

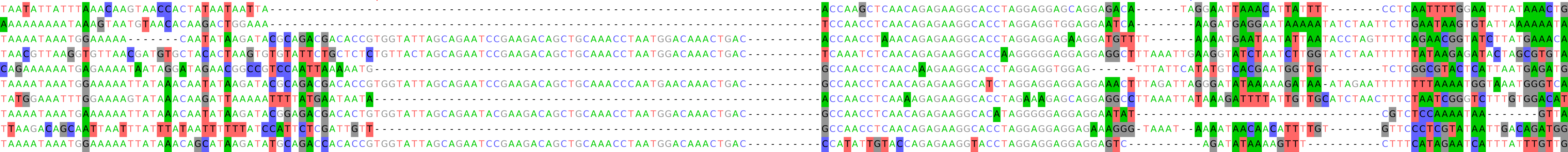
Start Crop Point



End Crop Point

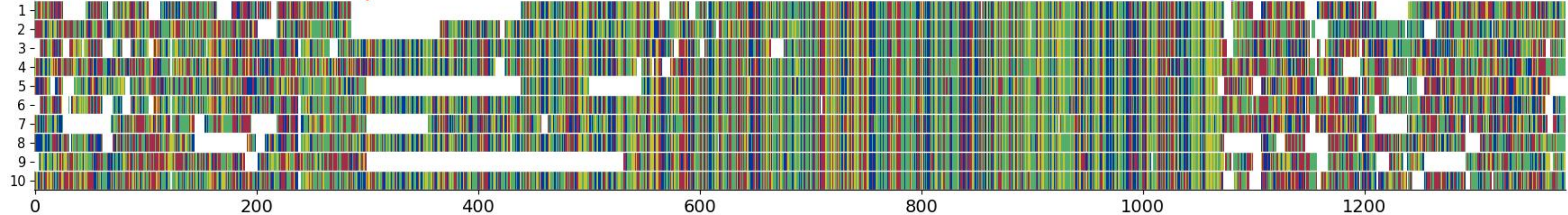


