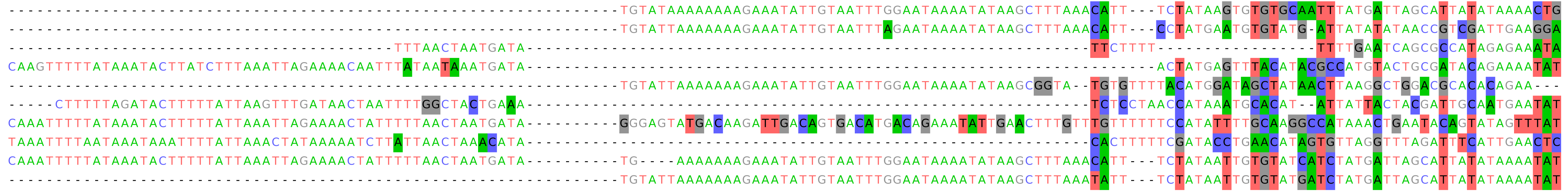
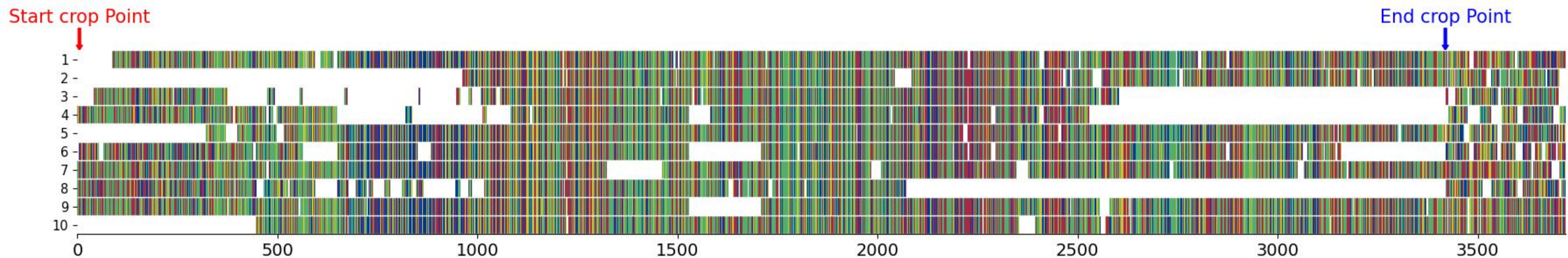


Start Crop Point

End Crop Point

