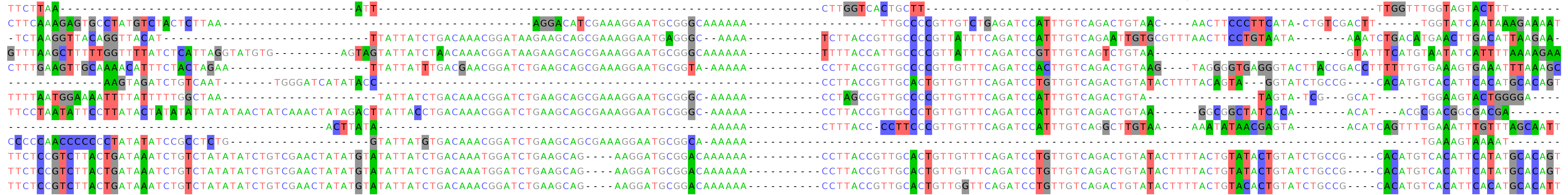
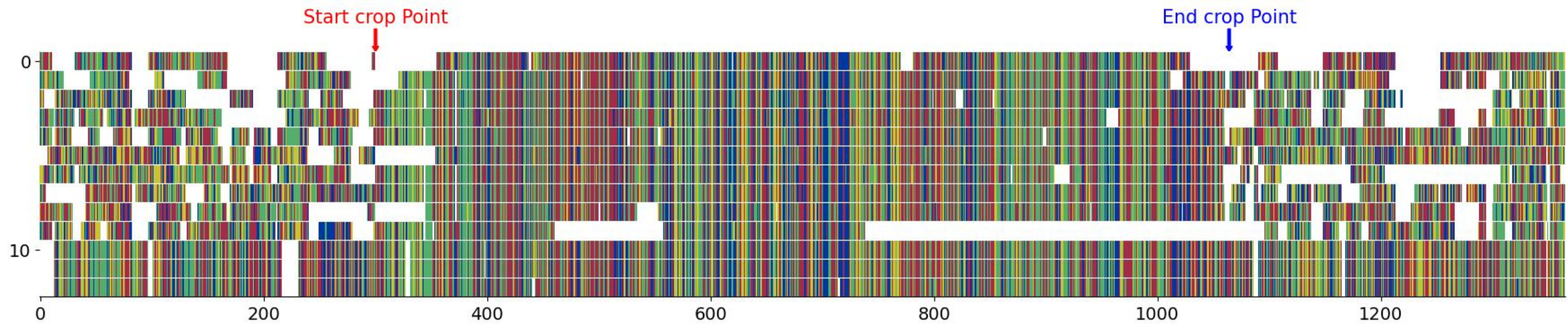


End crop Point

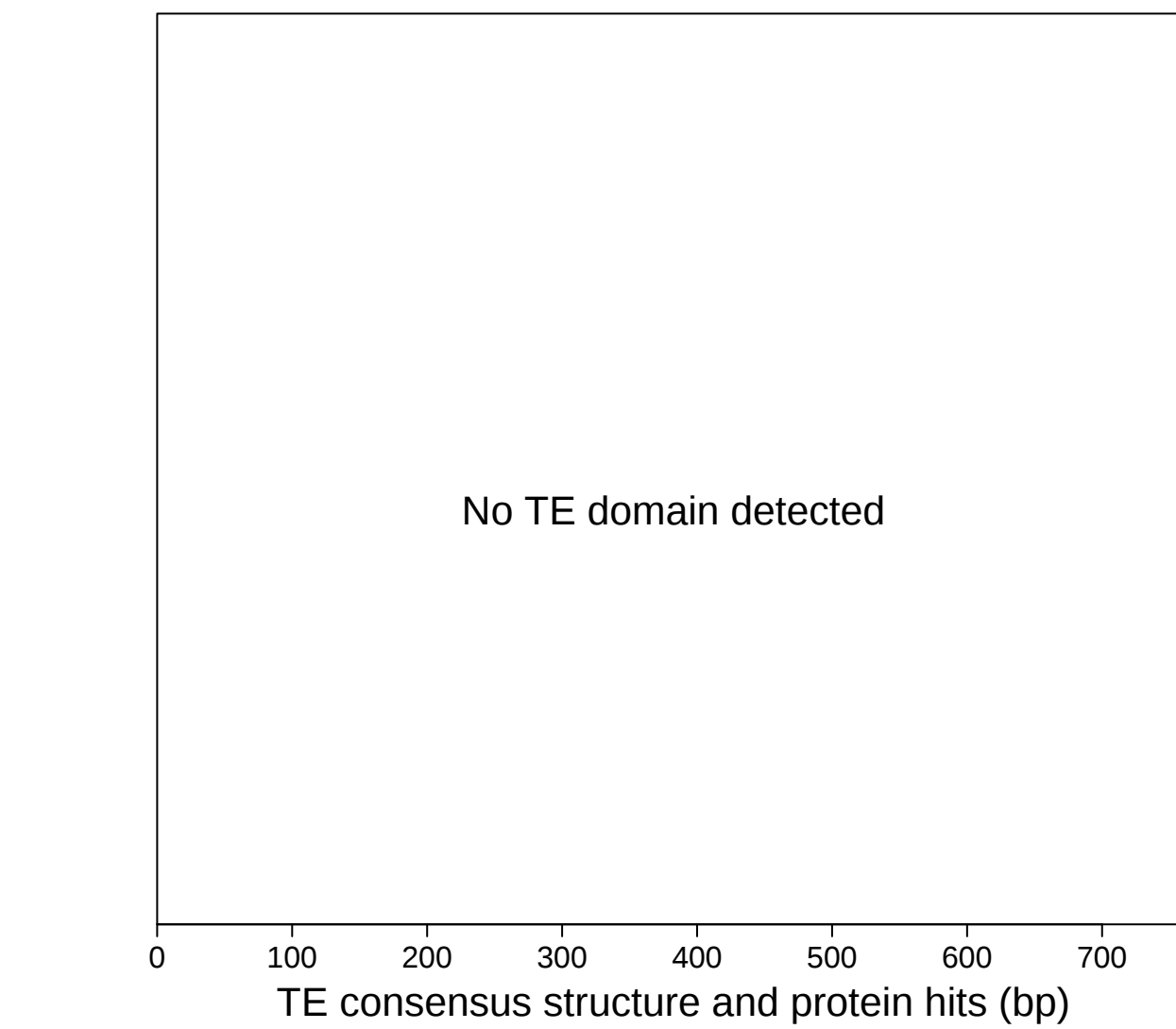
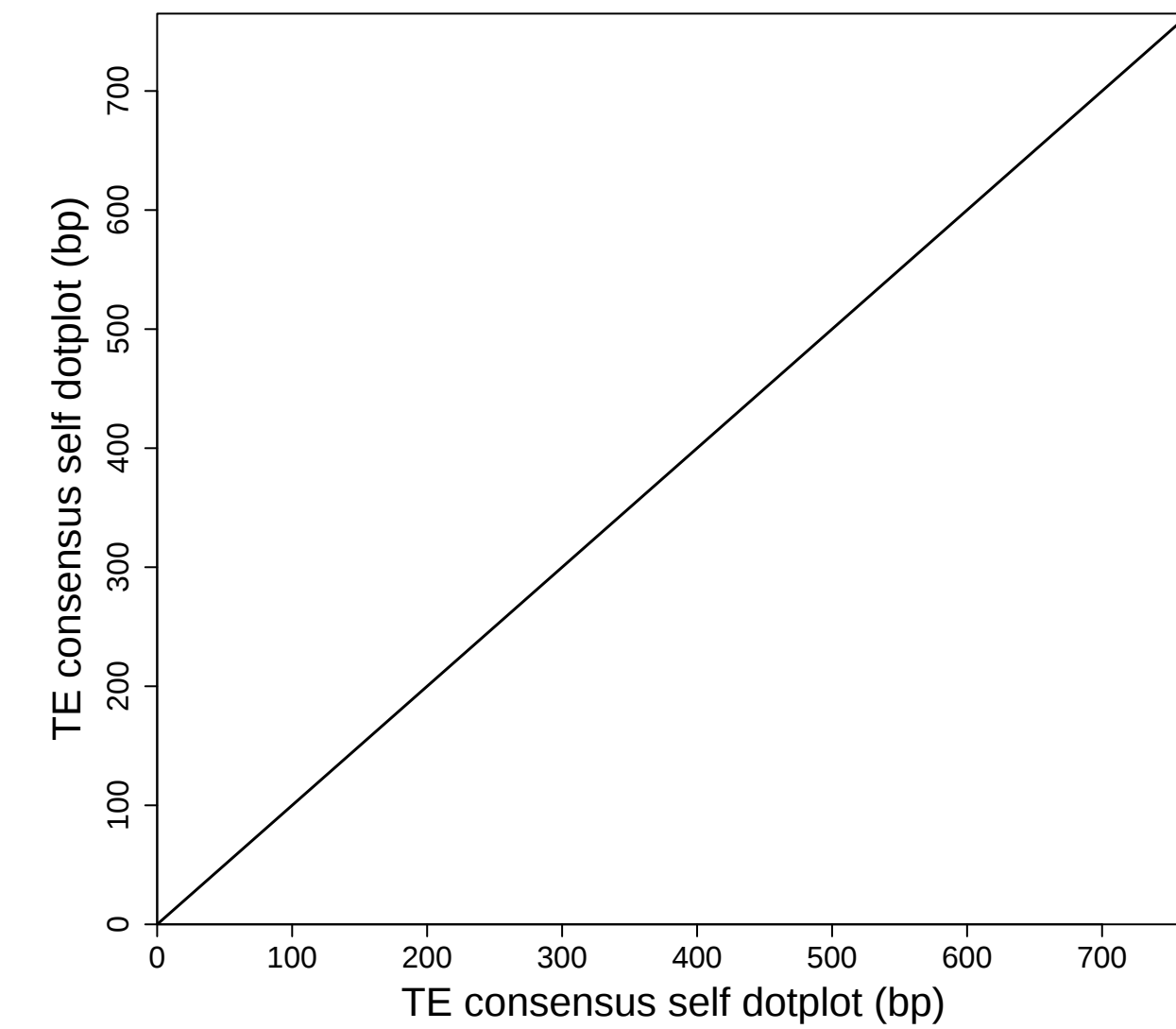