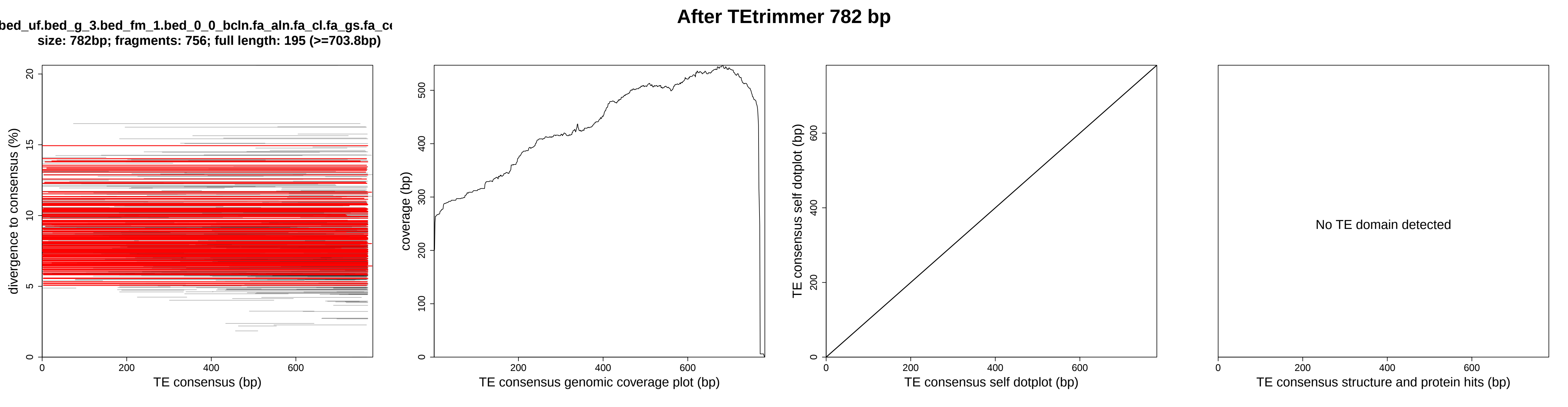
MSA length = 782



Start crop Point

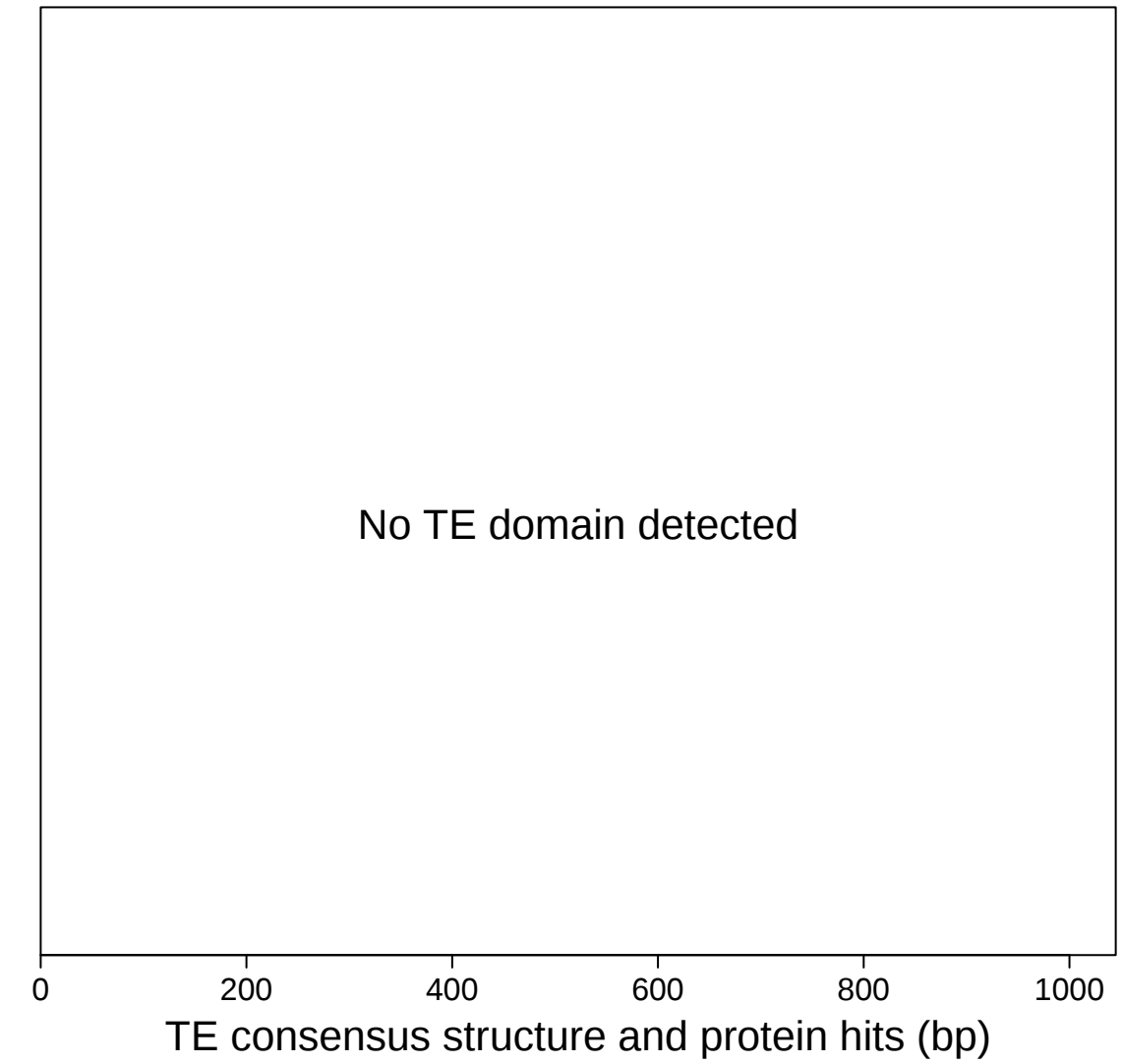
