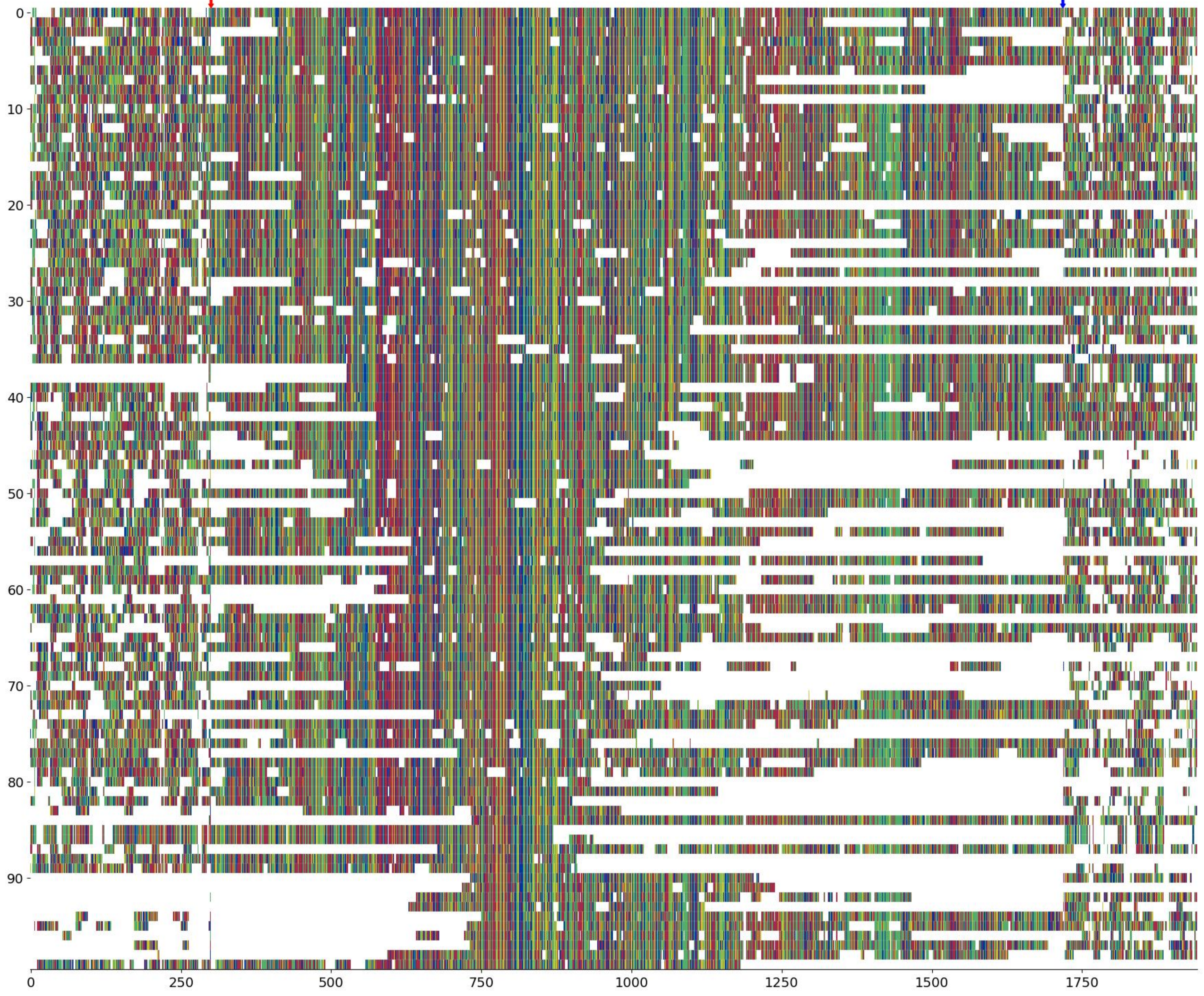




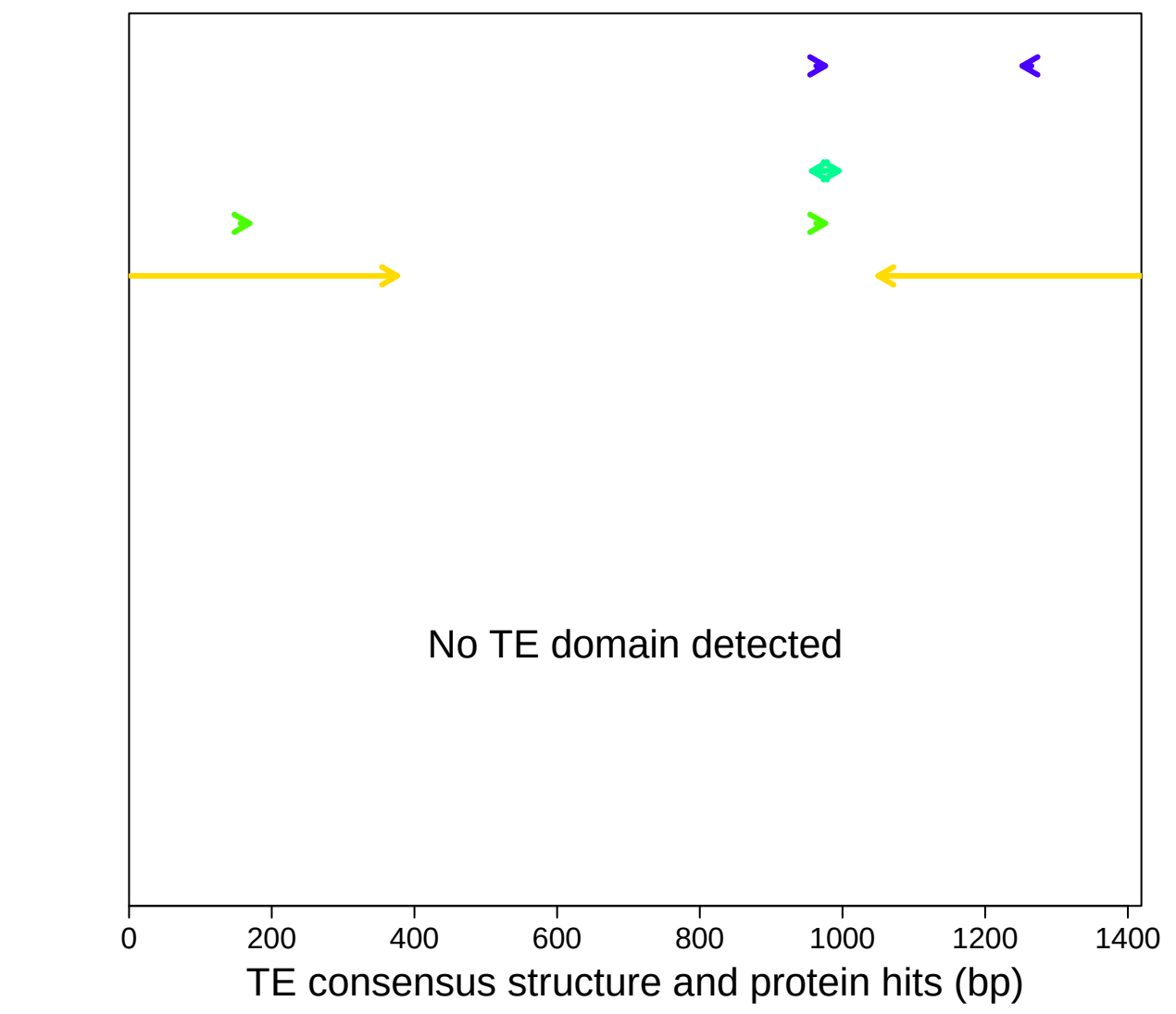
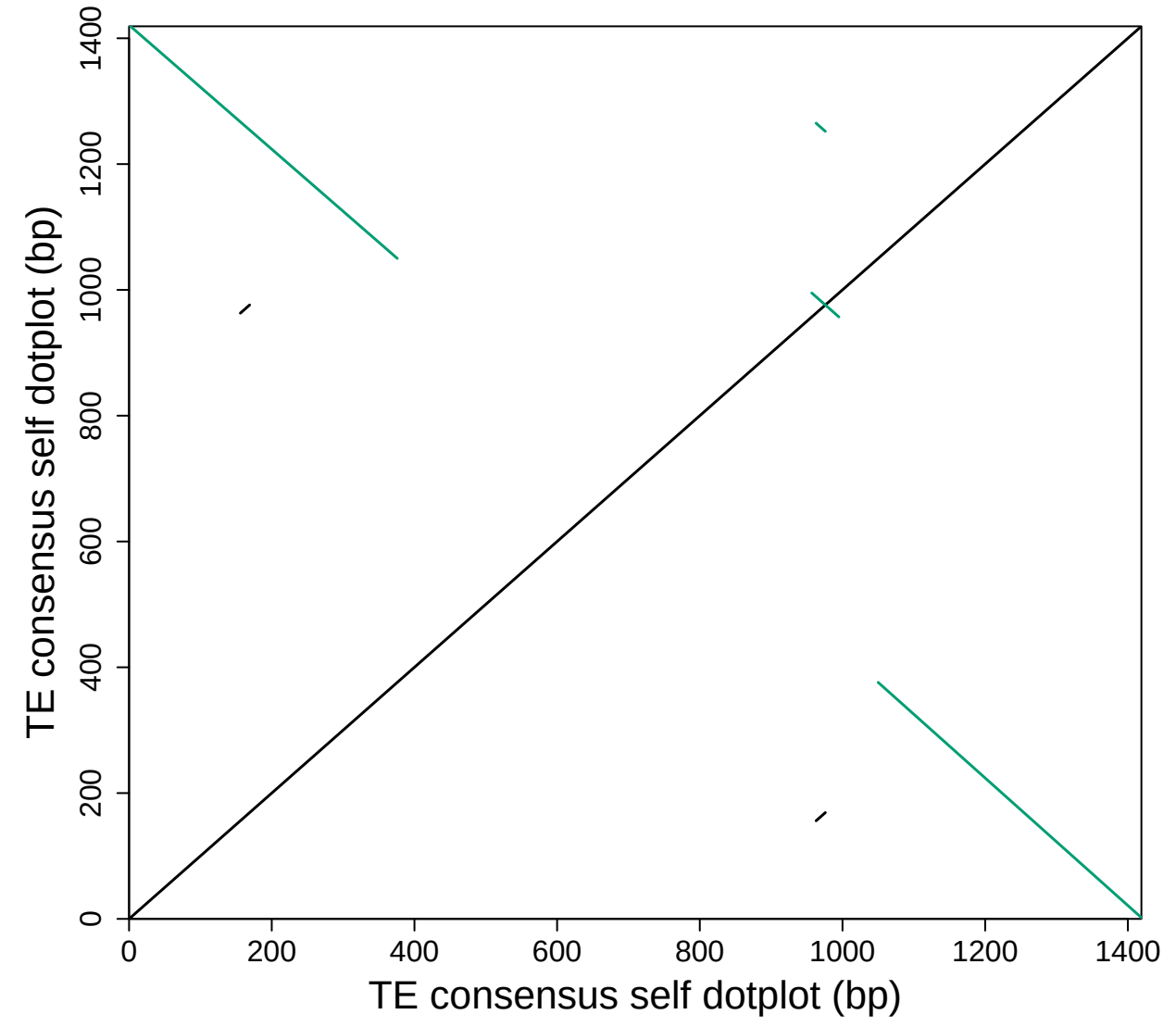
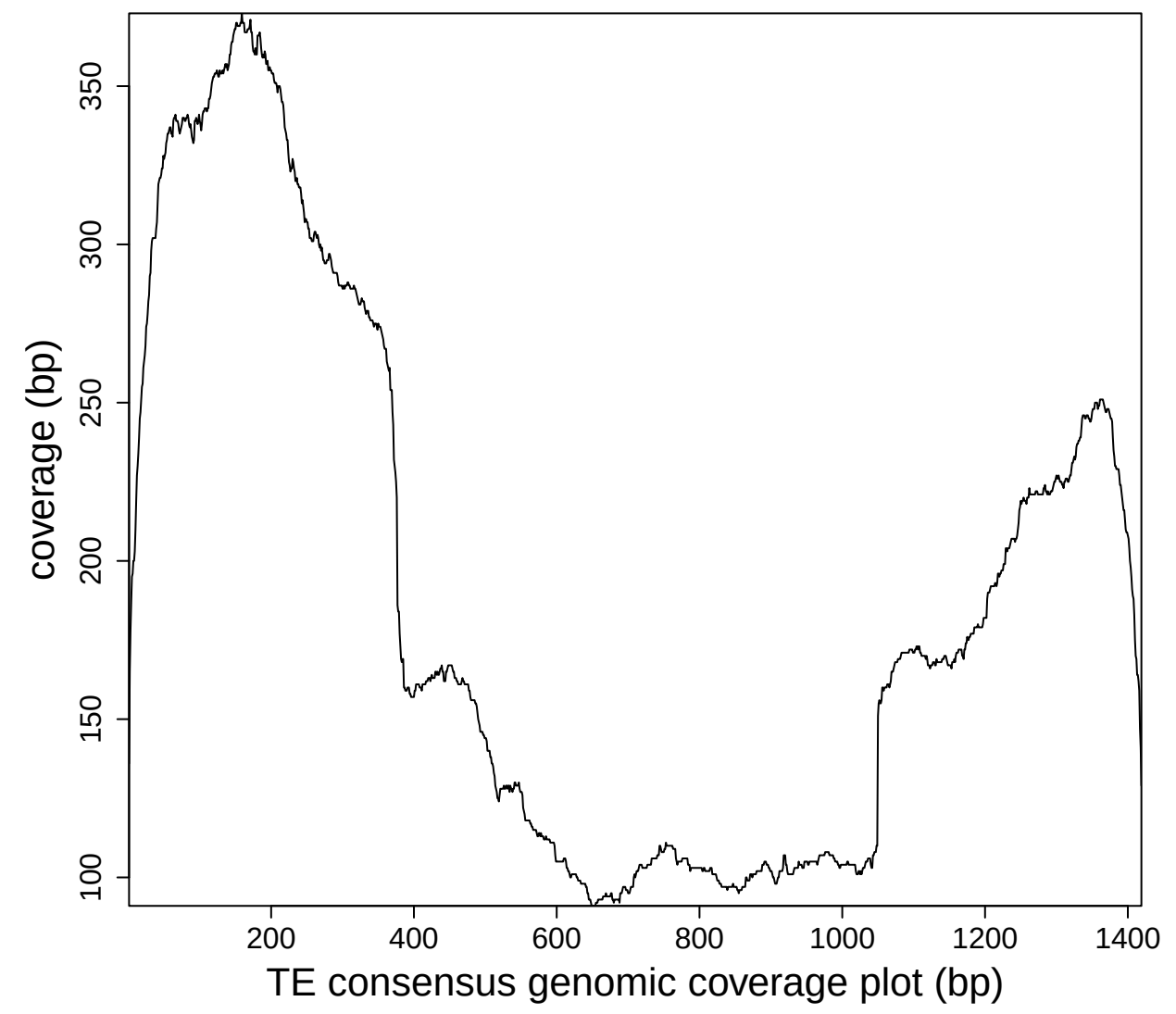