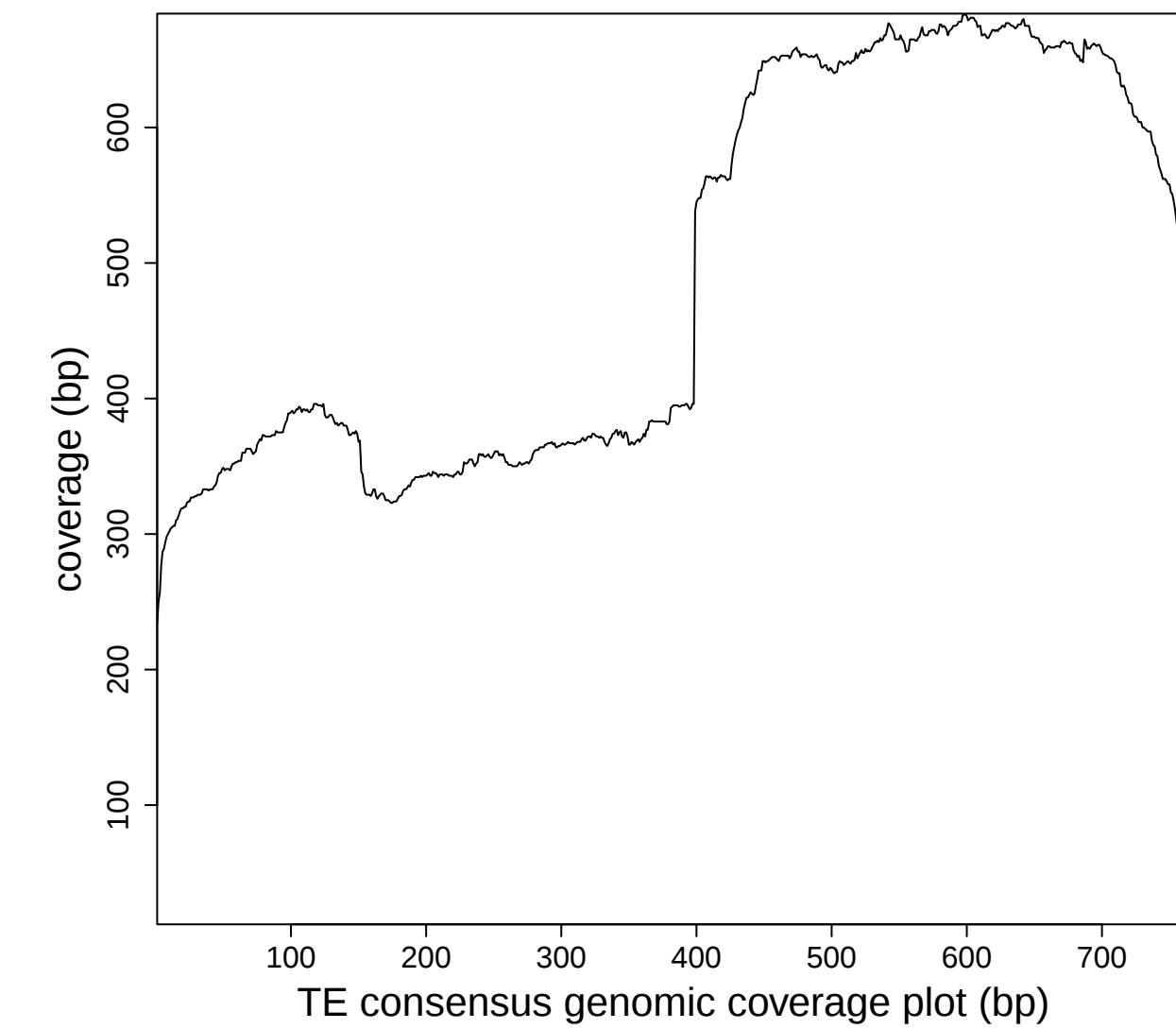
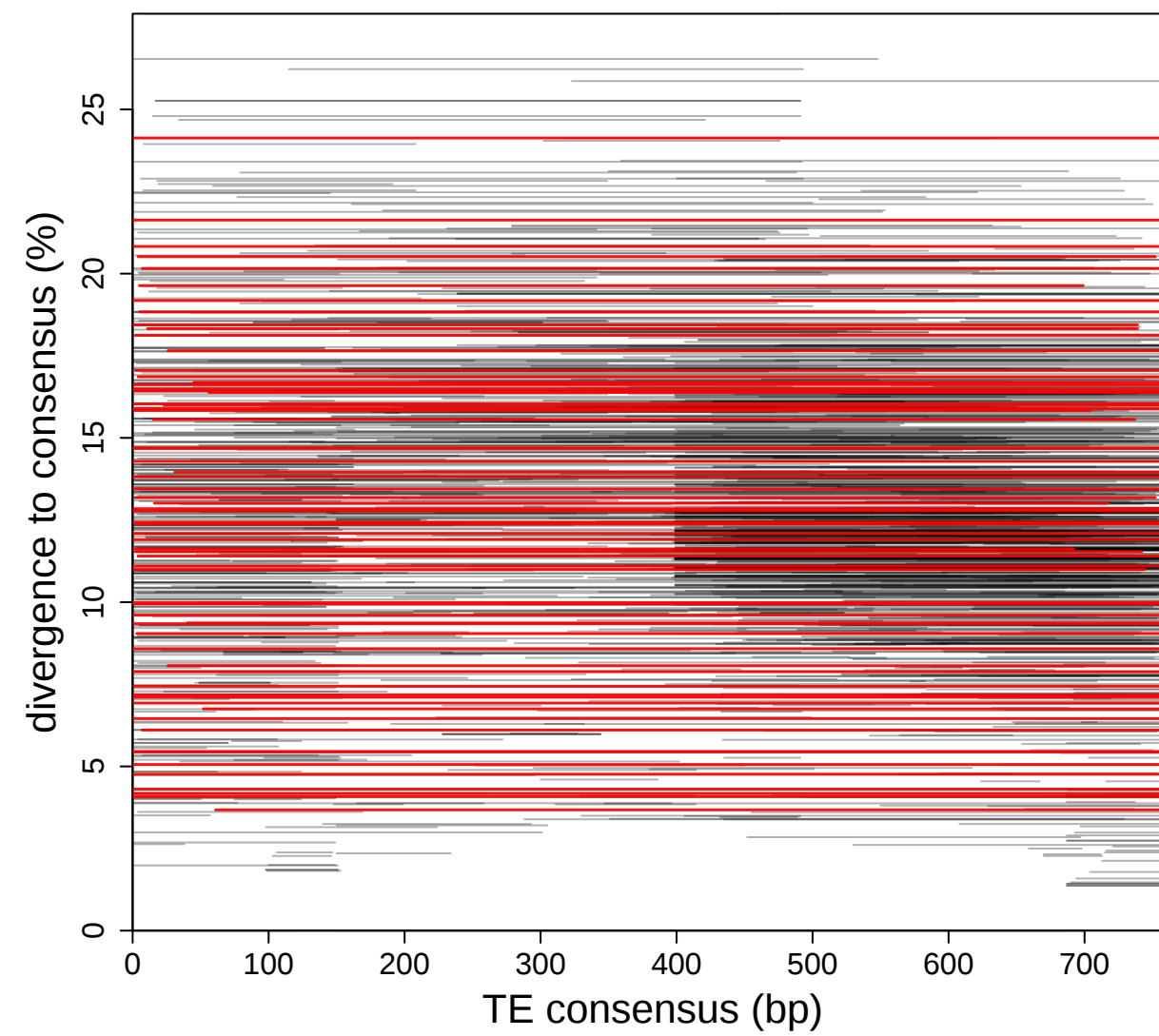
MSA length = 765





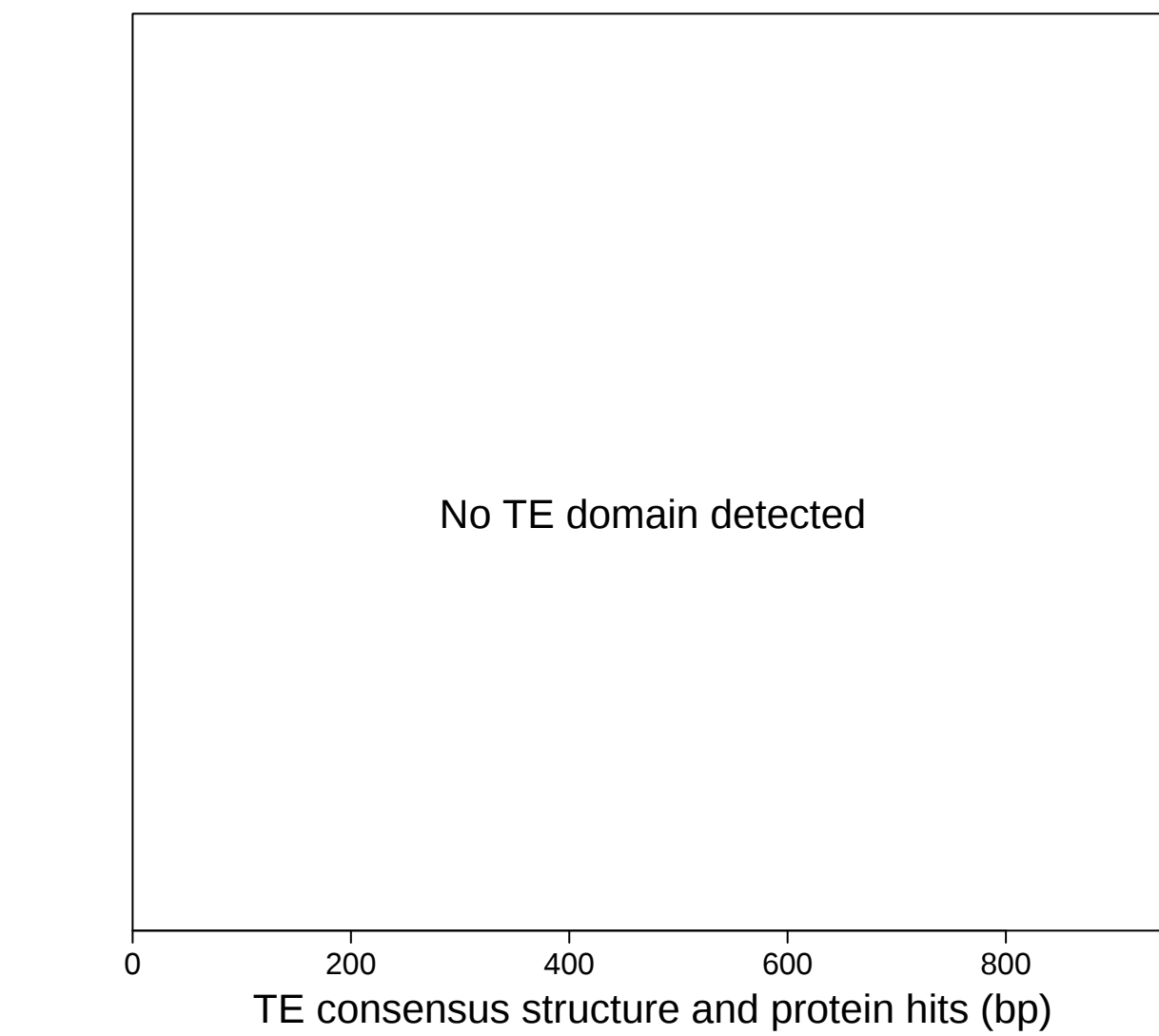