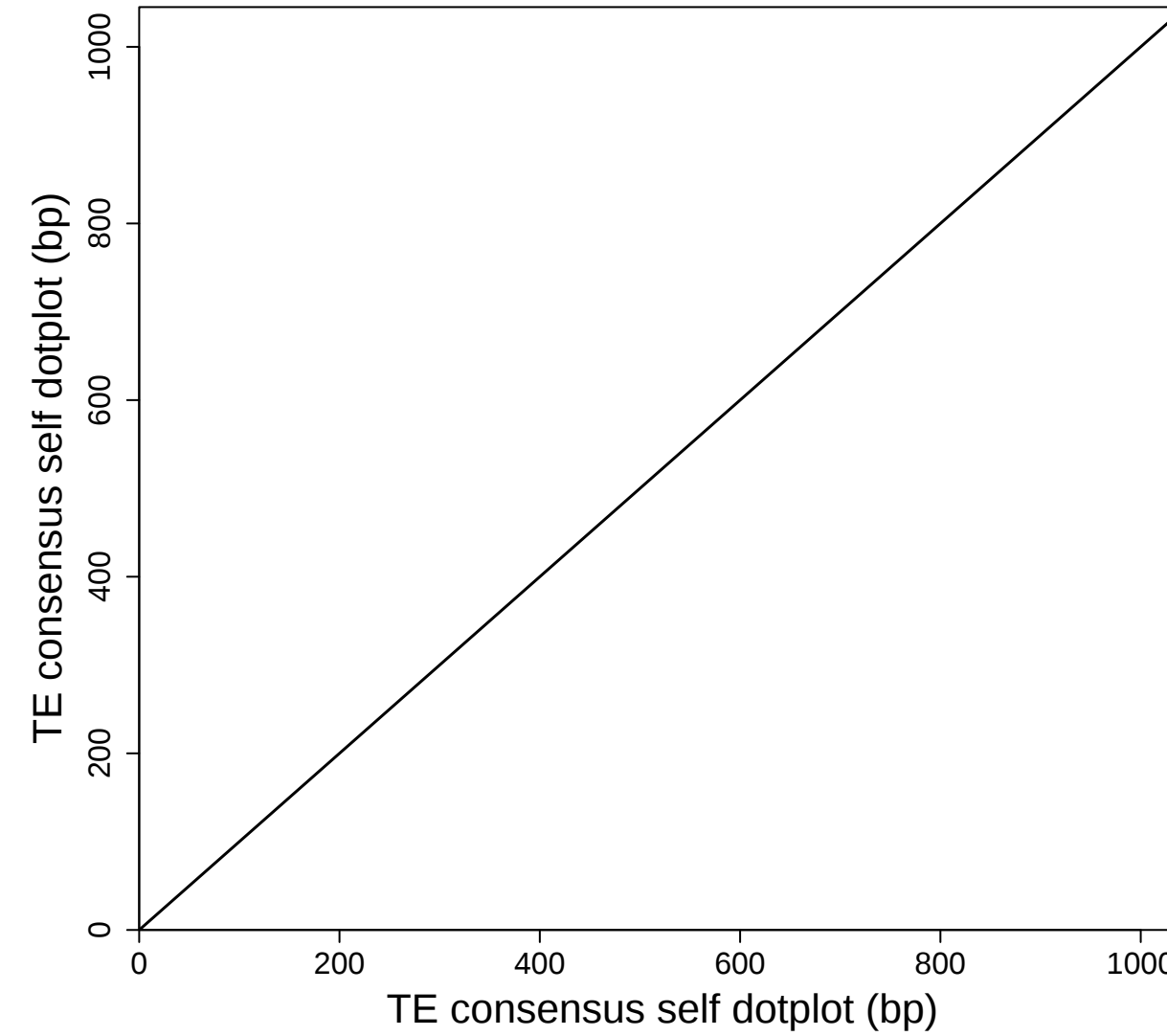
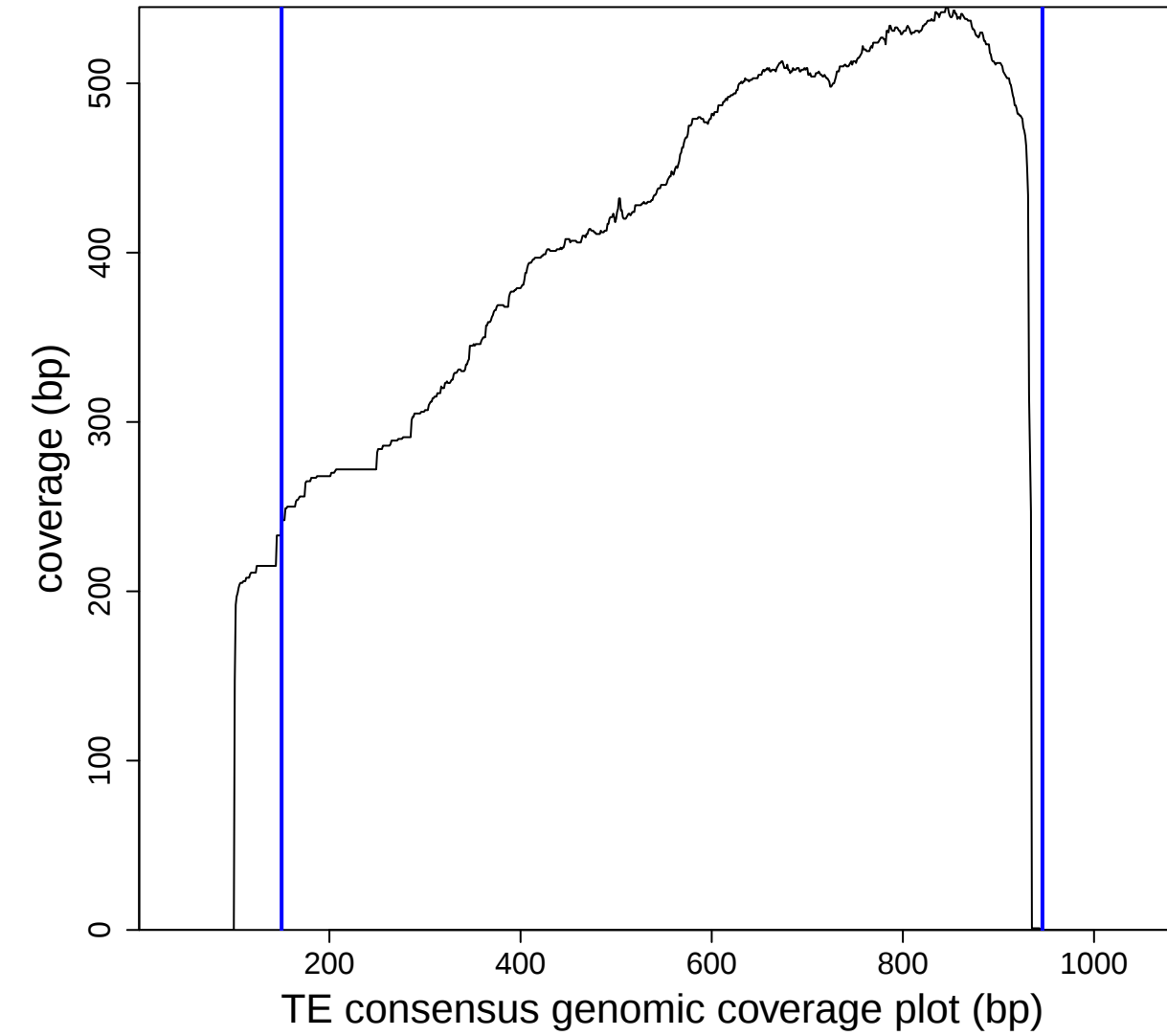
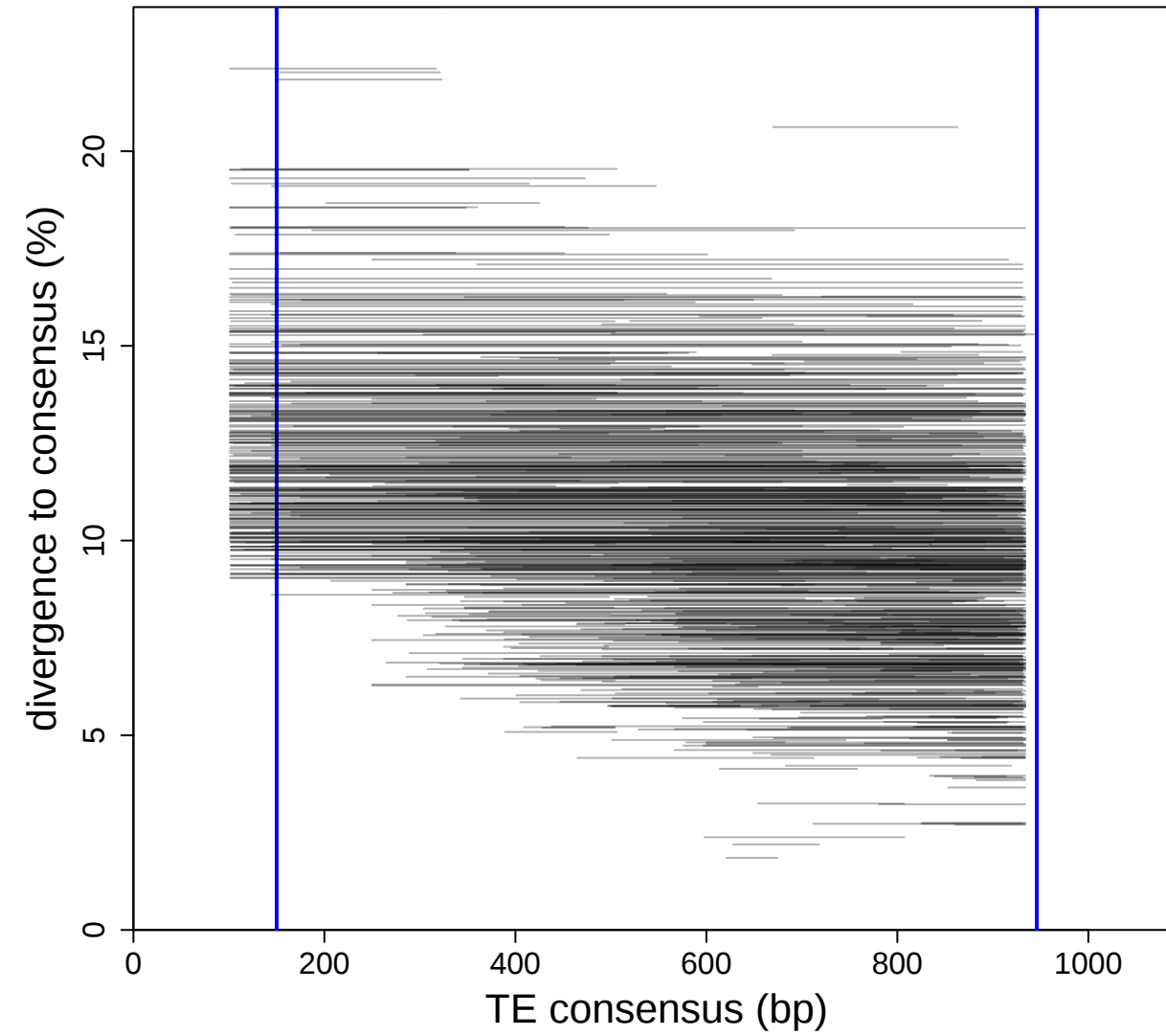
End crop Point