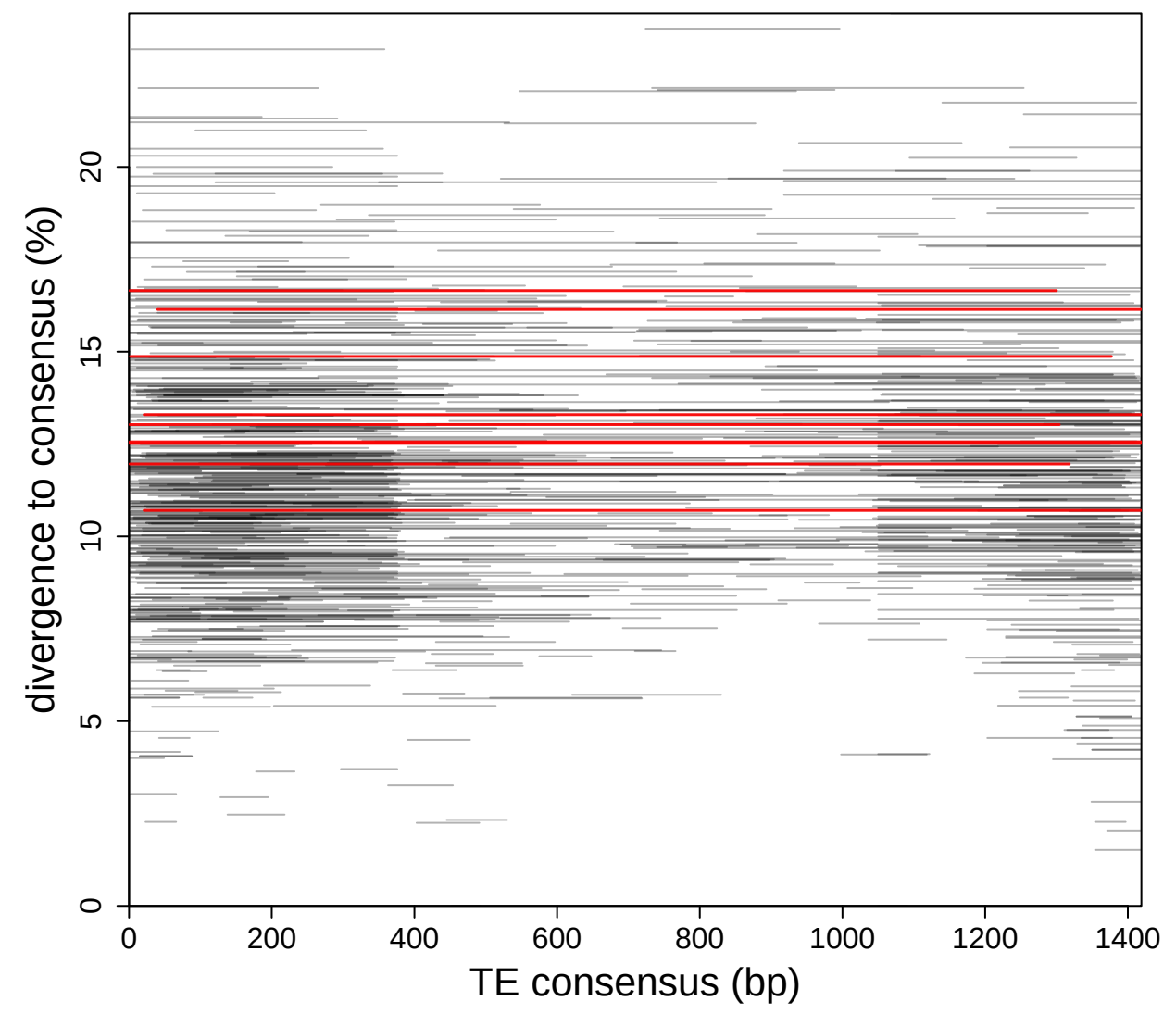
Start crop Point

End crop Point



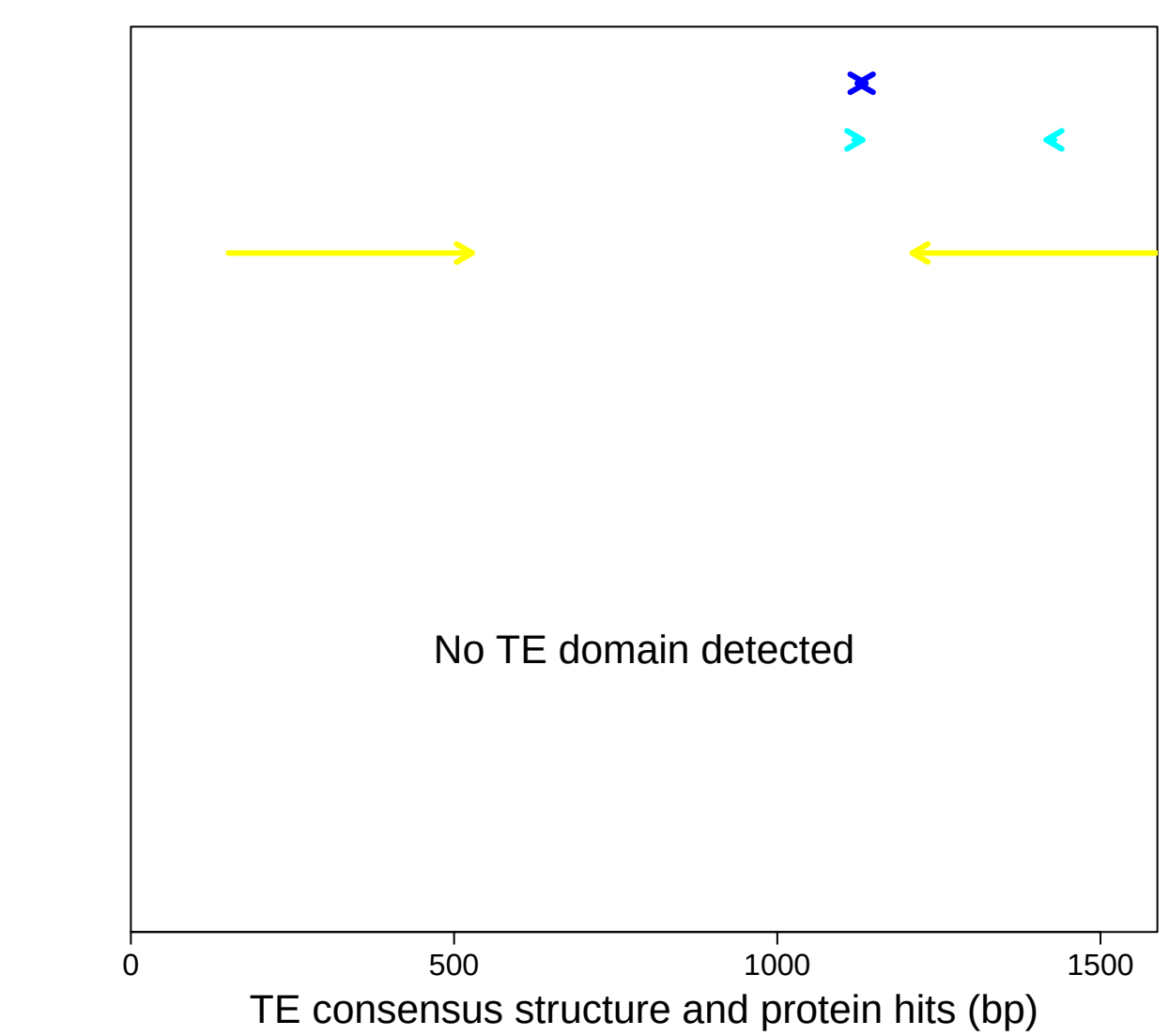
