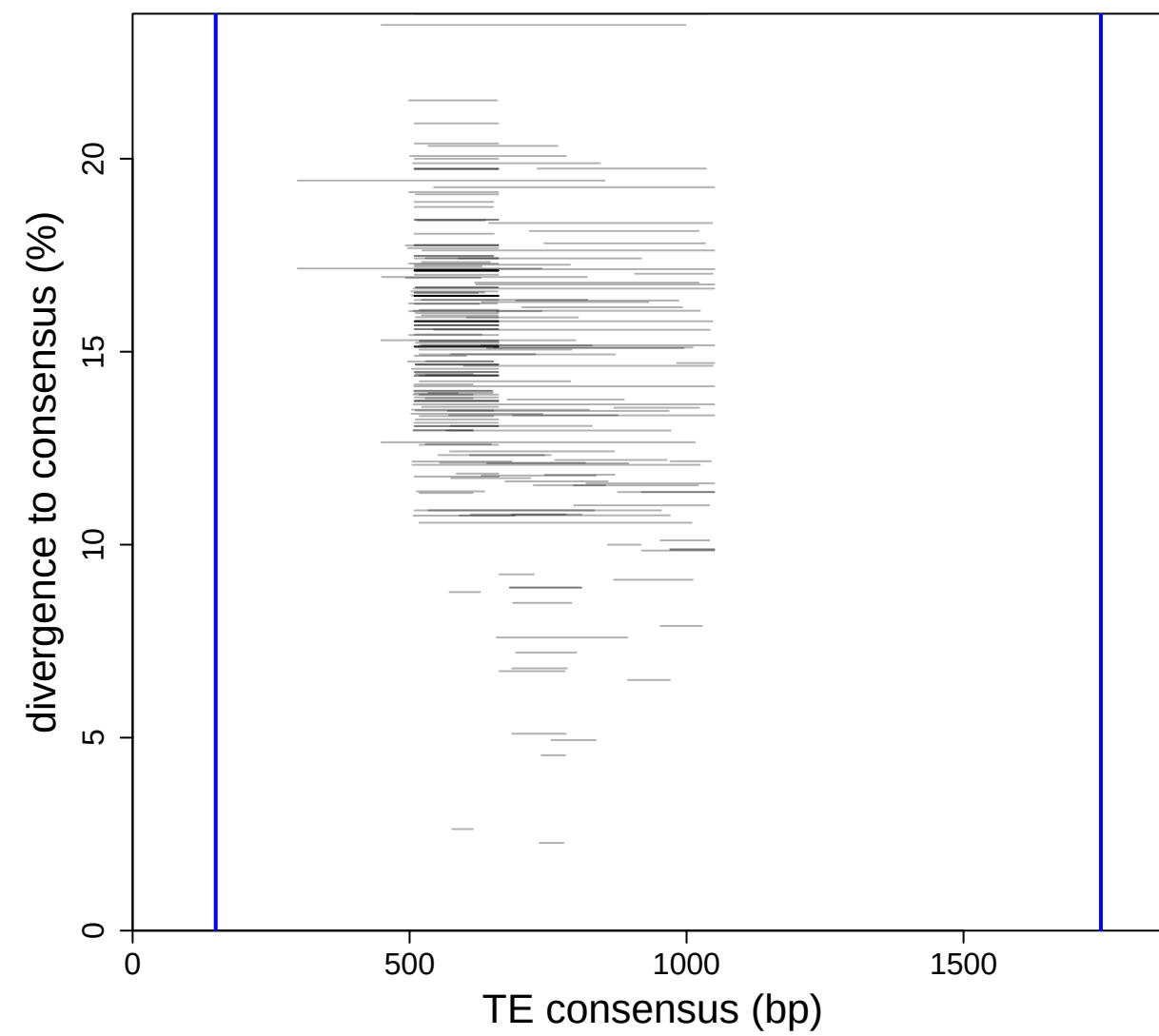
bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.fa_c
size: 1462bp; fragments: 