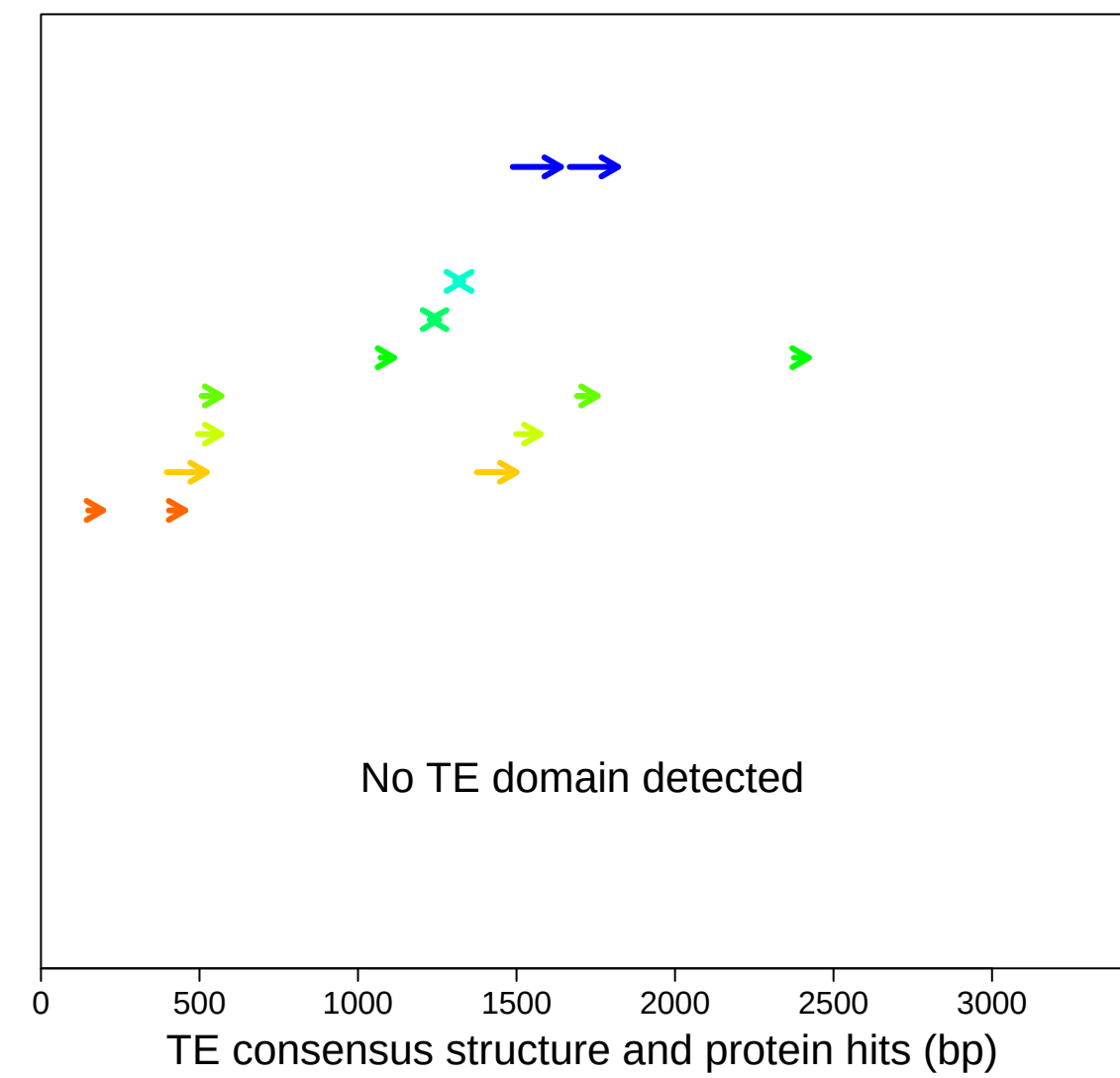
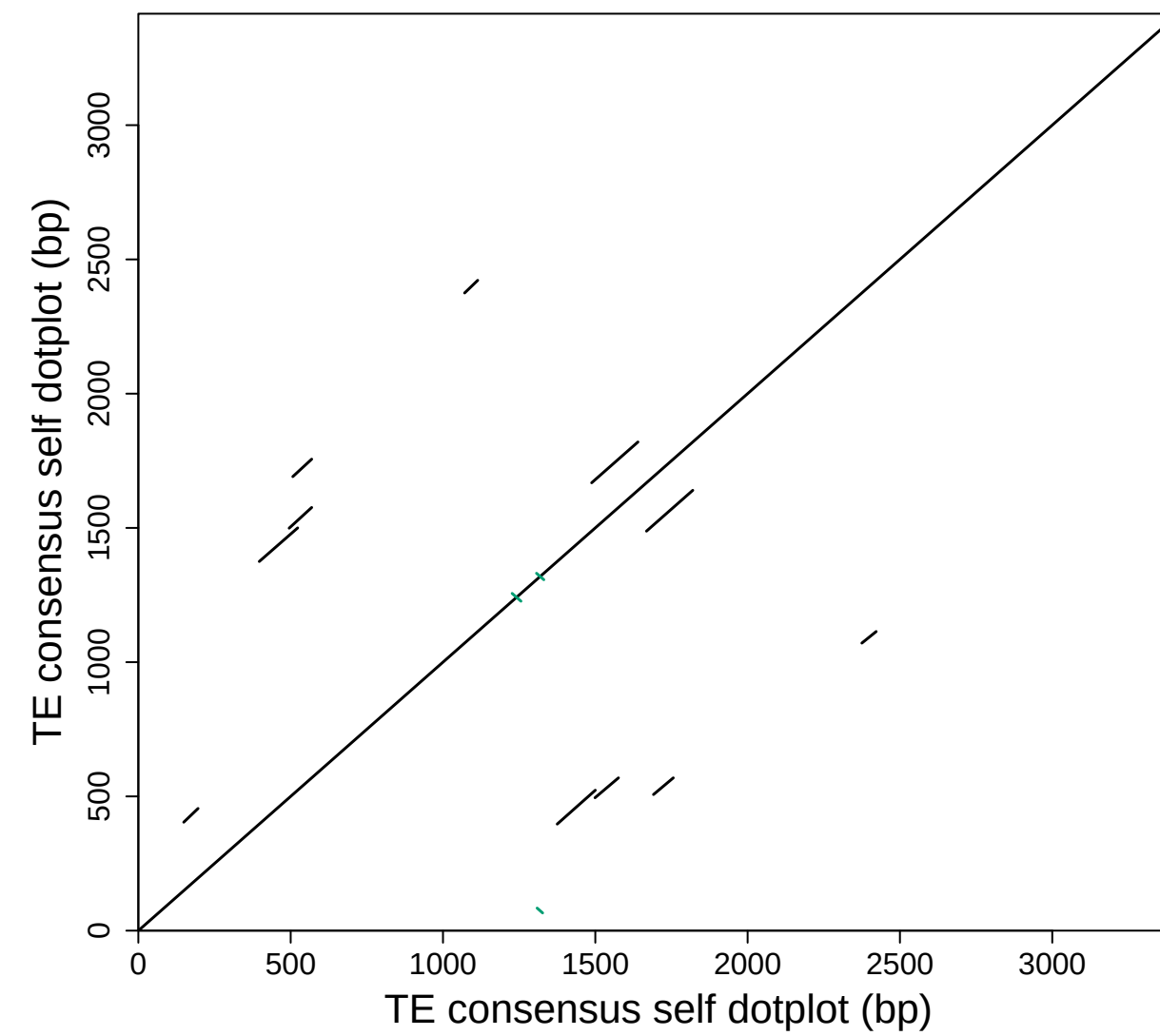
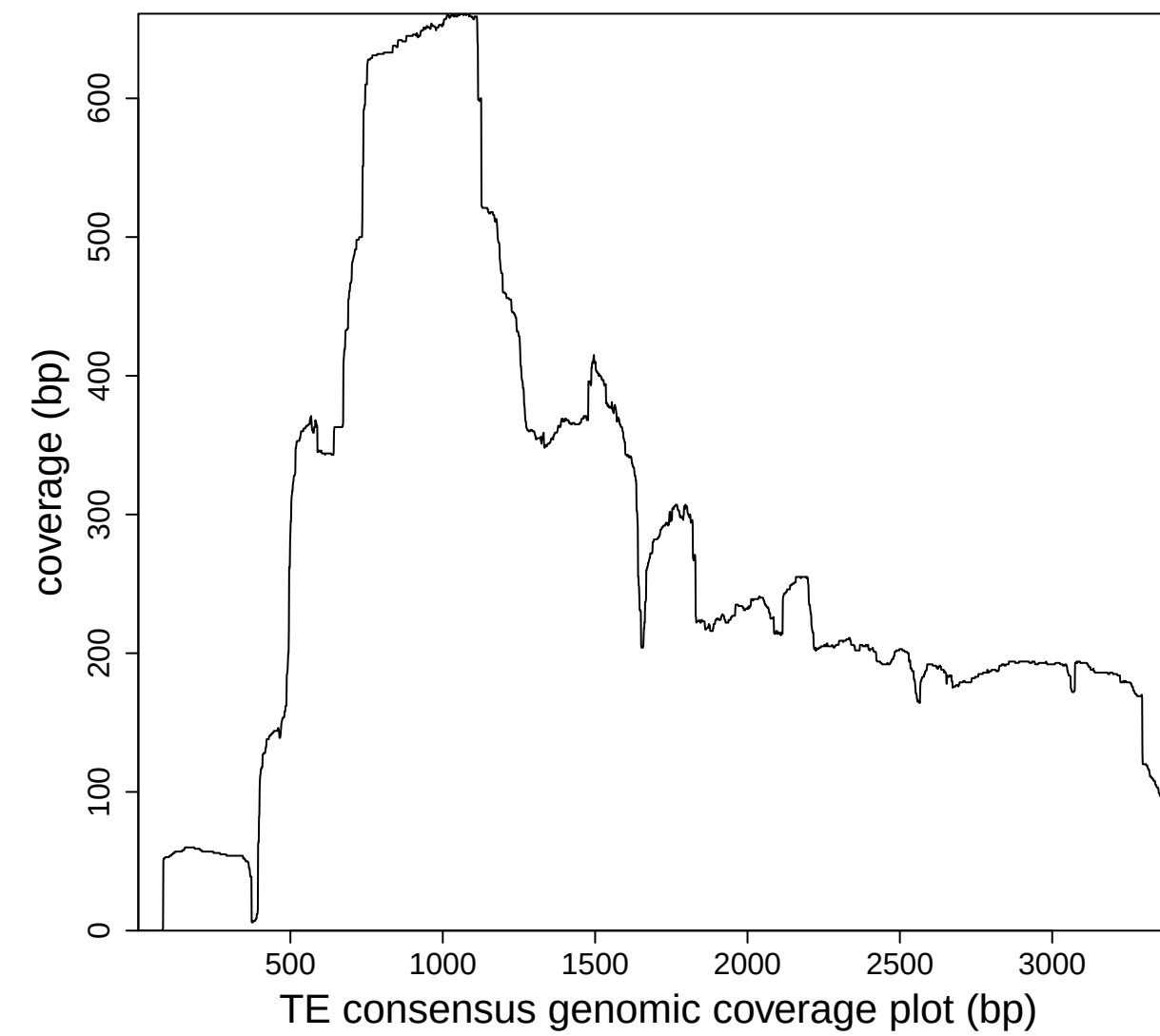
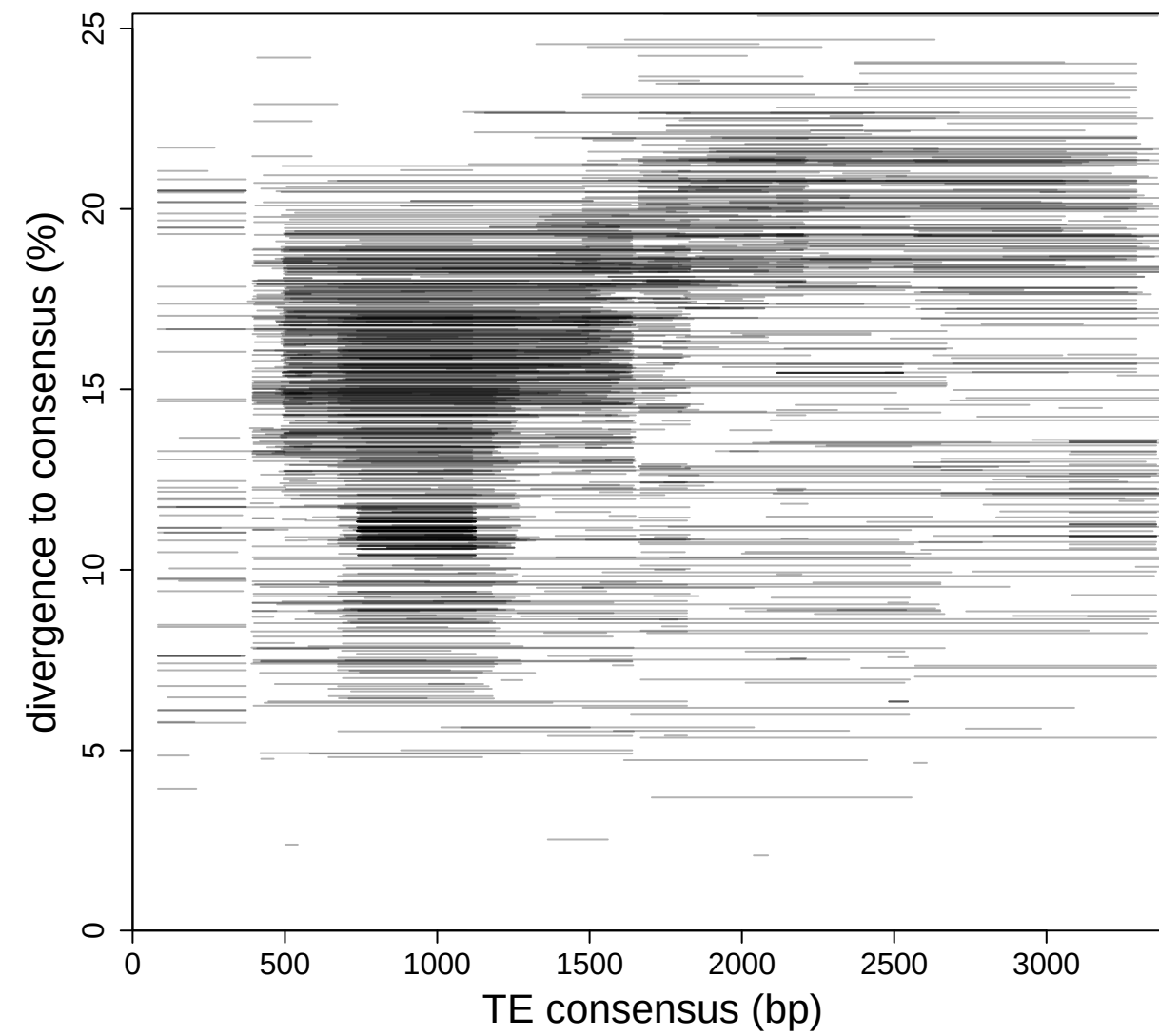