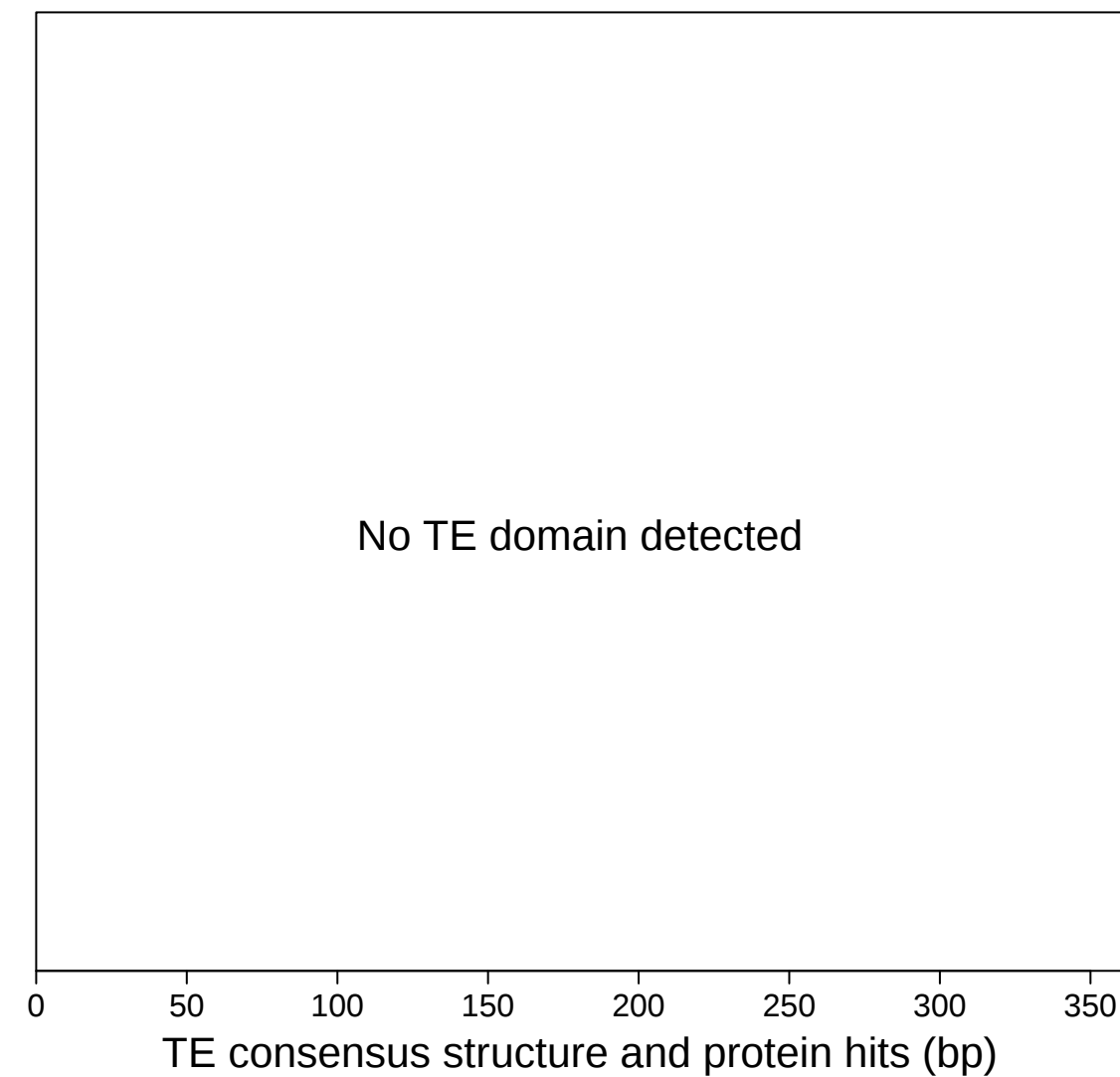