4426; full length: 0 (>=1315.8bp)

After TETrimmer 1462 bp

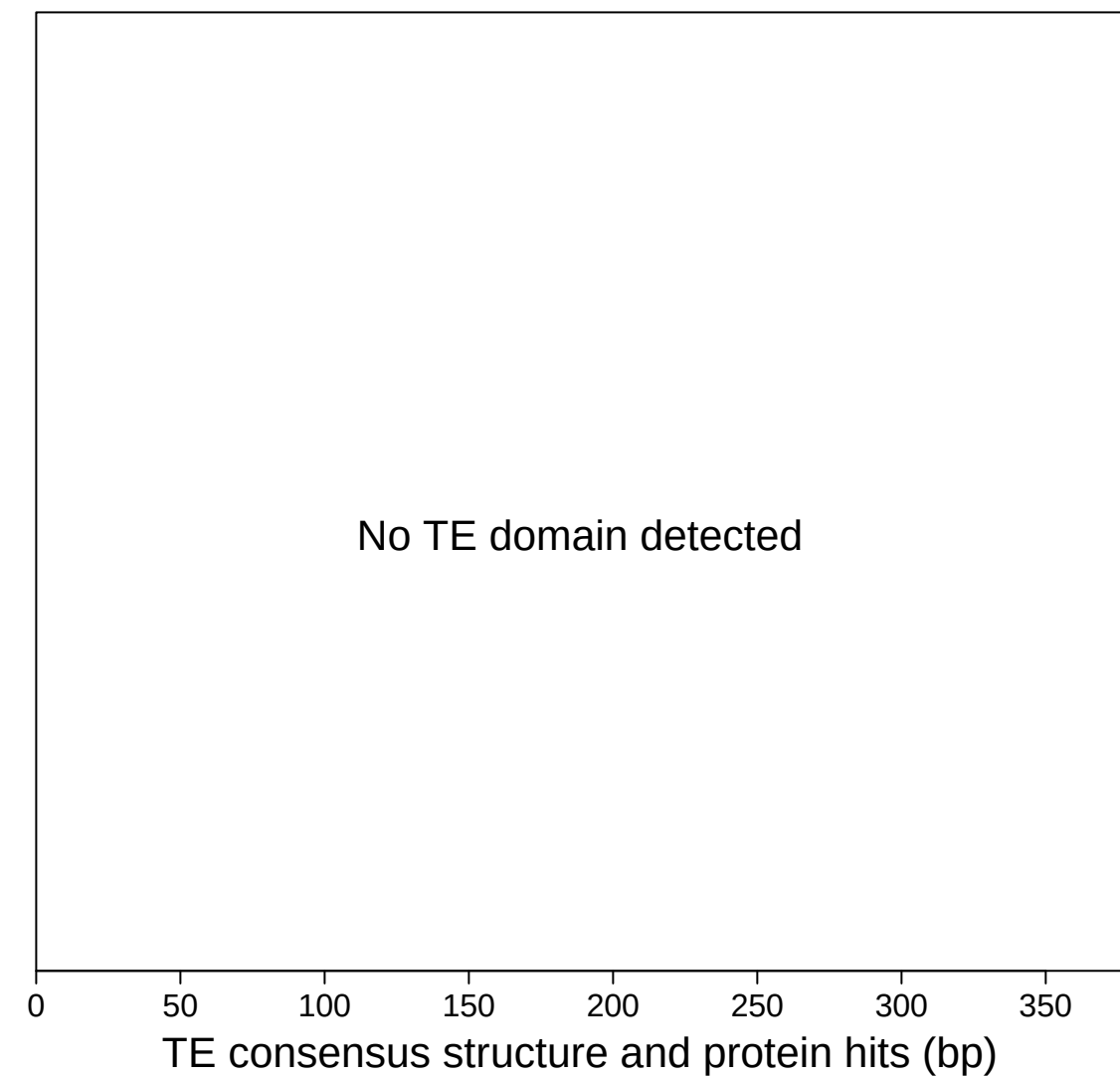
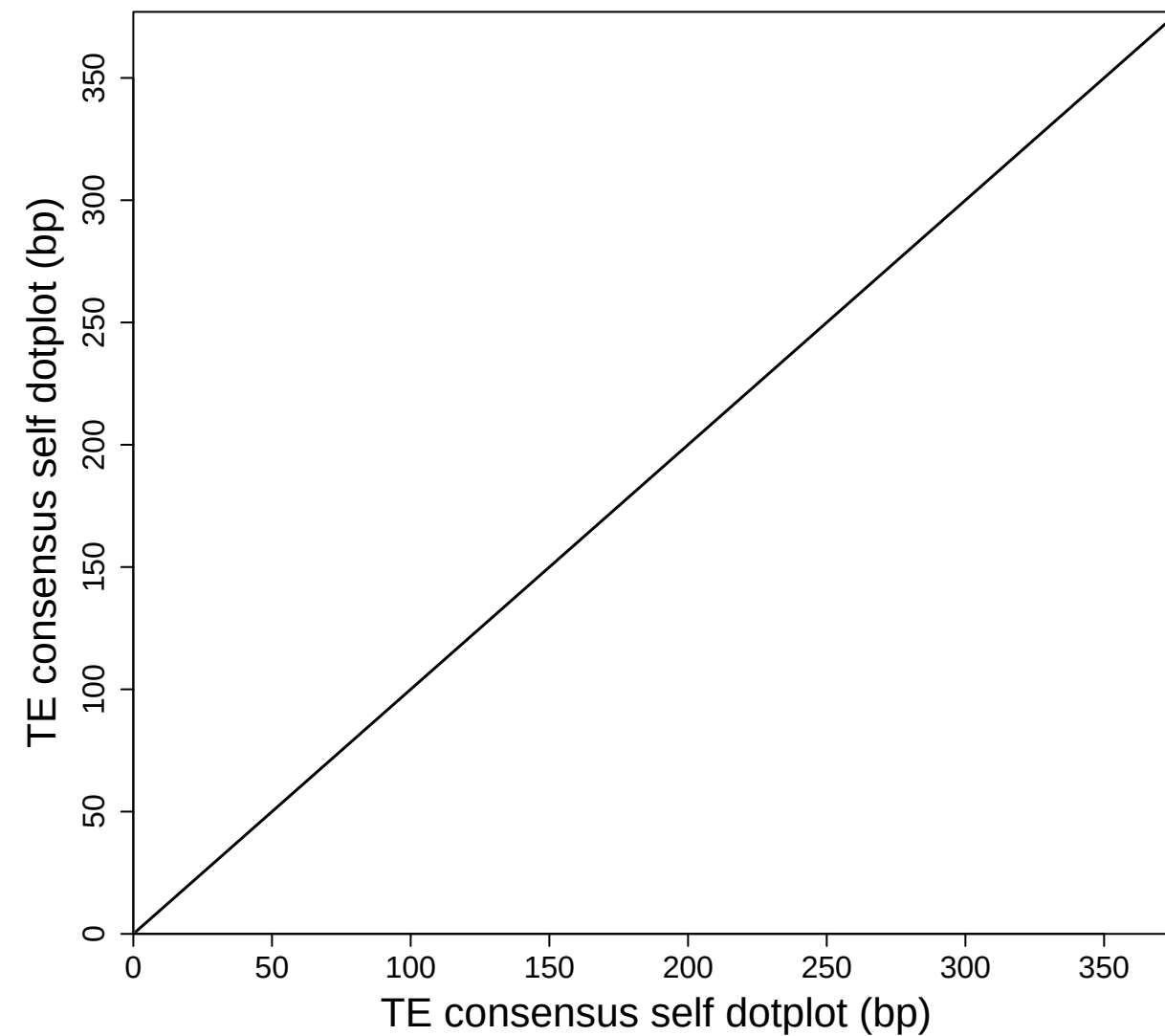