MSA length = 3415





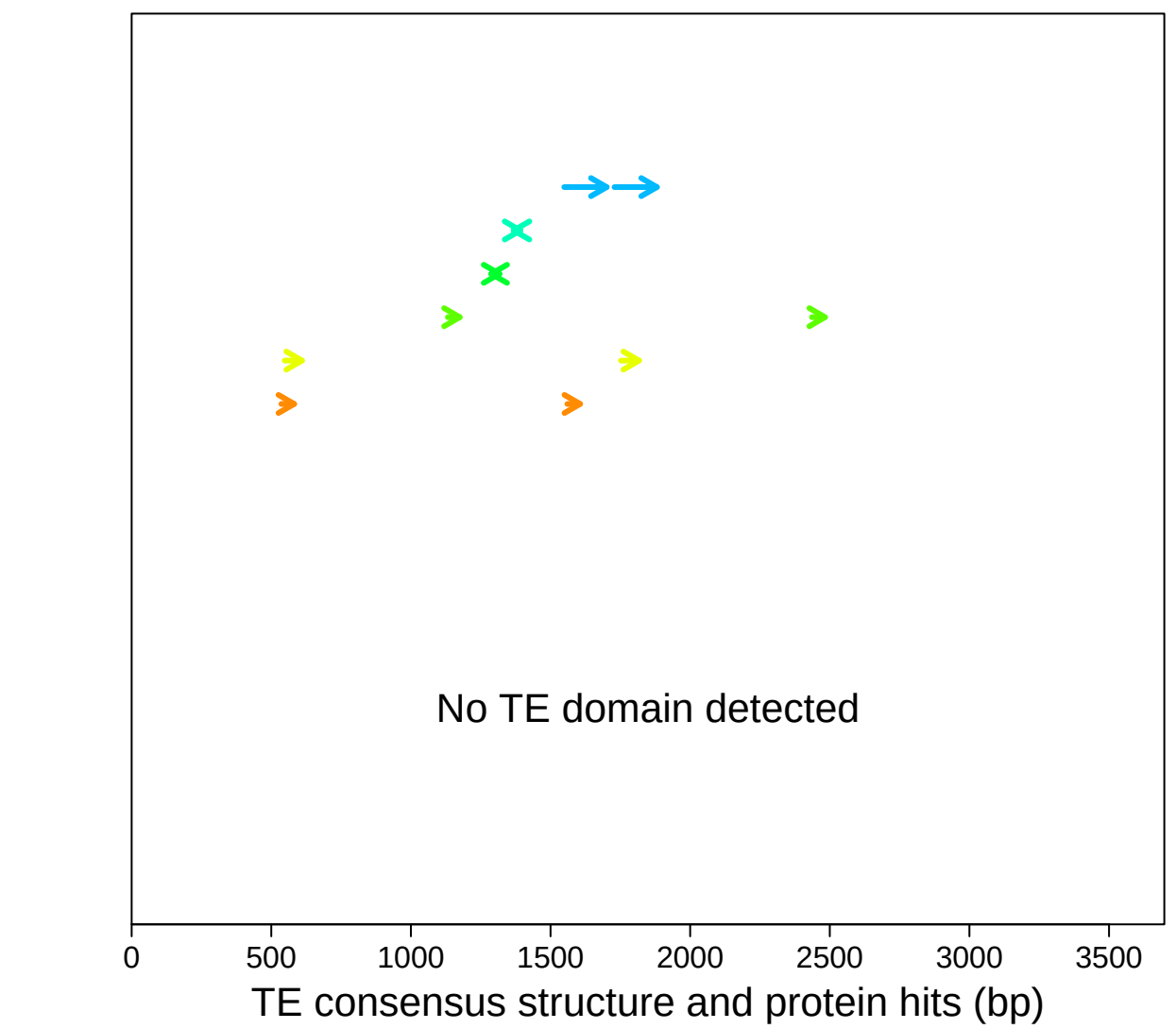
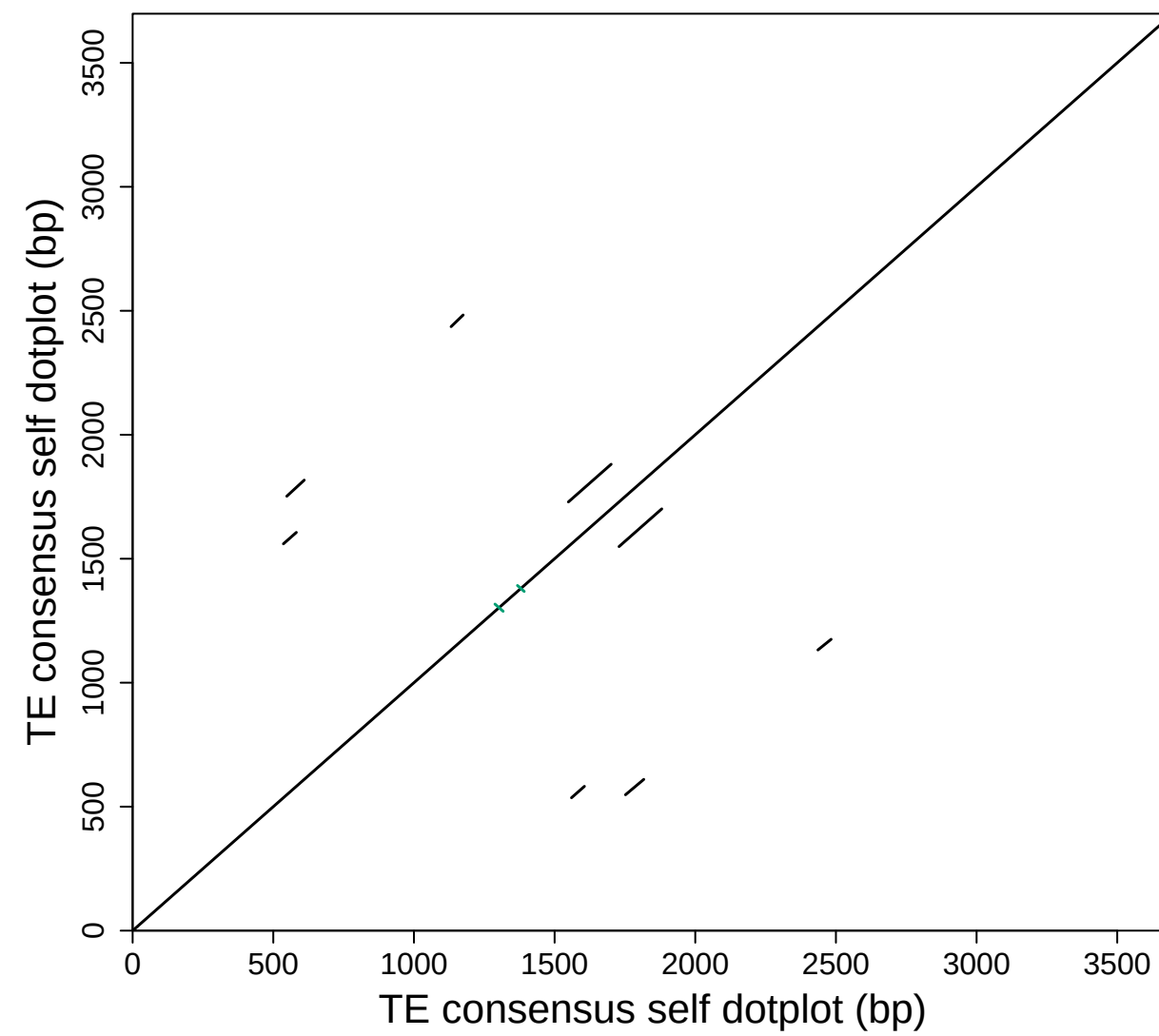
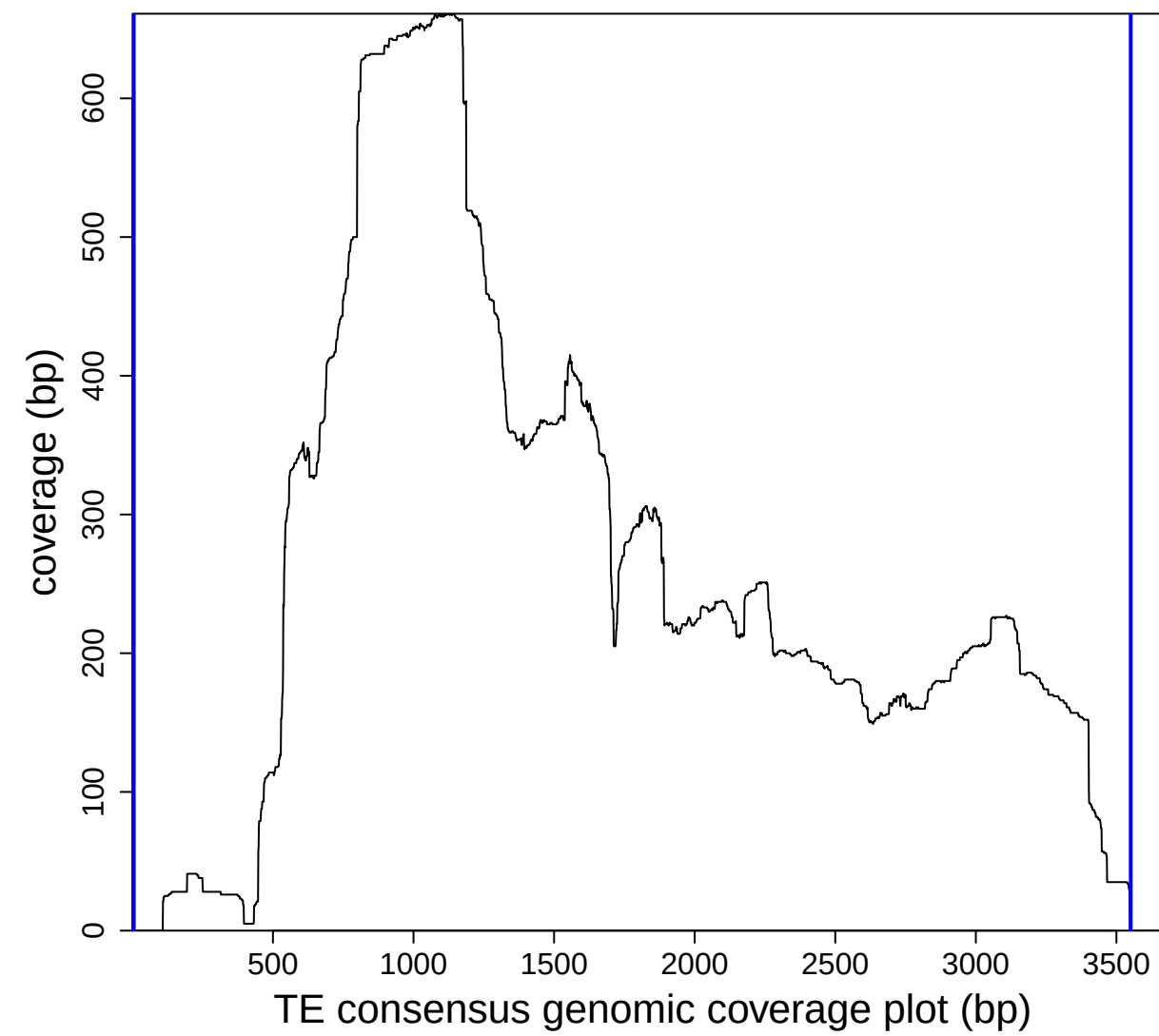