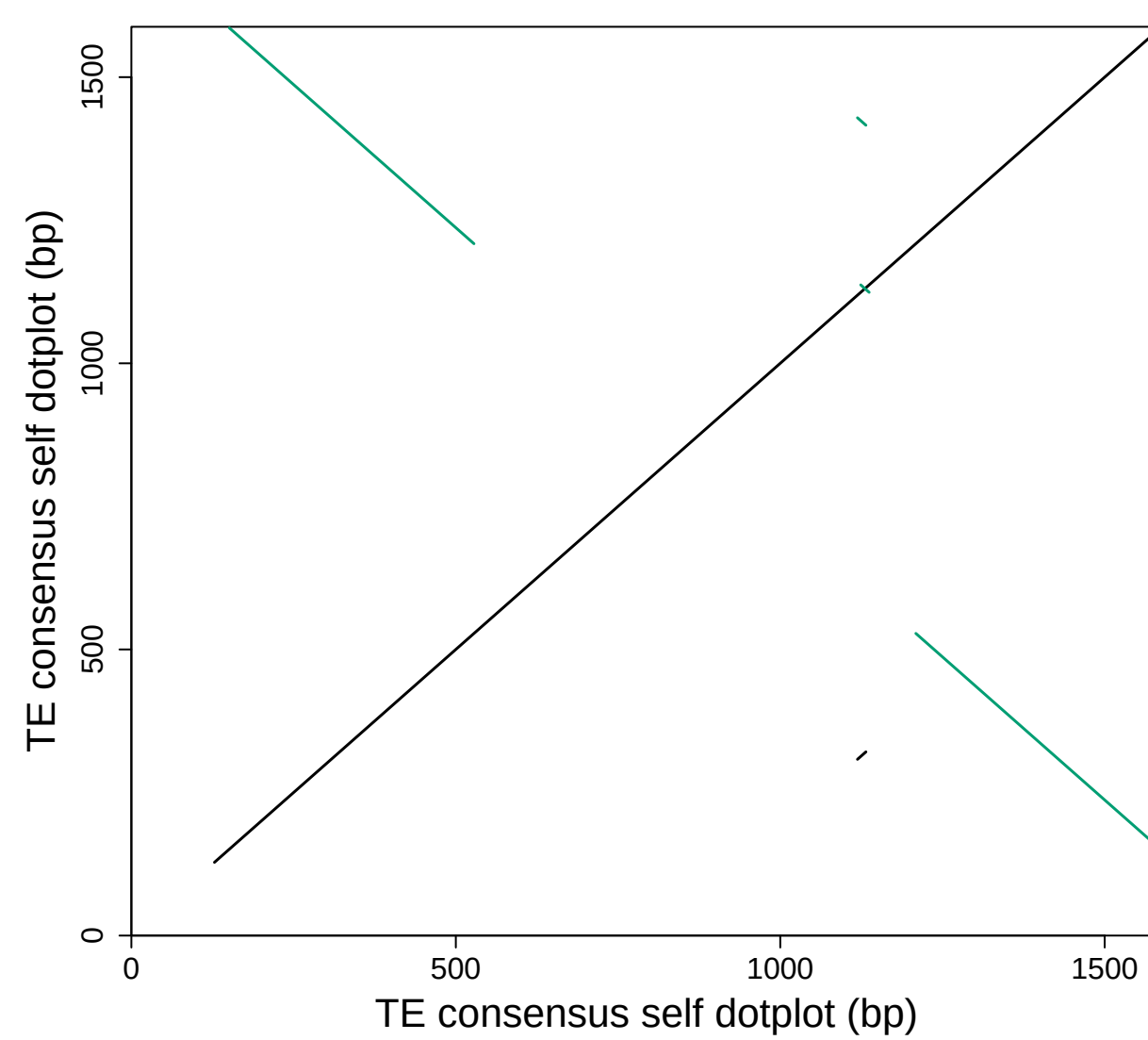
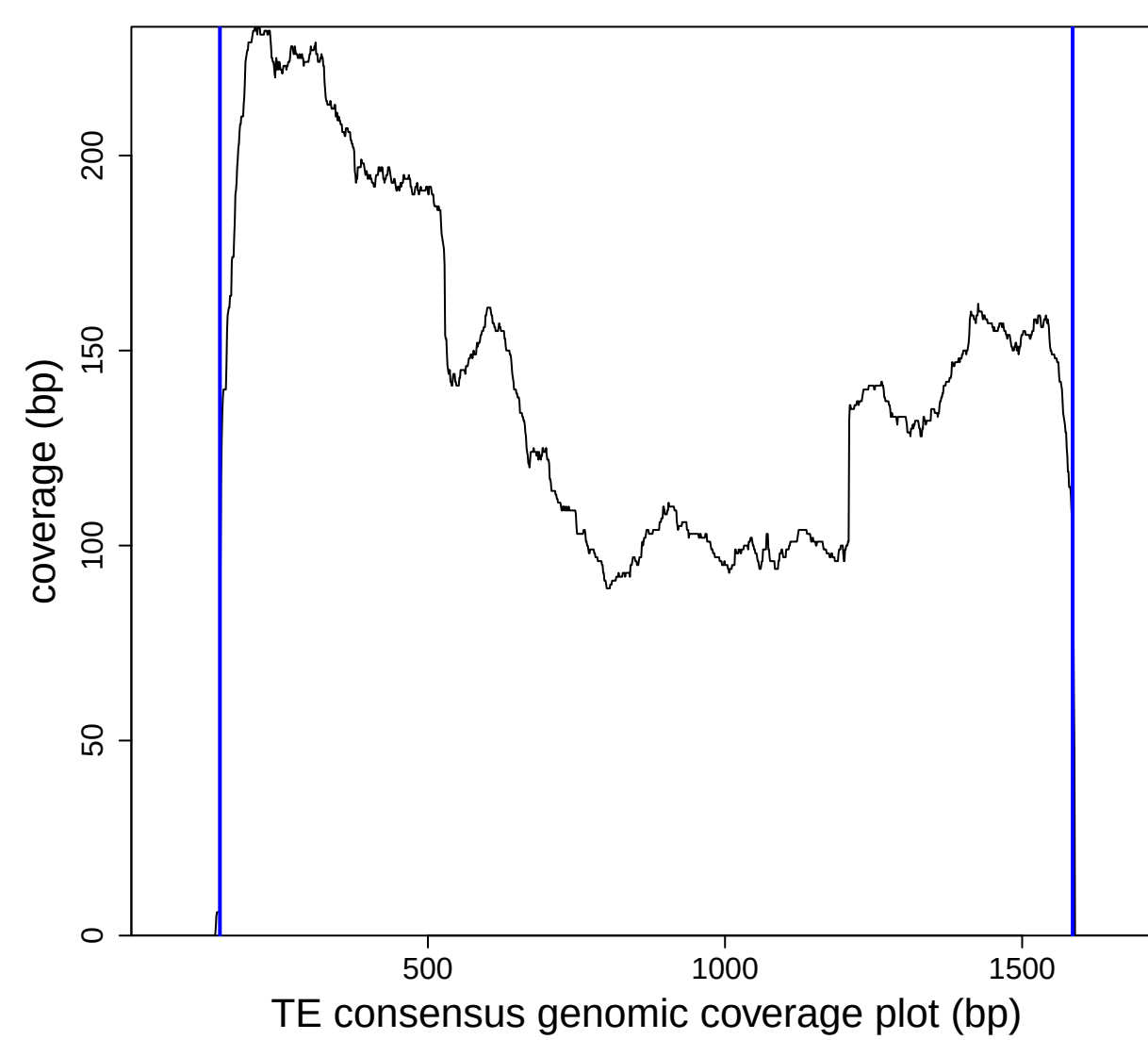
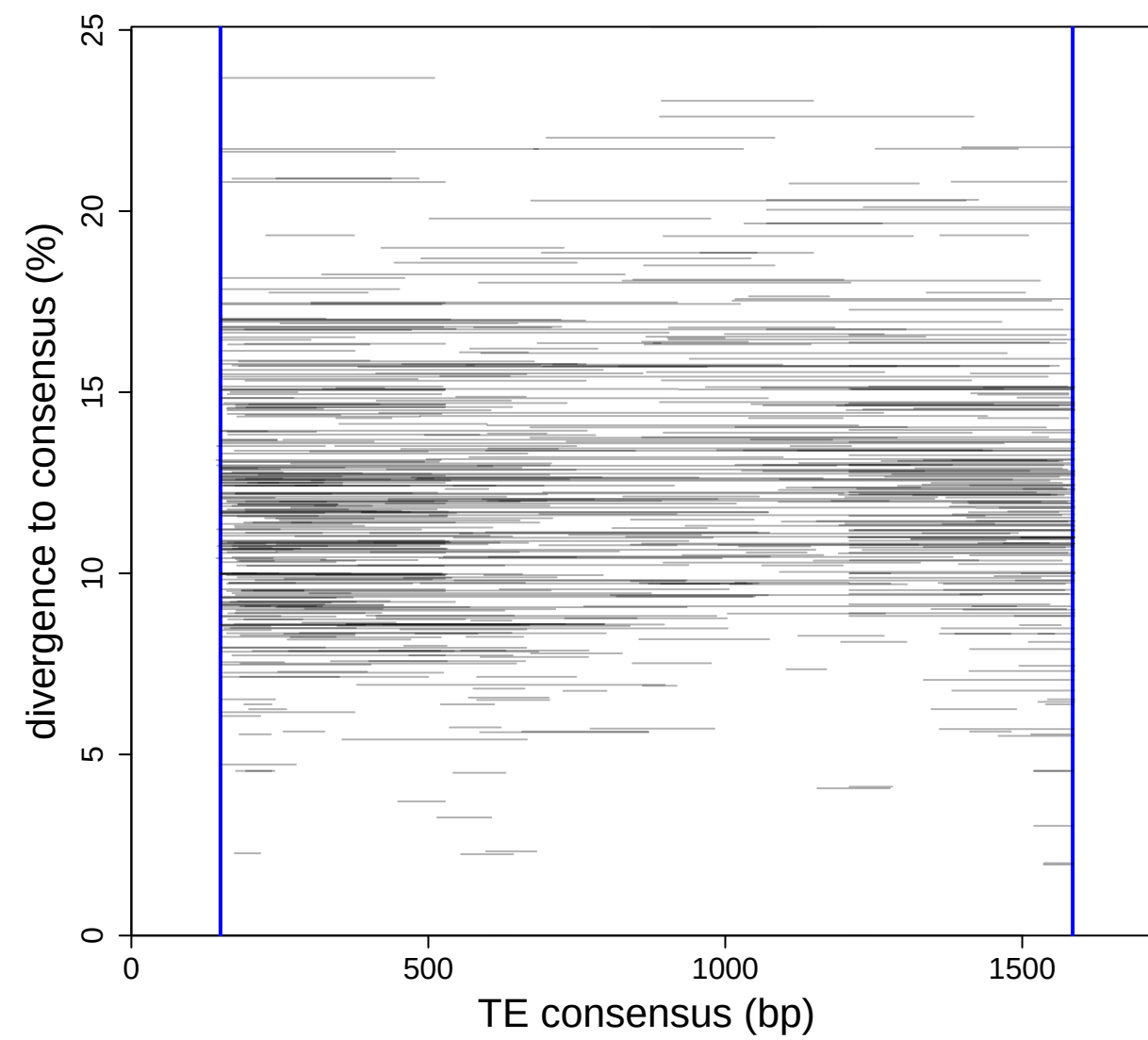
bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.fa_c