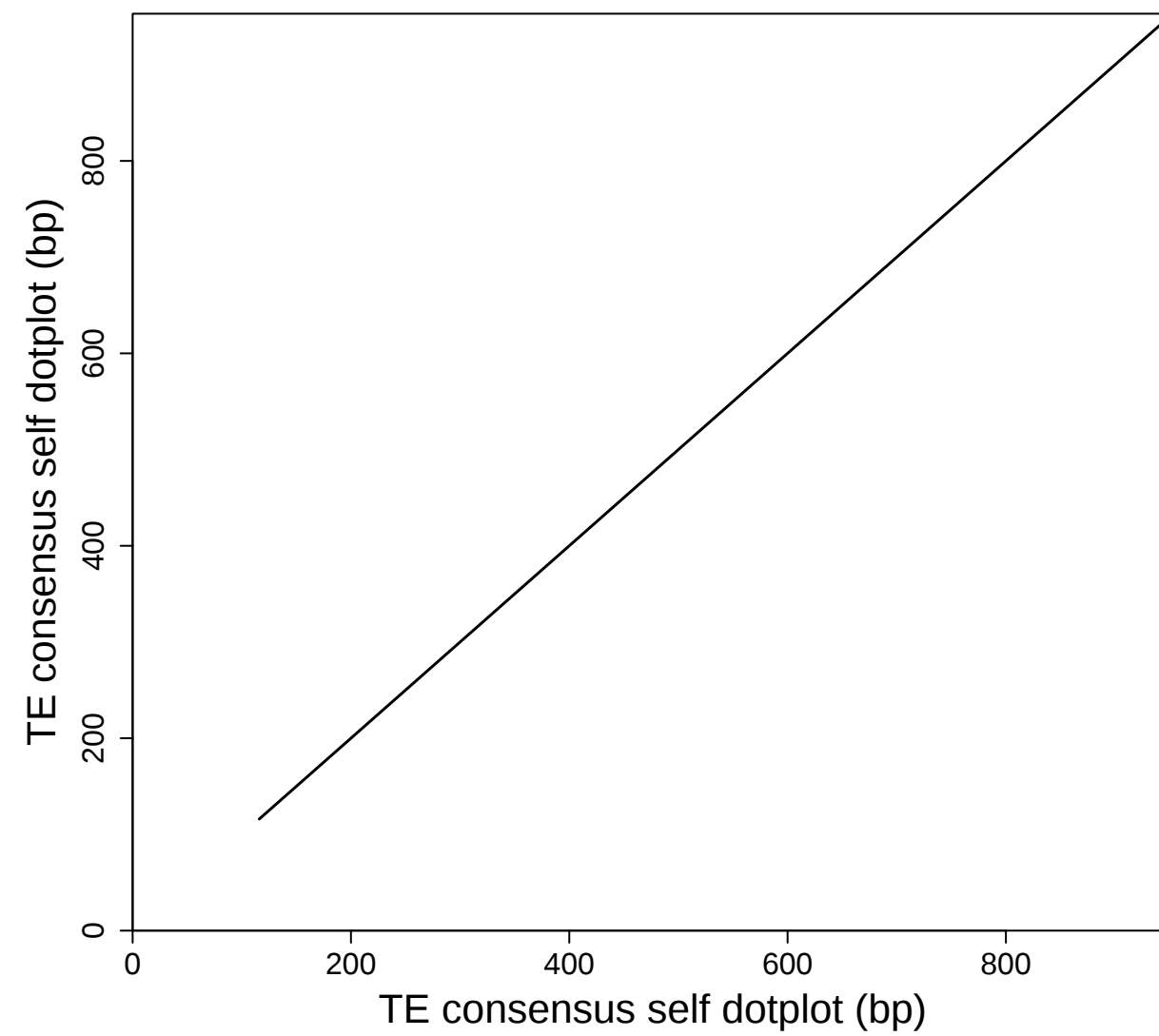
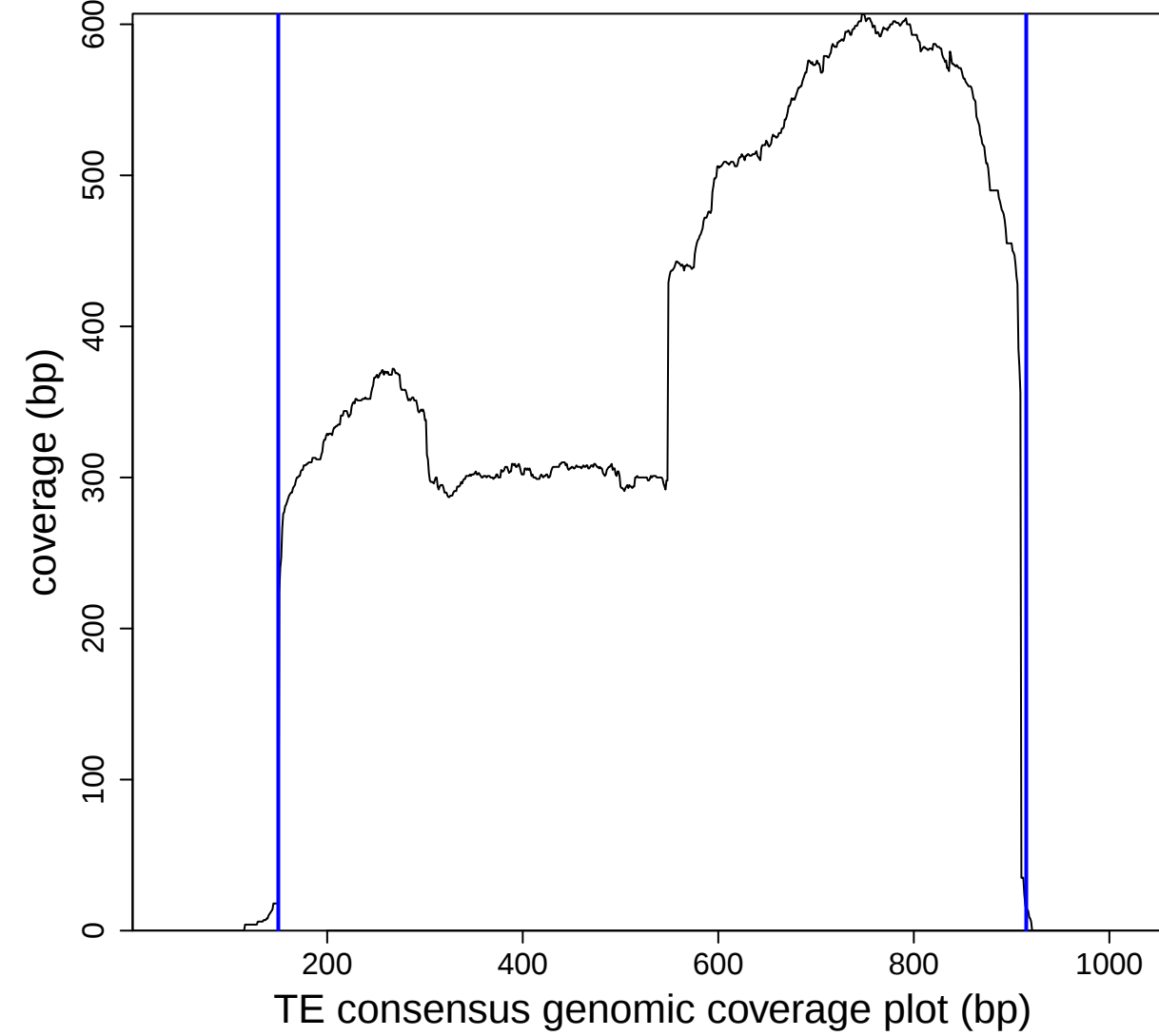
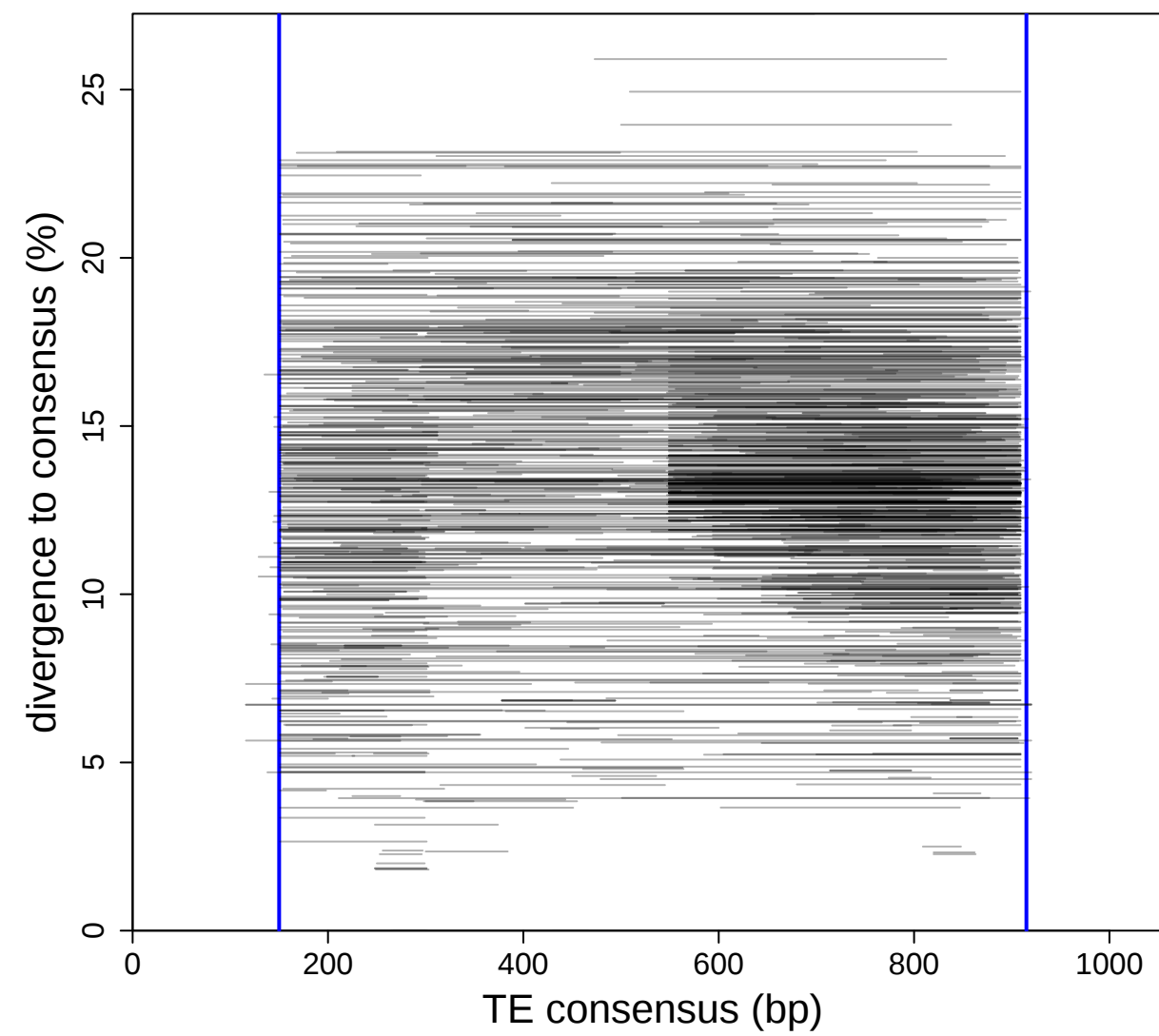
a.b.bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.f