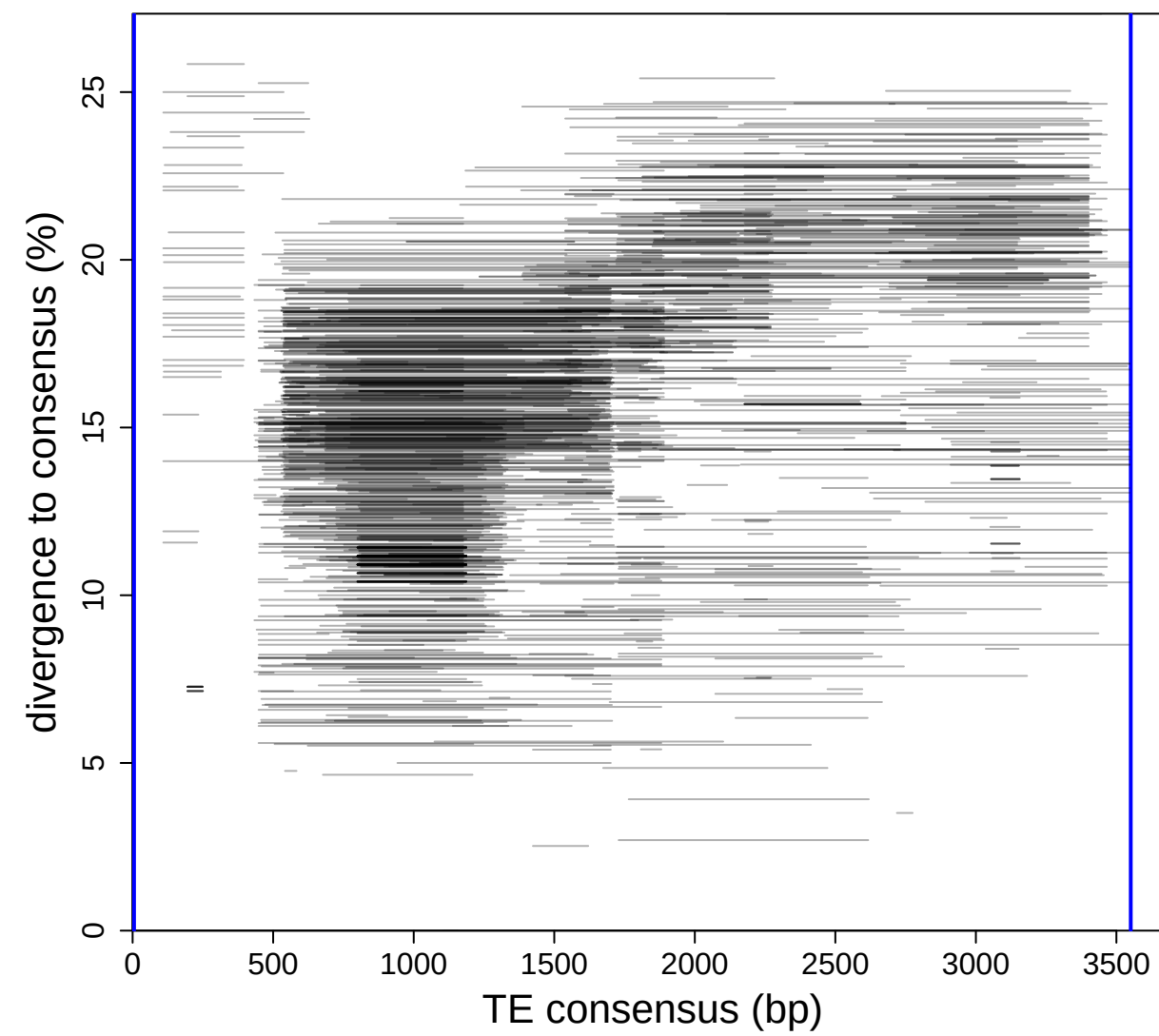
After TEtrimmer 3415 bp

b.bed uf.bed g 1.bed fm 1.bed 0 0 bcln.fa aln.fa cl.fa gs.fa