size: 1419bp; fragments: 878; full length: 9 (>=1277.1bp)

After TETrimmer 1419 bp

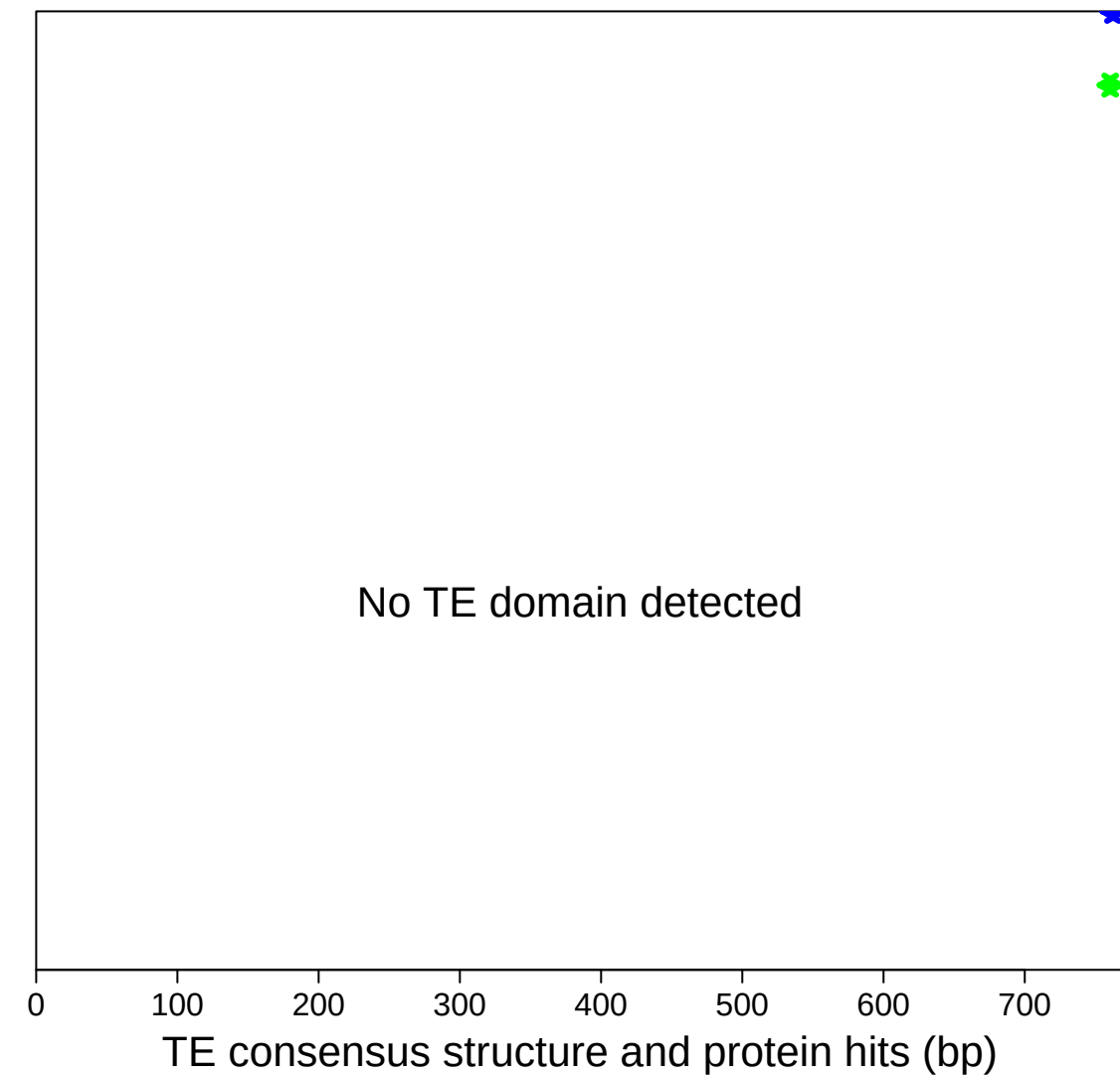