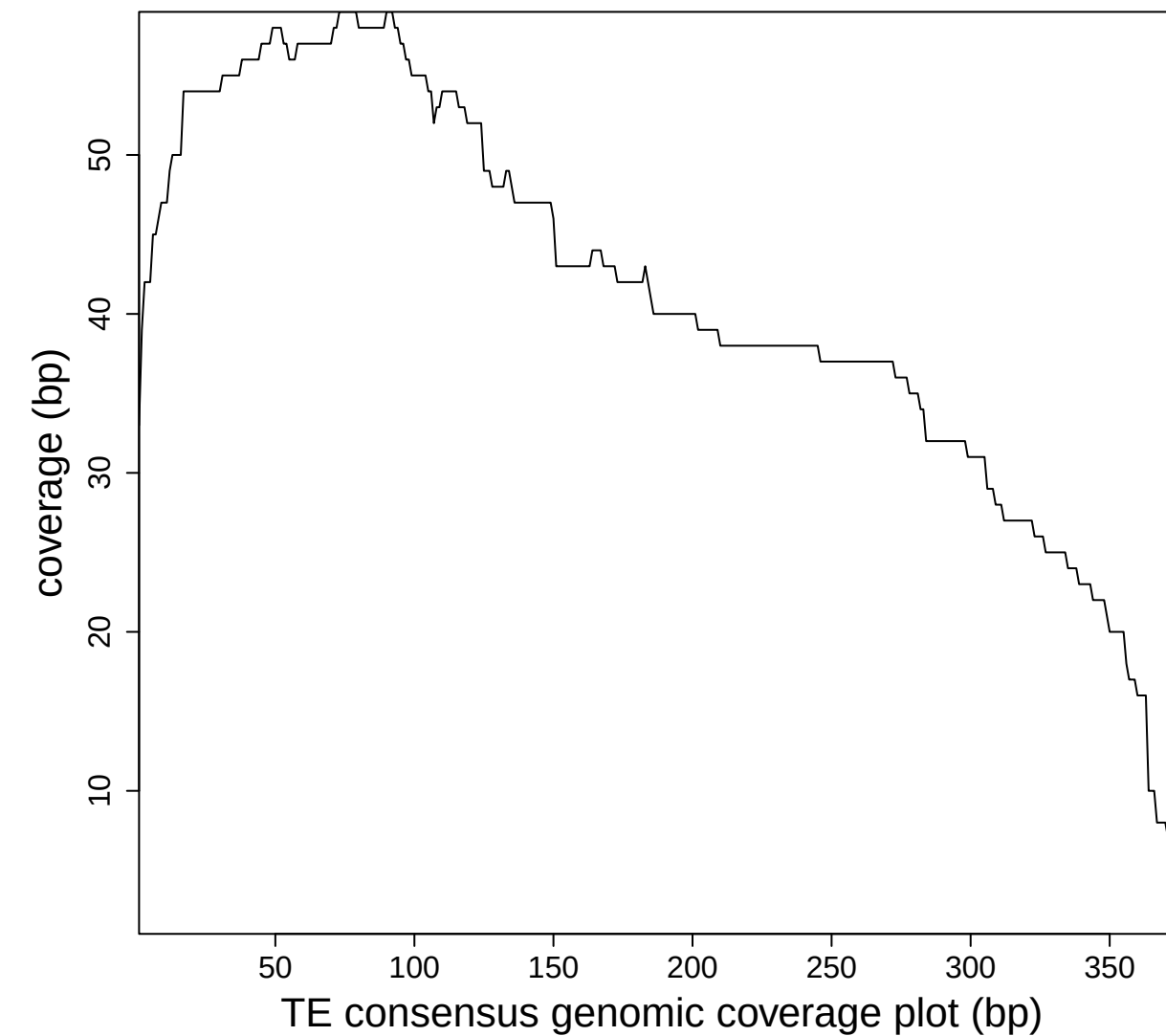
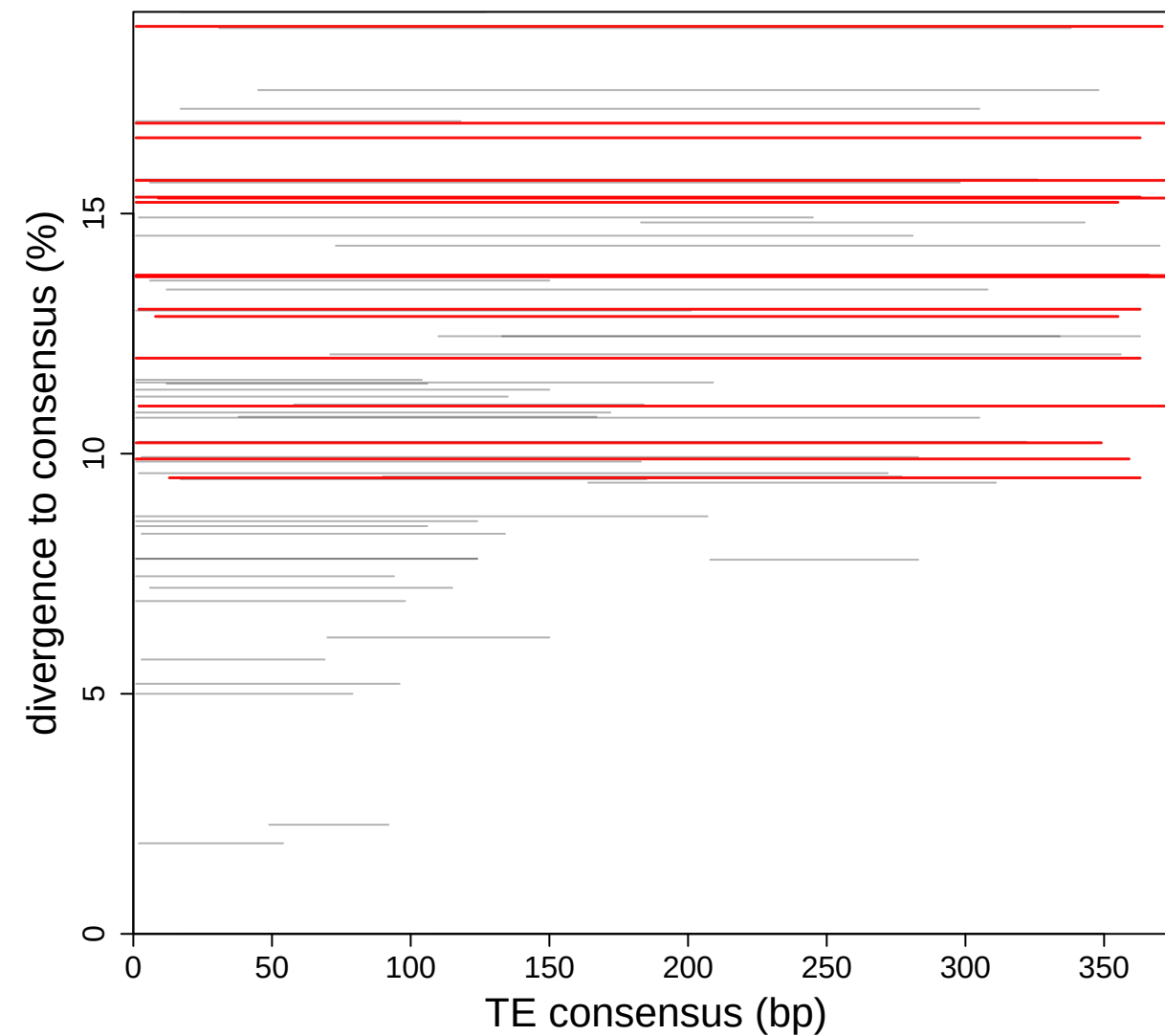


ly_5044.fasta.b.bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.f
size: 1878bp; fragments: 225; full length: 0 (>=1690.2bp)

After TEtrimmer Extended plot Blue lines are boundaries

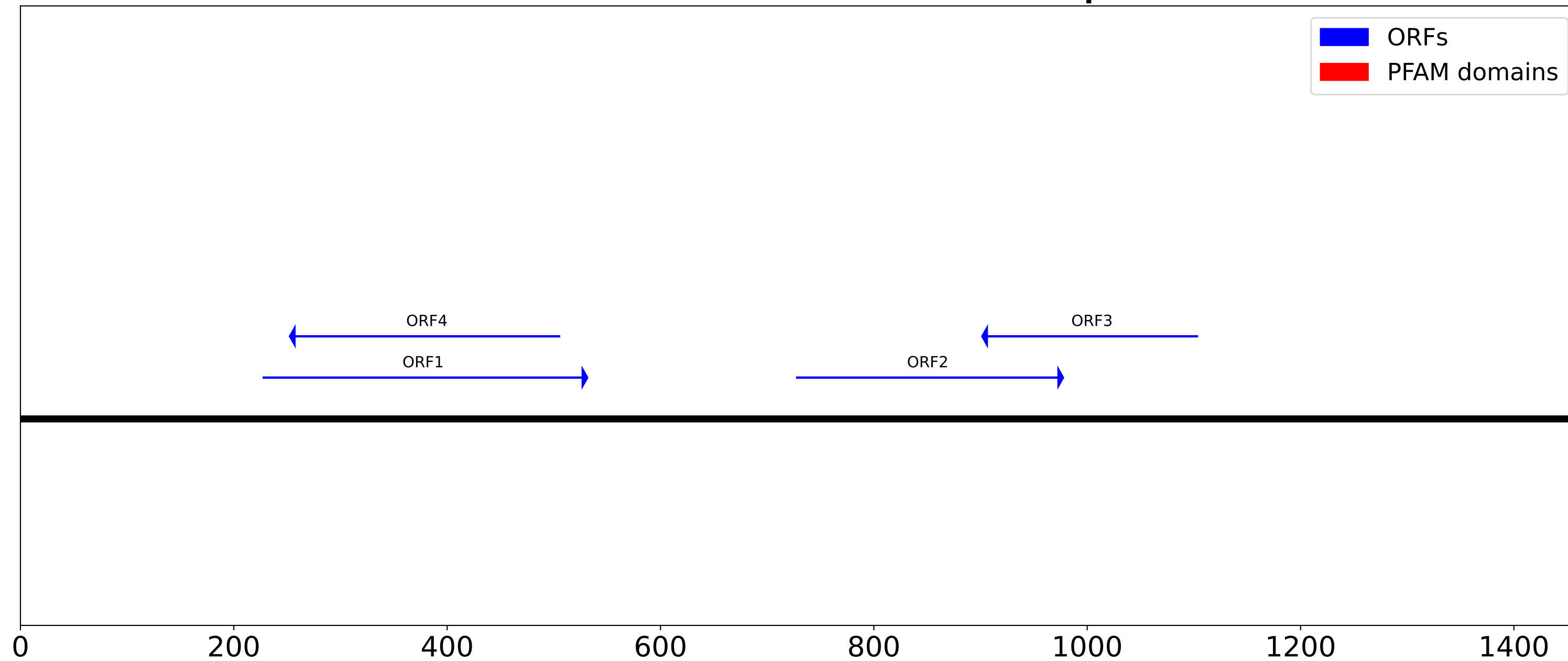


TE: rnd_5_family_5044