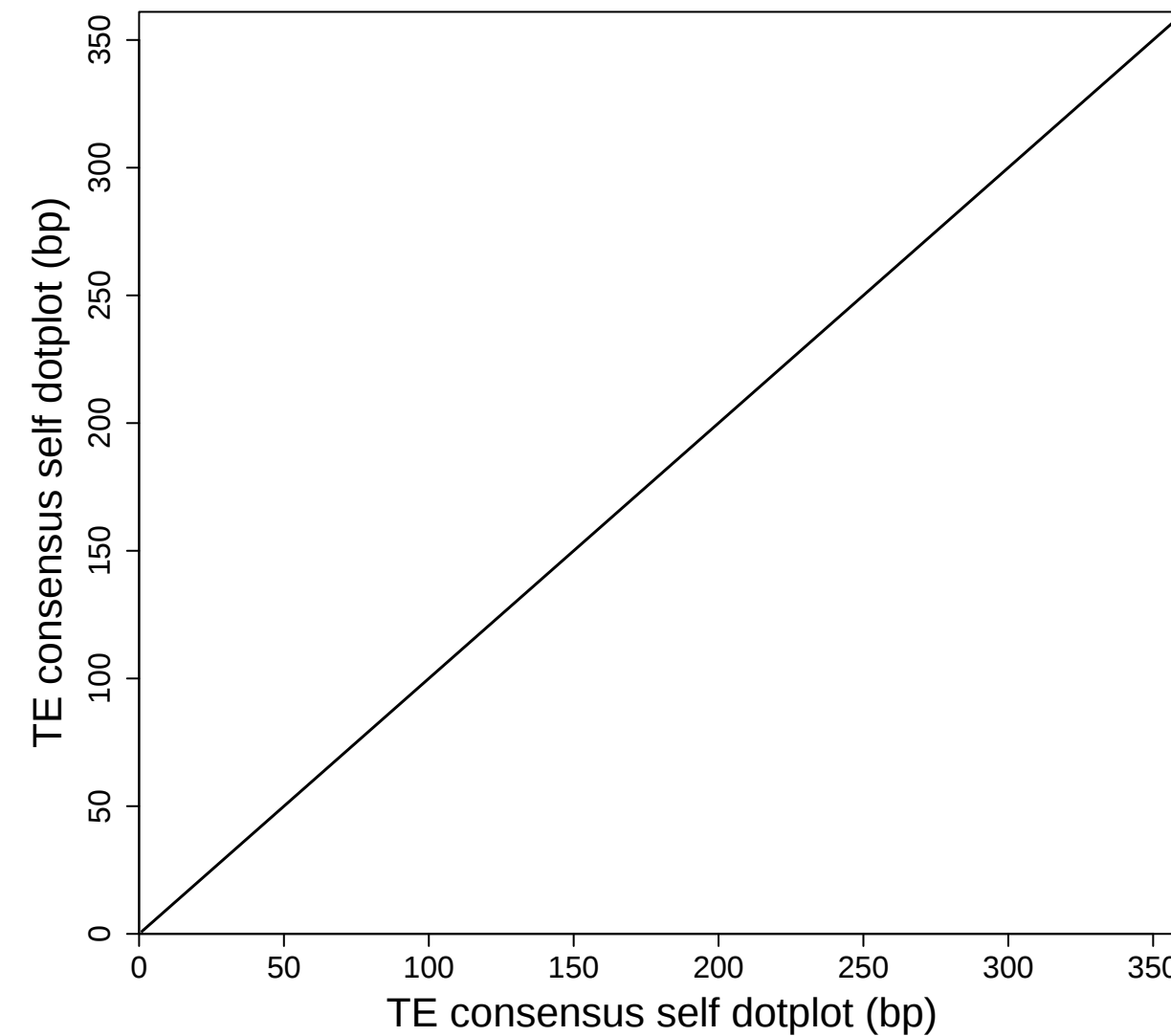
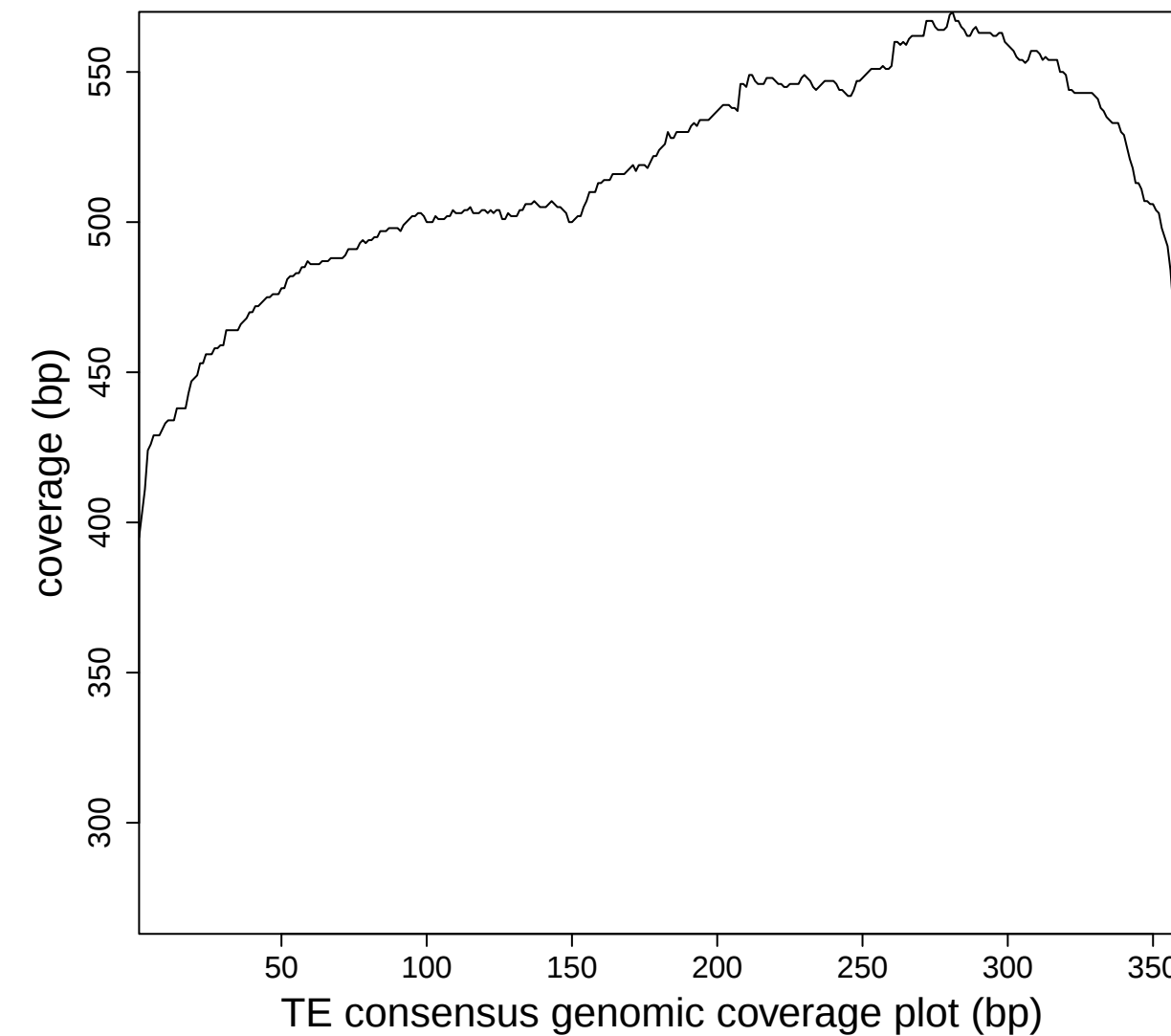