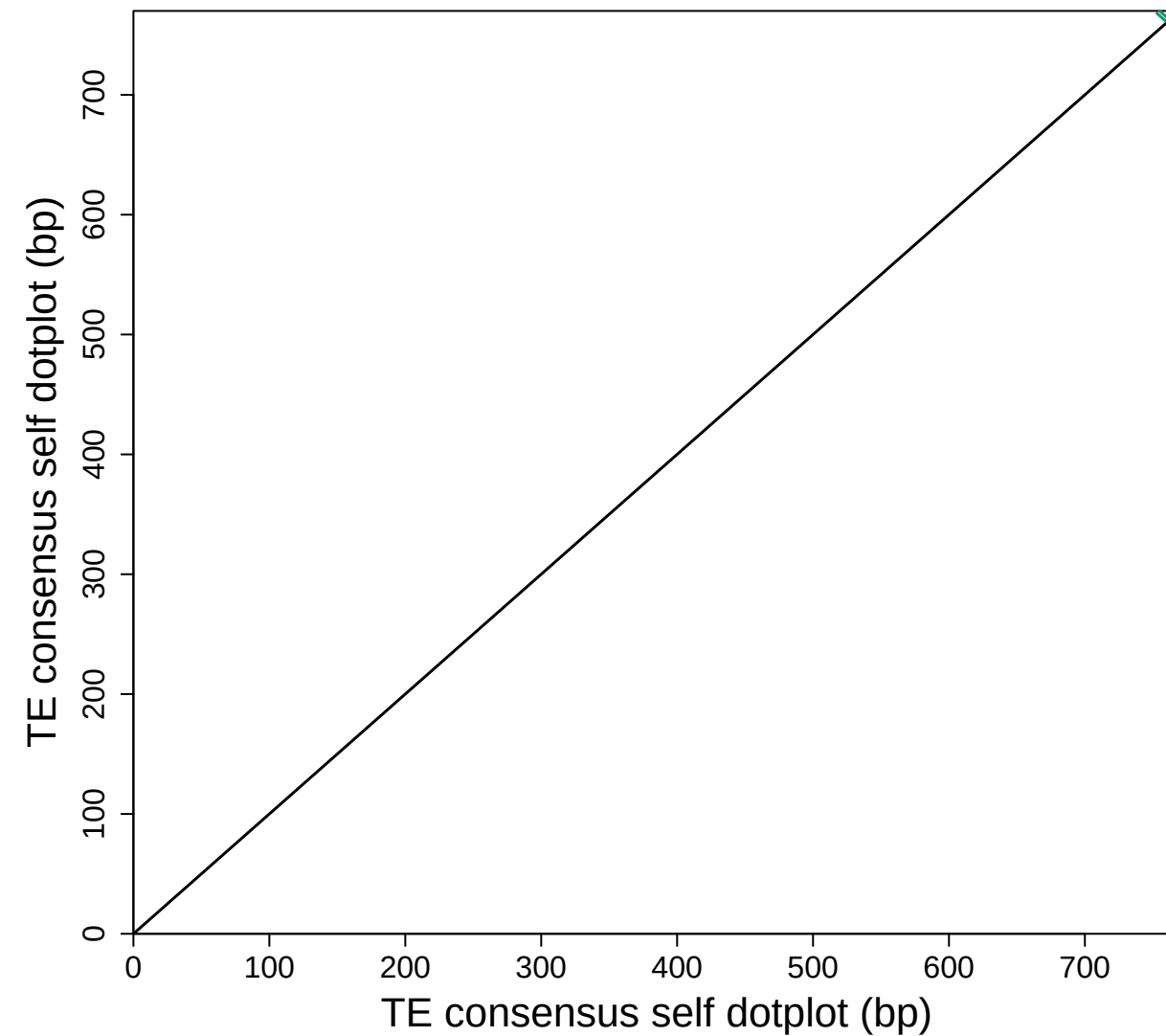
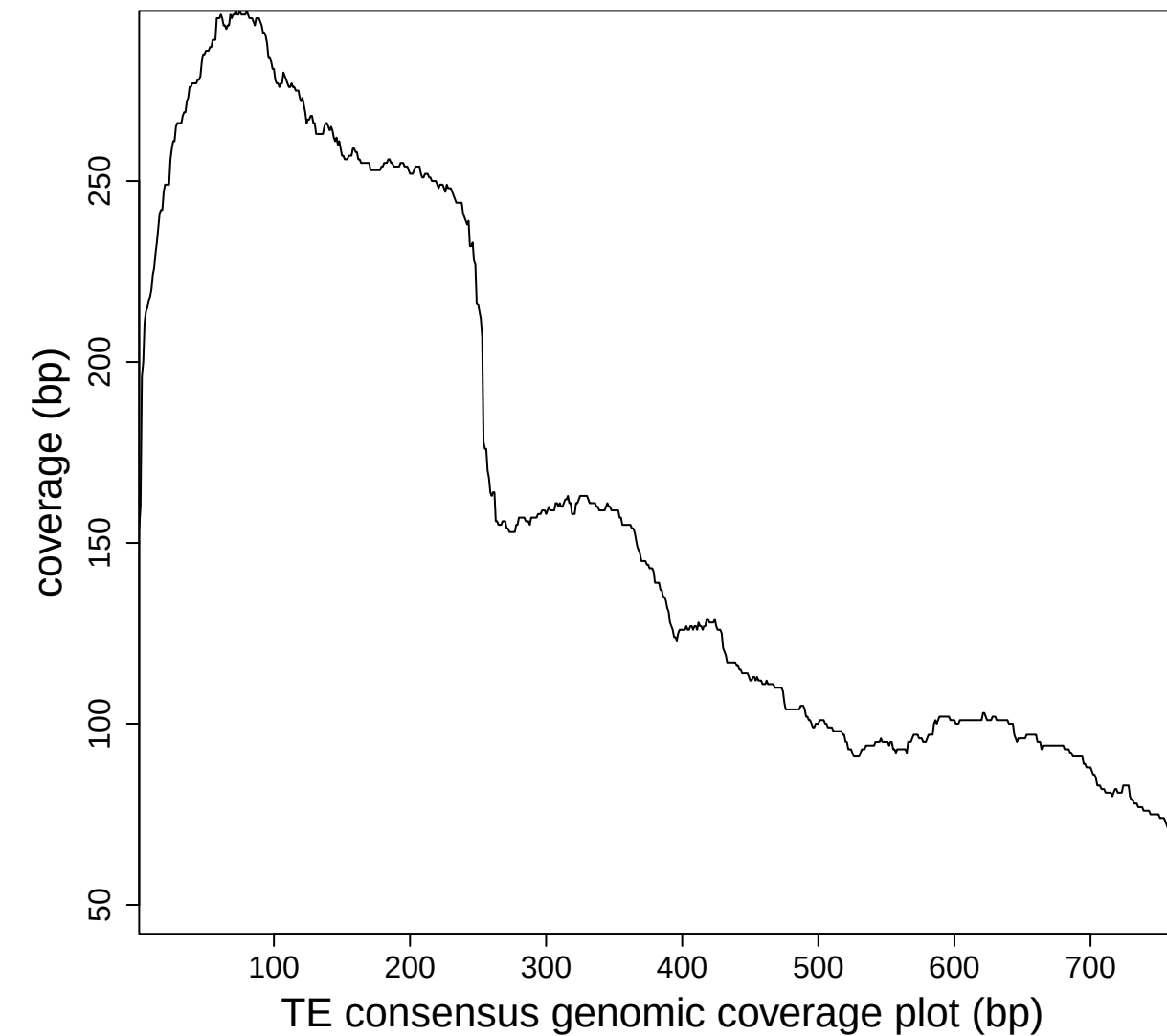
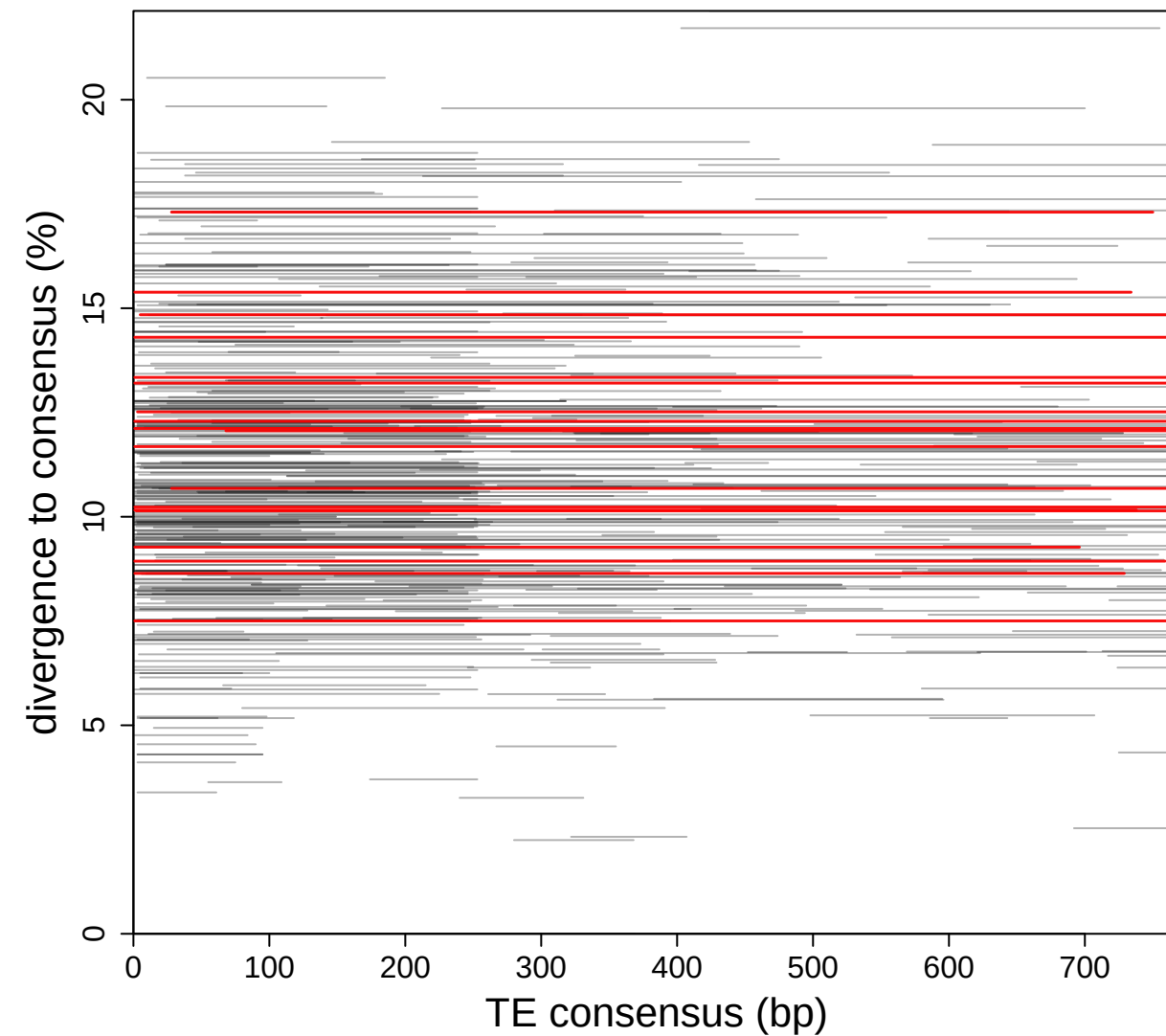


ly_3290.fasta.b.bed uf.bed g_1.bed fm_1.bed_0_0 bcIn.fa aln.f