size: 765bp; fragments: 1261; full length: 67 (≥ 688.5 bp)

After TEtrimmer 765 bp

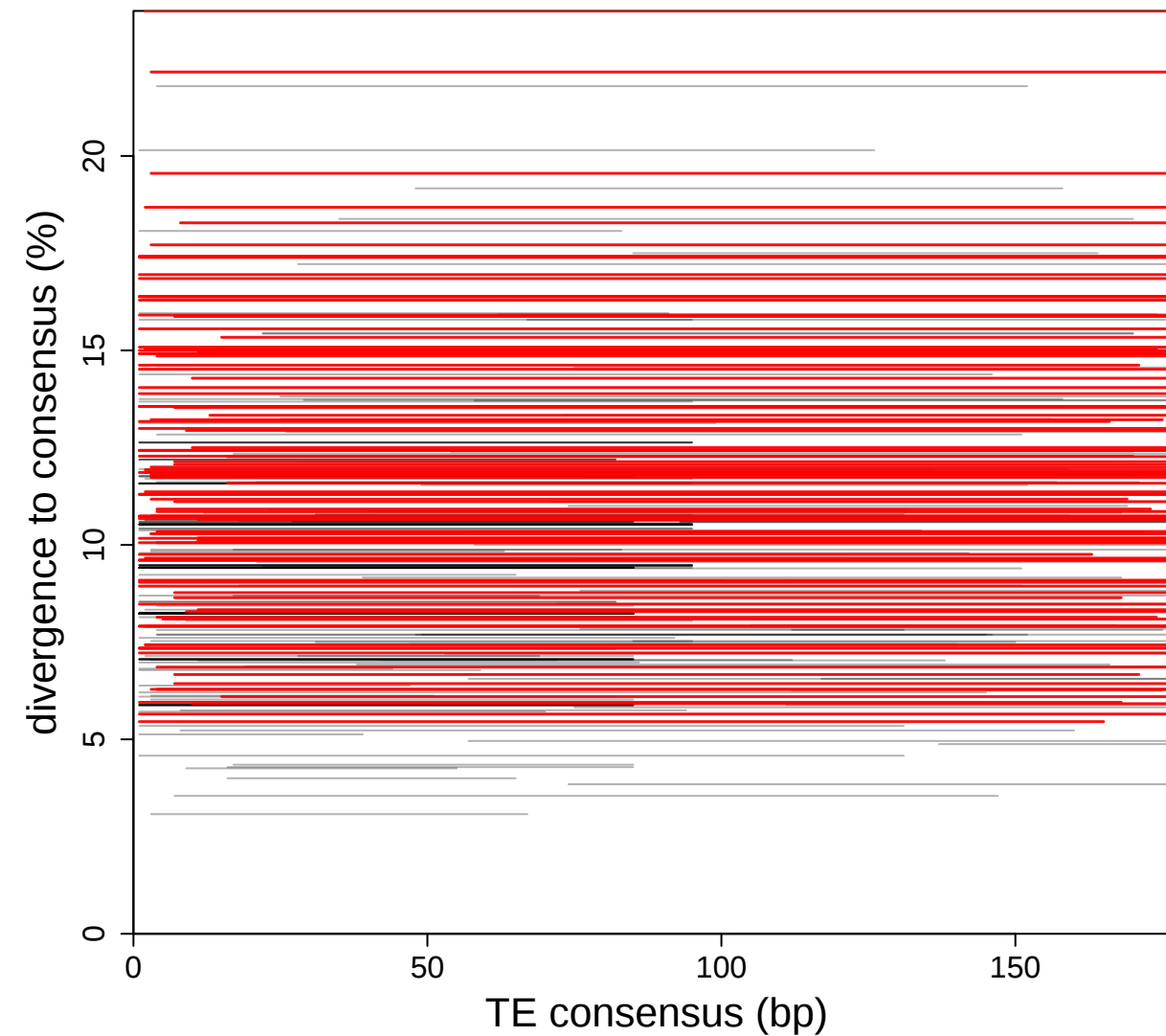