size: 377bp; fragments: 69; full length: 18 (≥ 339.3 bp)

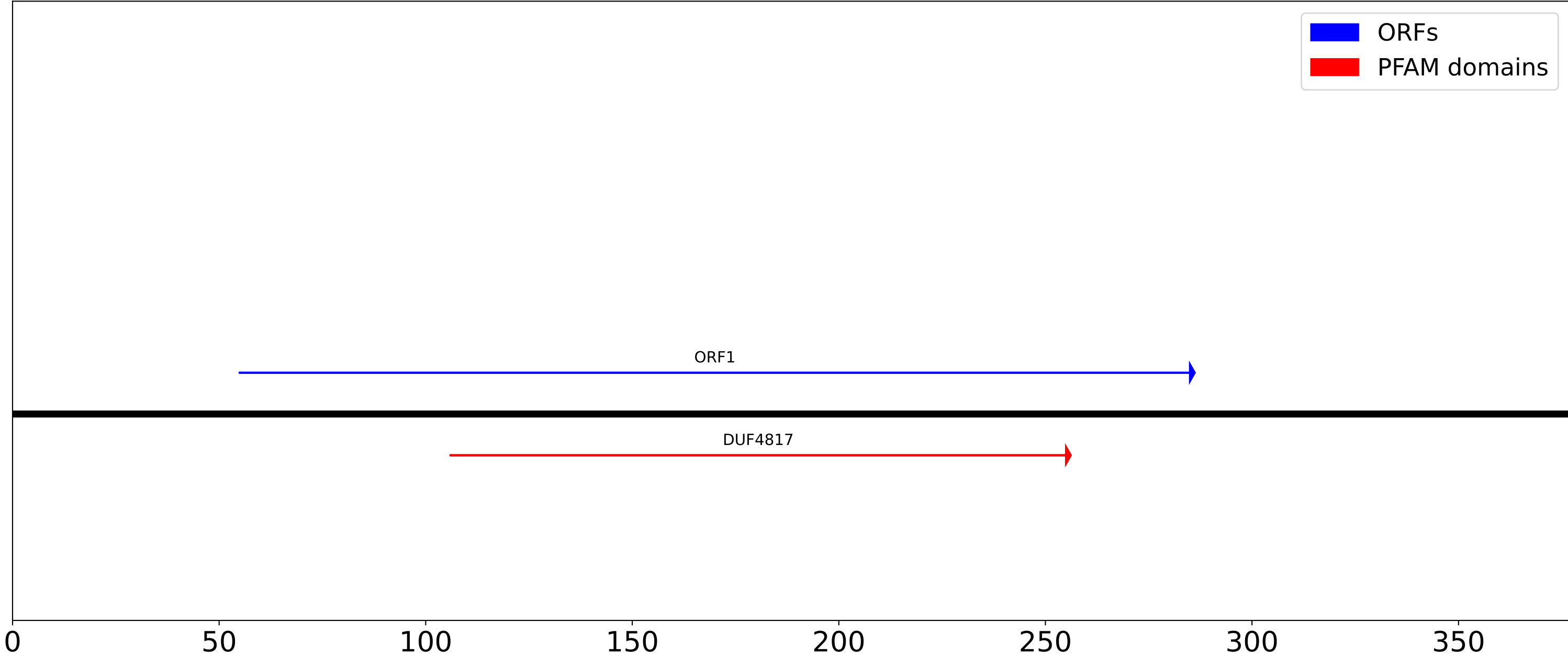


Before TEtrimmer 377 bp