size: 1735bp; fragments: 613; full length: 0 (>=1561.5bp)

After TEtrimmer Extended plot Blue lines are boundaries

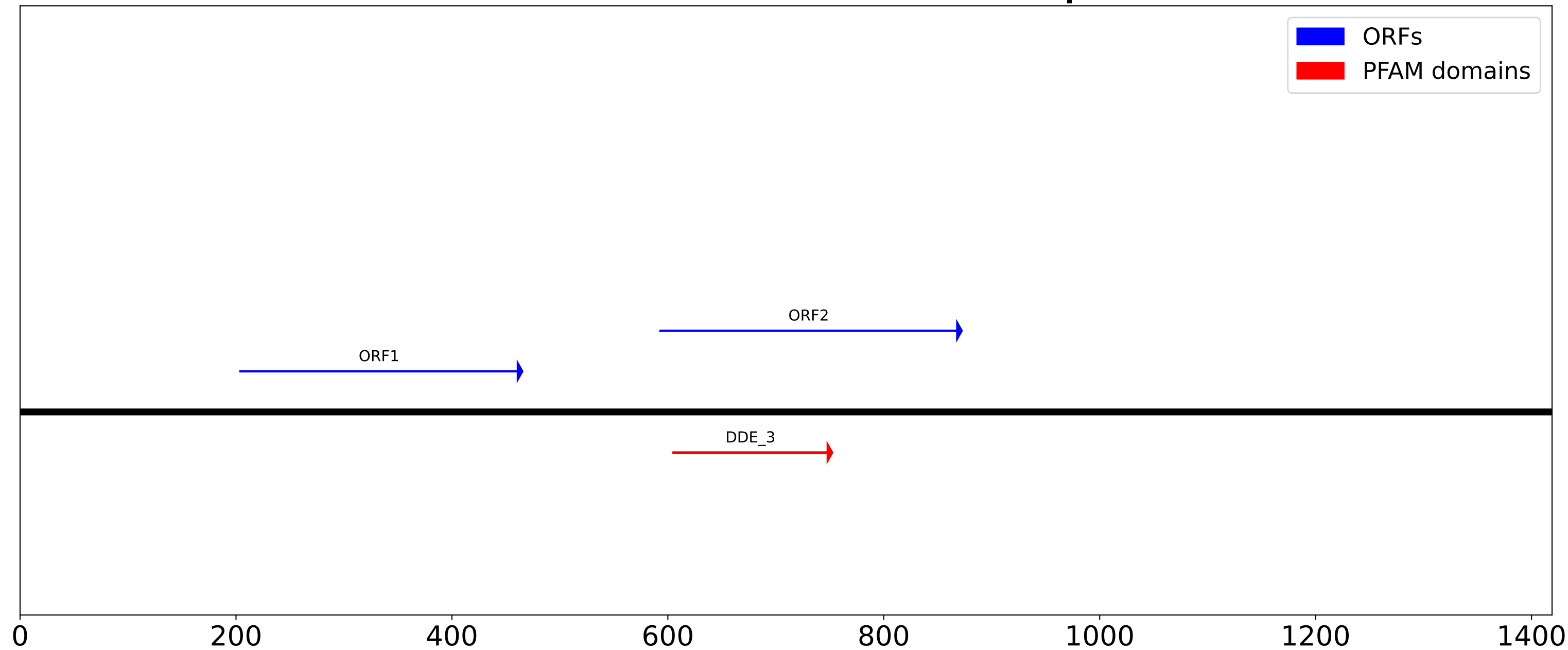