6.fasta.b.bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.f
size: 1065bp; fragments: 1065; full length: 0 (>=958.5bp)

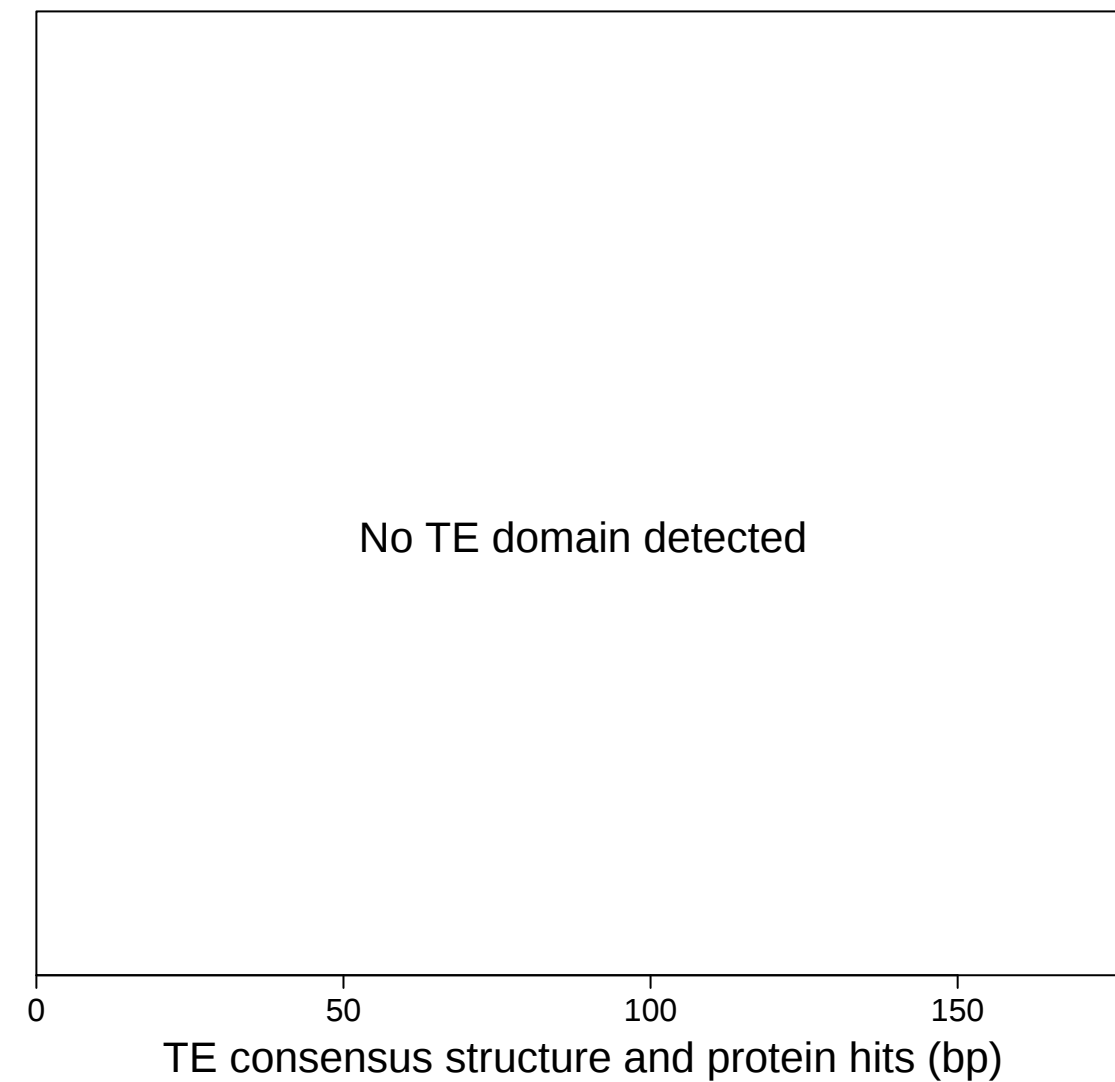