After TEtrimmer ORF and PFAM domain plot



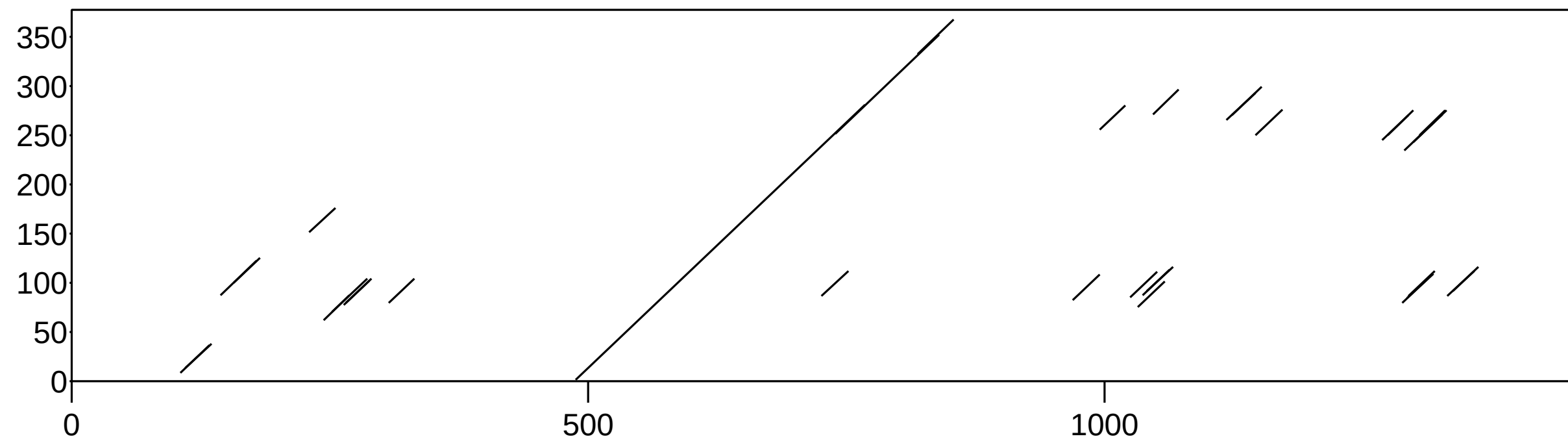
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)