TE: rnd_5_family_3290
size: 770bp; fragments: 452; full length: 19 (≥ 693 bp)

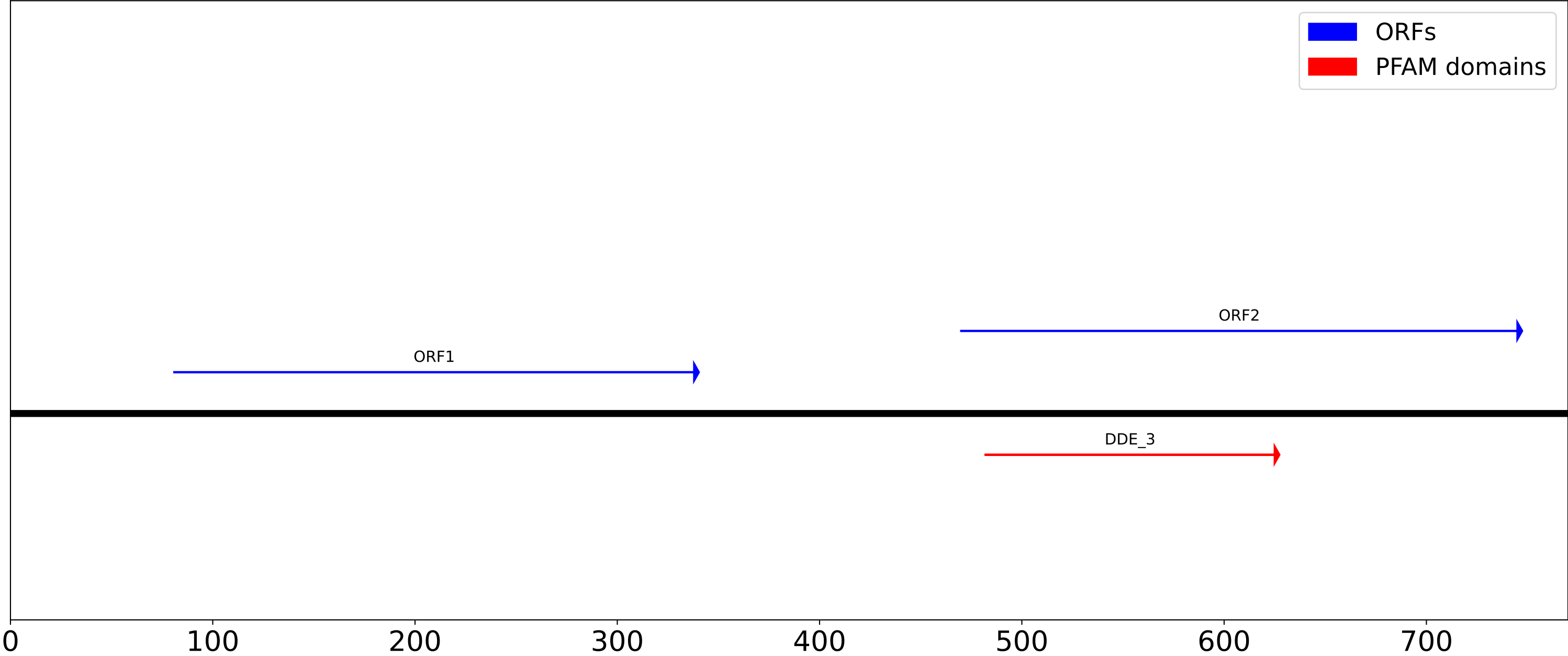