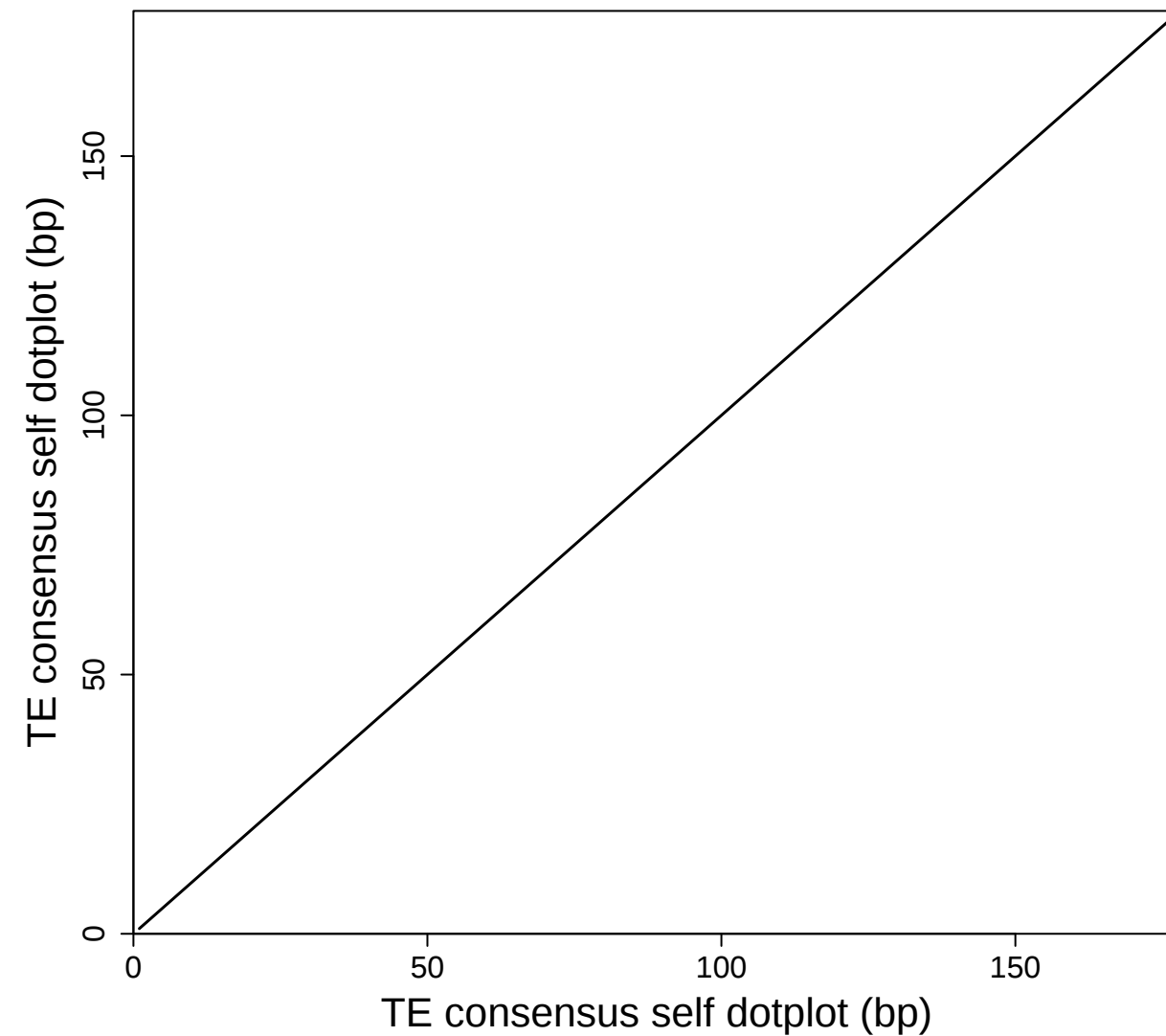
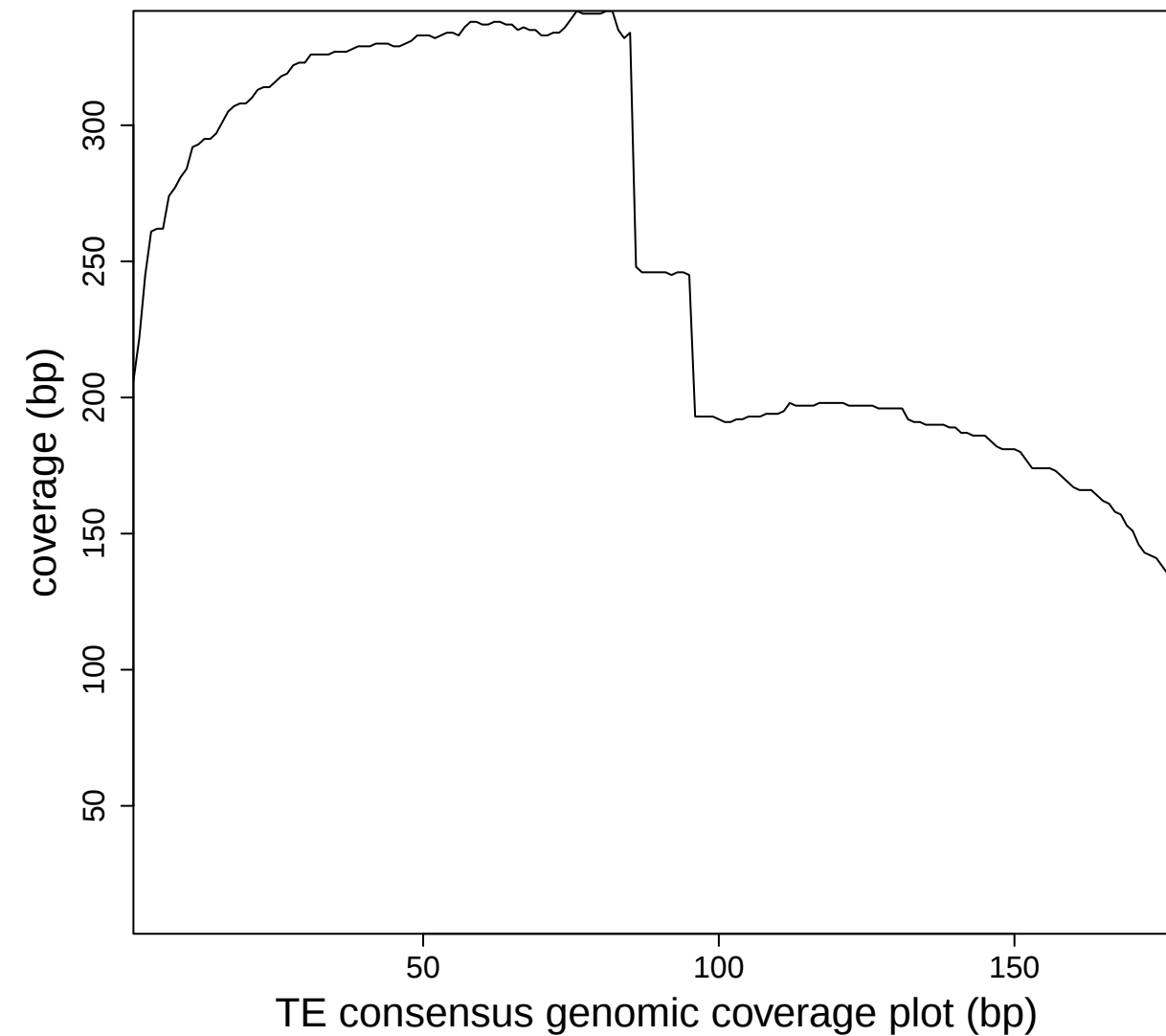
After TEtrimmer Extended plot Blue lines are boundaries



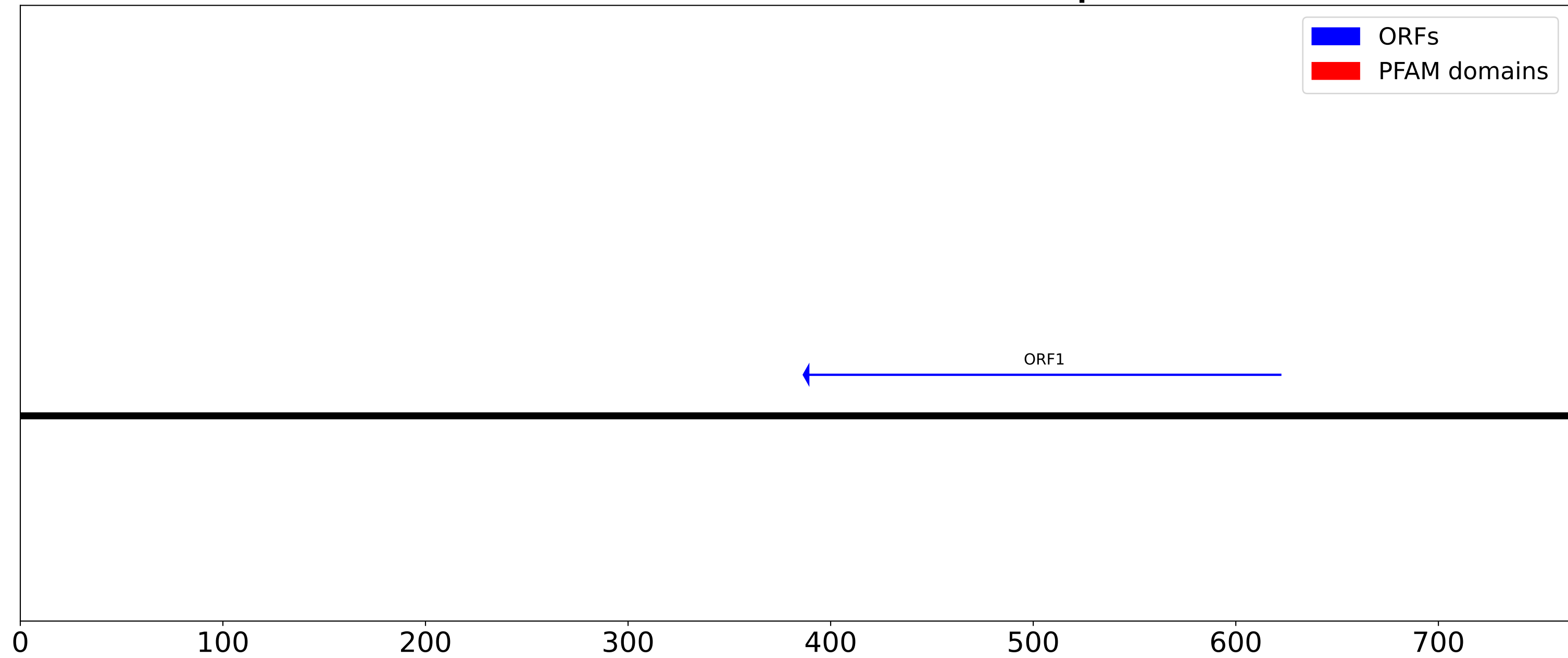