size: 3415bp; fragments: 1348; full length: 0 (≥ 3073.5 bp)

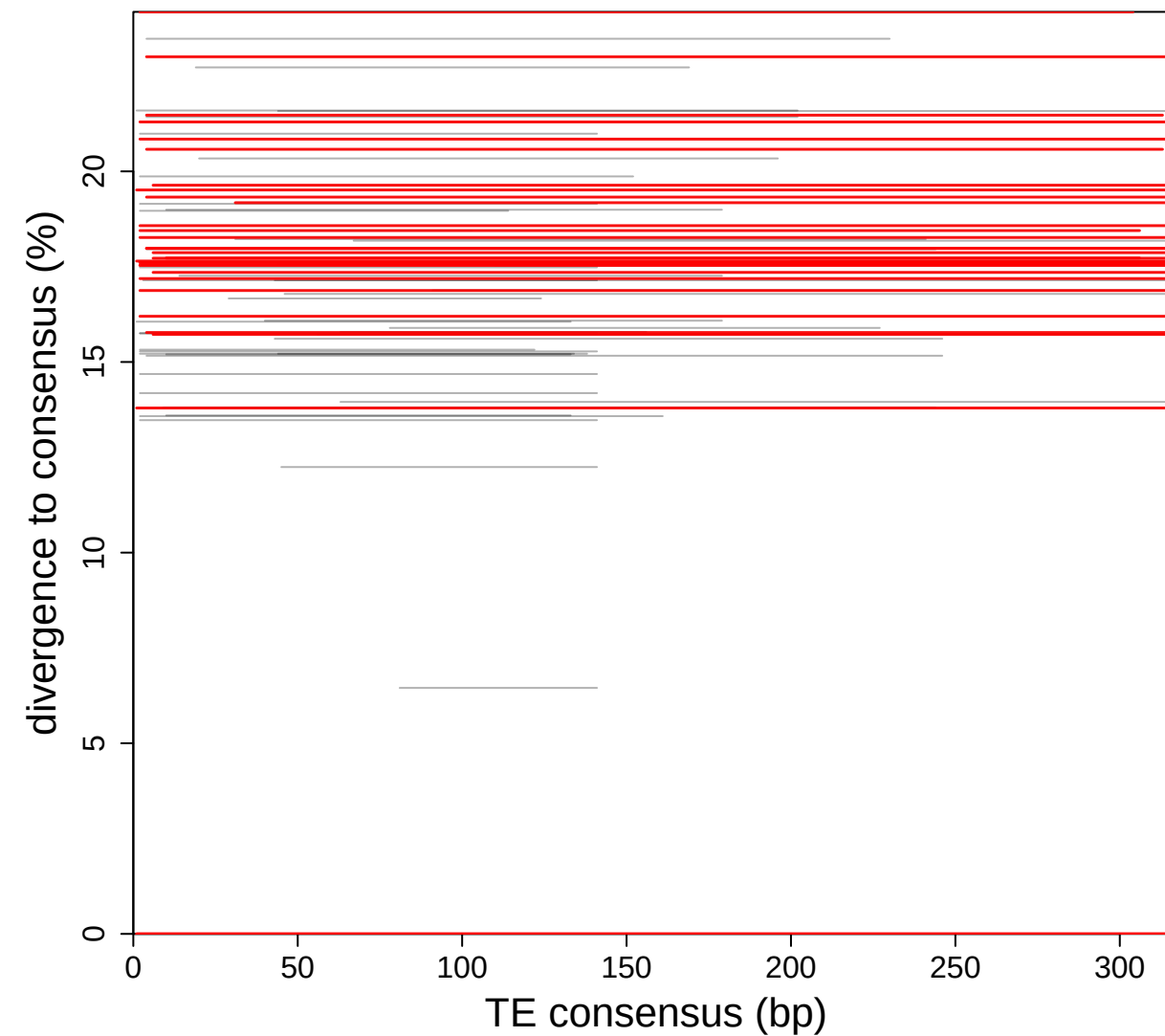


.fasta.b.bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa
size: 3701bp; fragments: 1324; full length: 0 (>=3330.9bp)

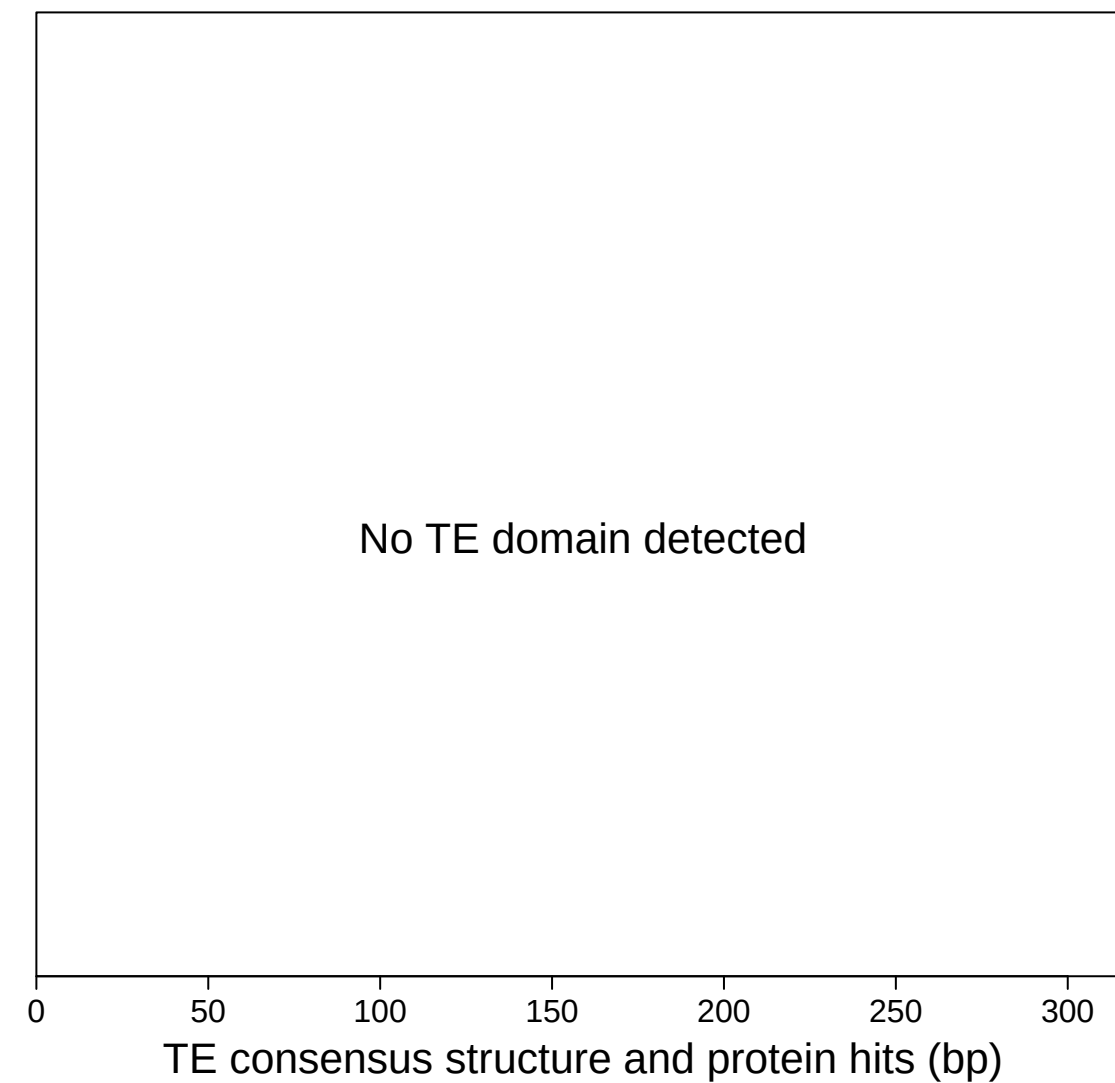