Before TEtrimmer 770 bp

After TEtrimmer ORF and PFAM domain plot



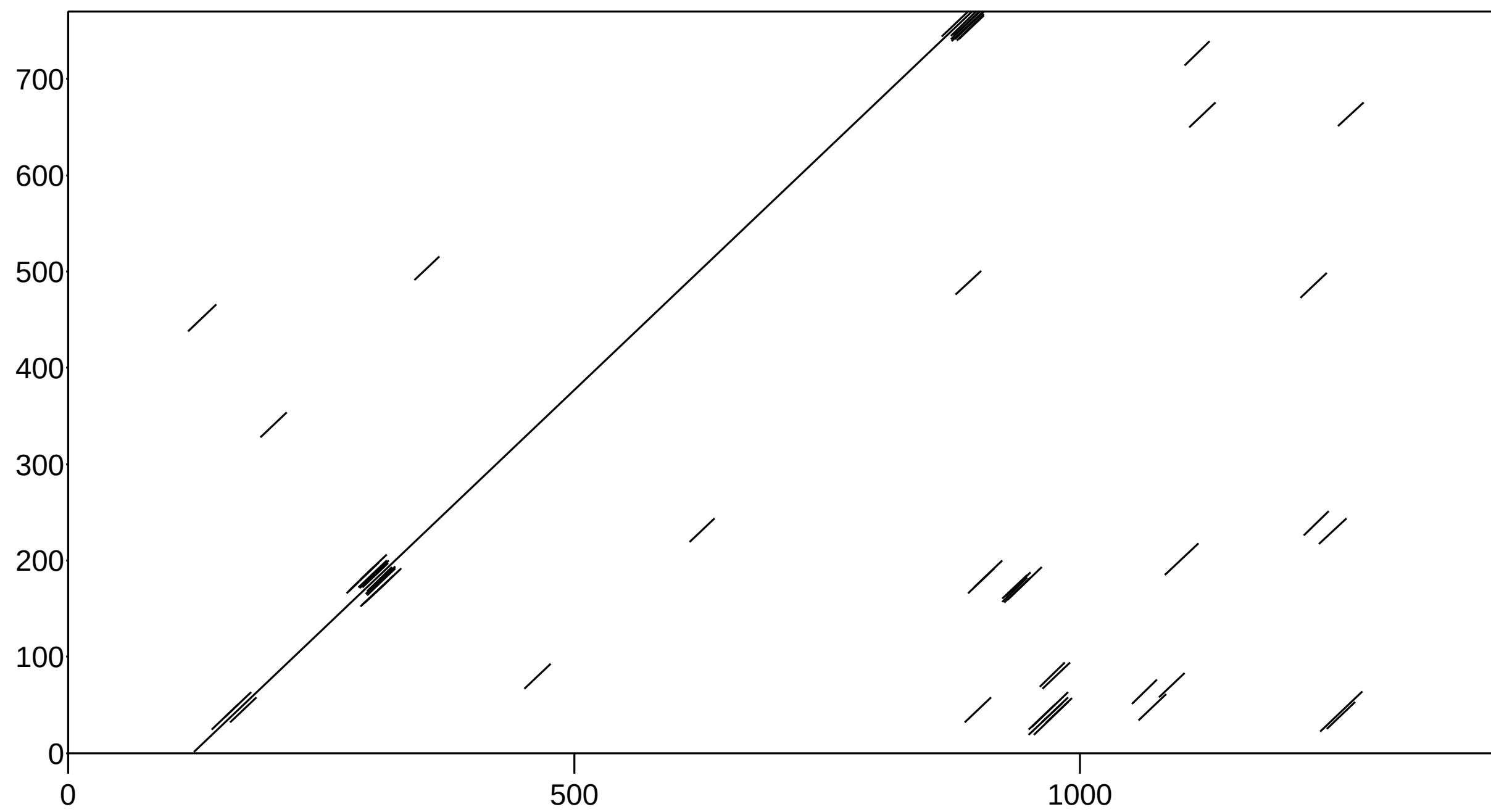
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)