TE: rnd_5_family_5696
size: 178bp; fragments: 370; full length: 111 (≥ 160.2 bp)



Before TEtrimmer 178 bp



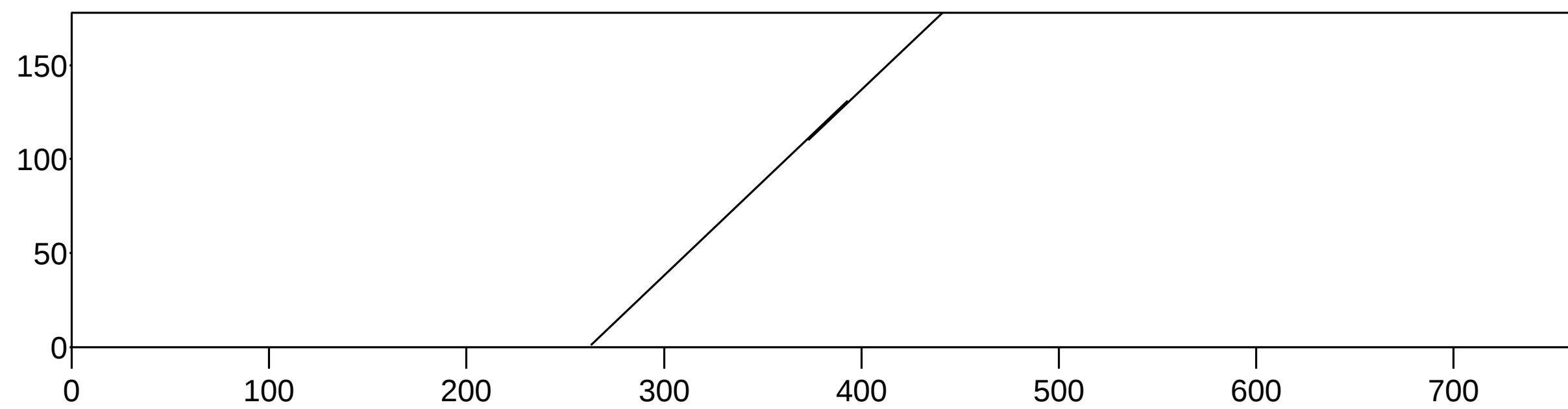
After TEtrimmer ORF and PFAM domain plot



Dotplot

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

onsensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)