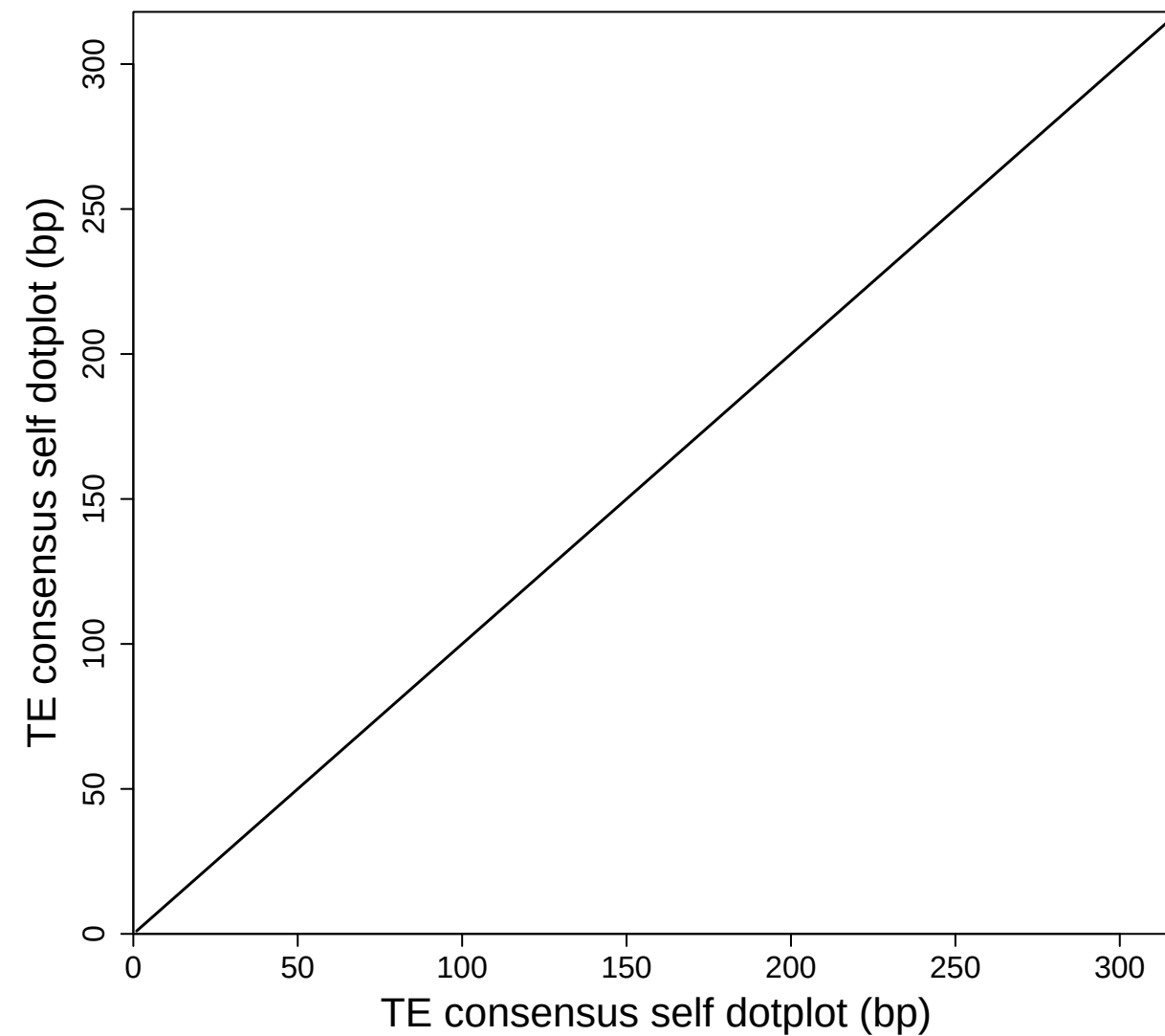
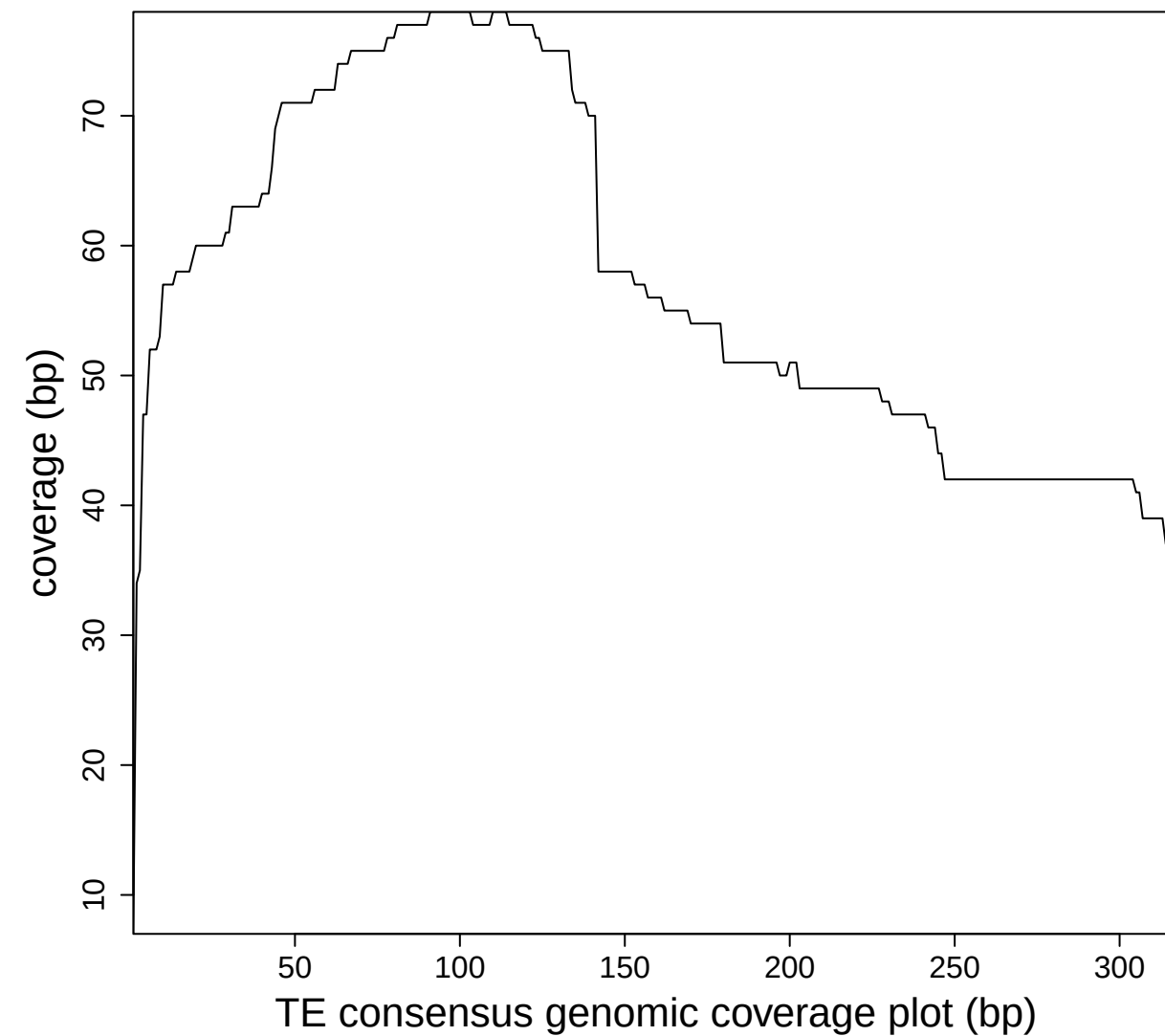
After TEtrimmer Extended plot Blue lines are boundaries