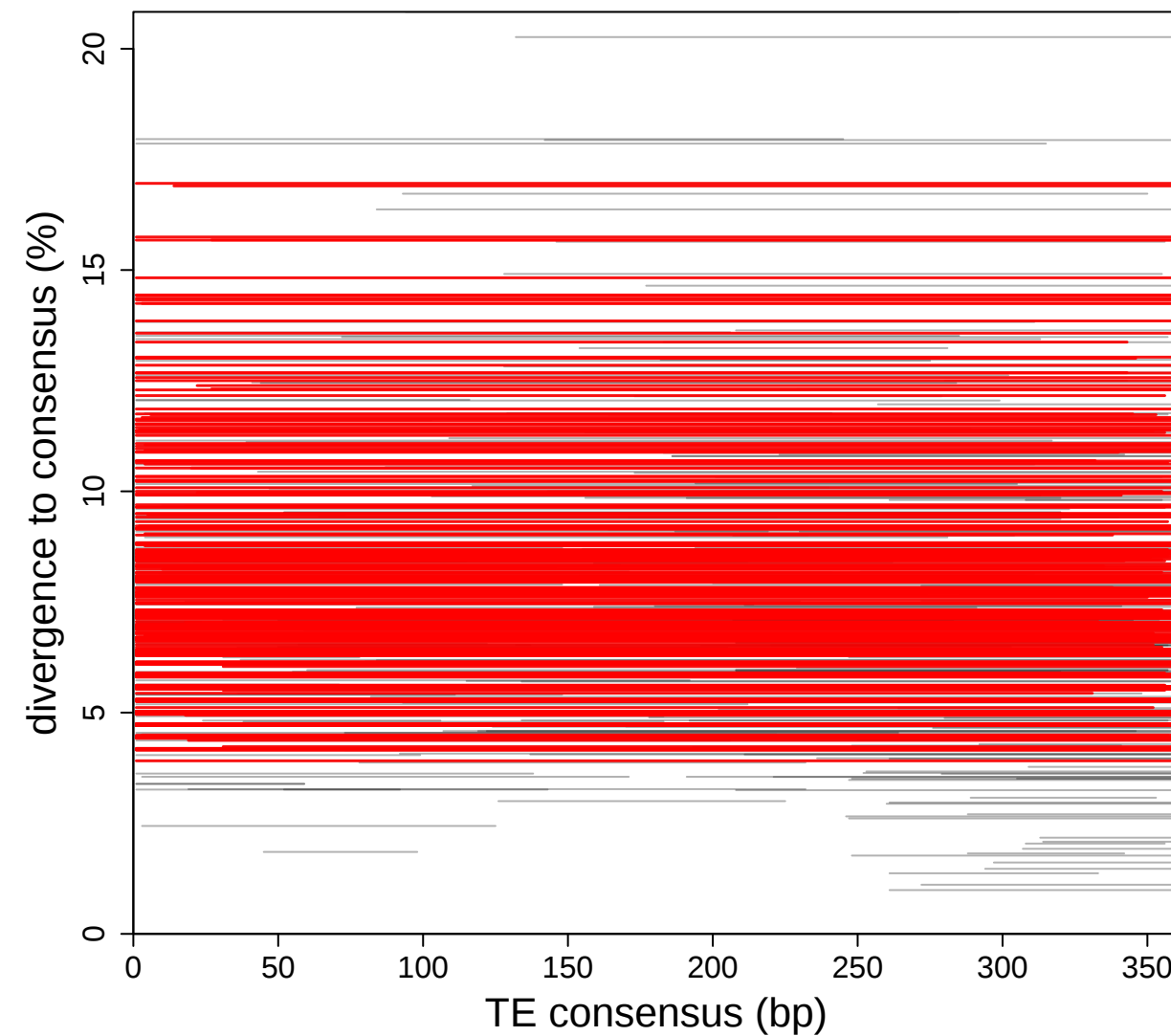


6.fasta.b.bed_uf.bed_g_3.bed_fm_1.bed_0_0_bc\nln.fa_aln.fa_cl.f
size: 1096bp; fragments: 726; full length: 0 (>=986.4bp)

After TEtrimmer Extended plot Blue lines are boundaries

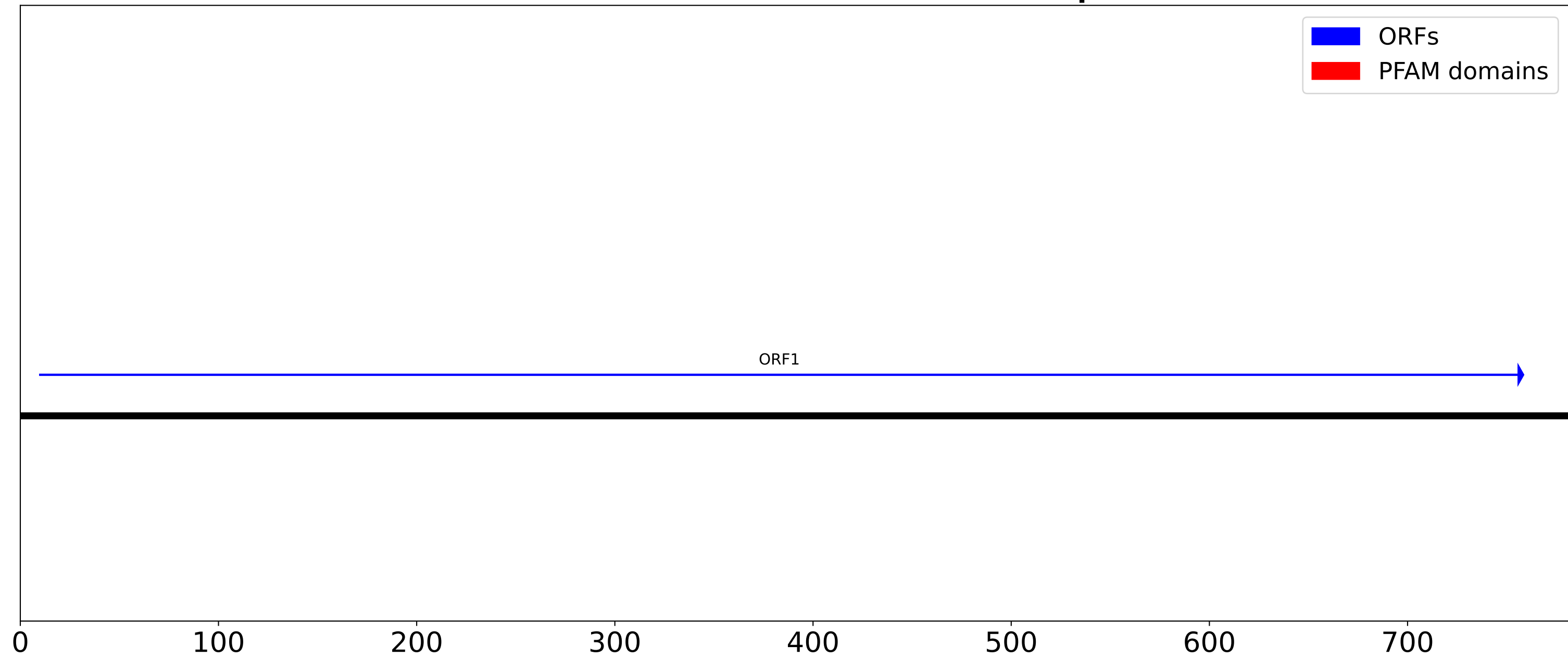


TE: rnd_5_family_1796
size: 361bp; fragments: 676; full length: 353 (≥ 324.9 bp)

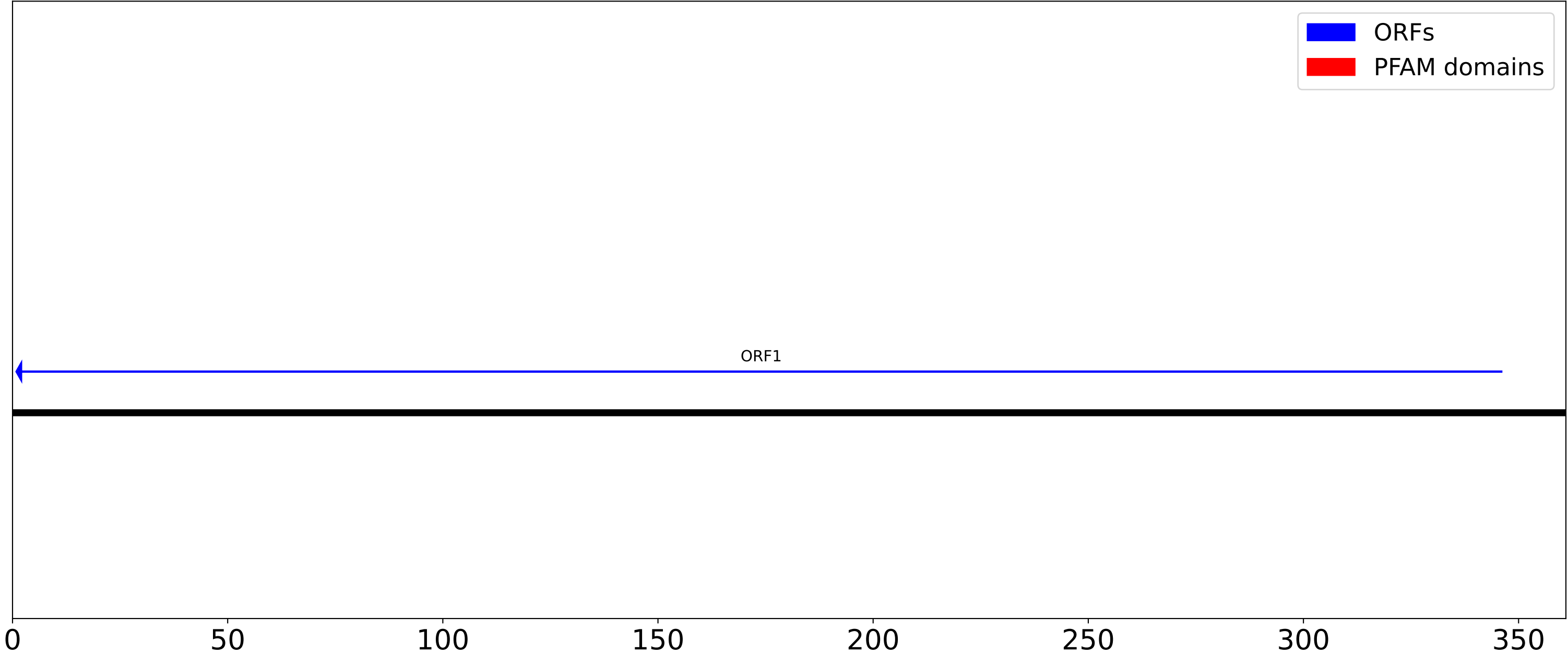


Before TEtrimmer 361 bp

After TEtrimmer ORF and PFAM domain plot



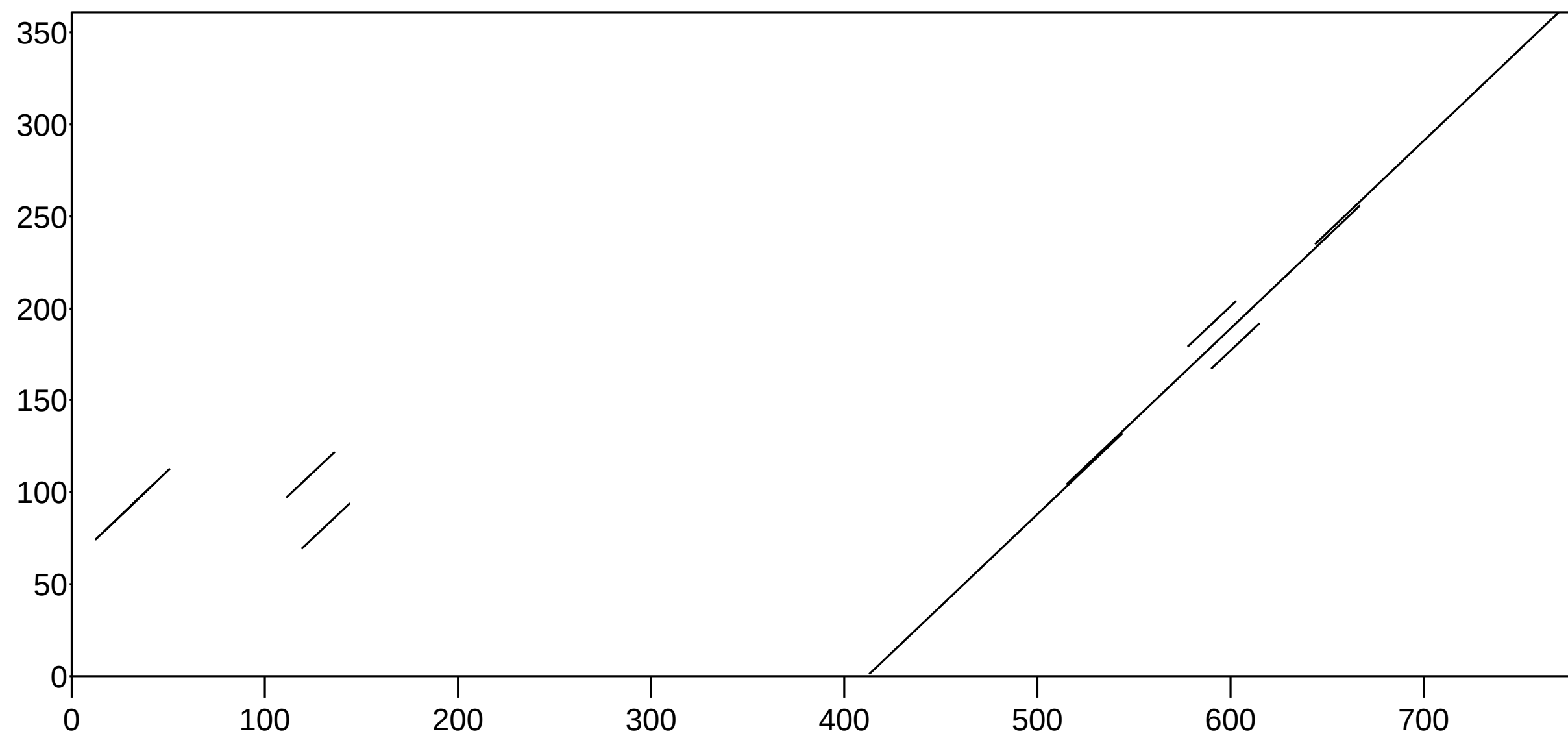
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)