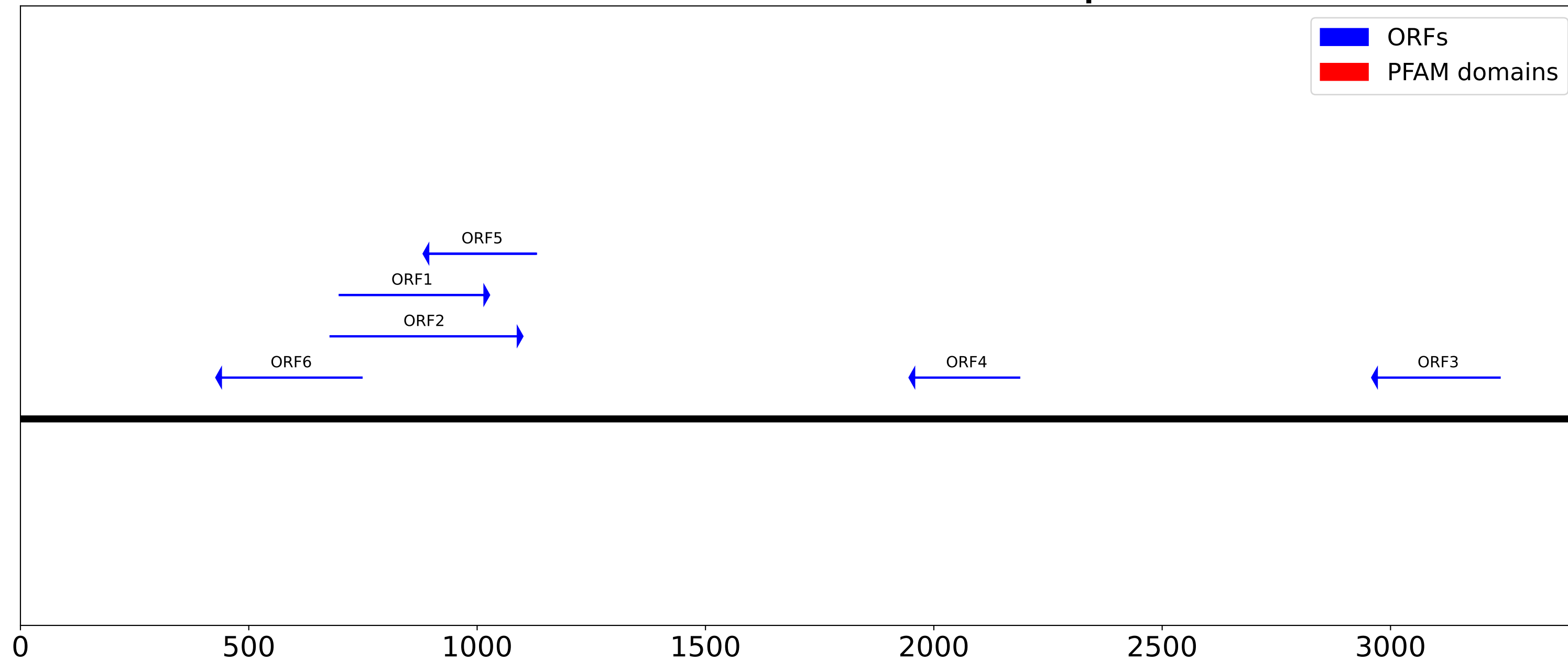
TE: ltr_1_family_226
size: 318bp; fragments: 80; full length: 32 (≥ 286.2 bp)



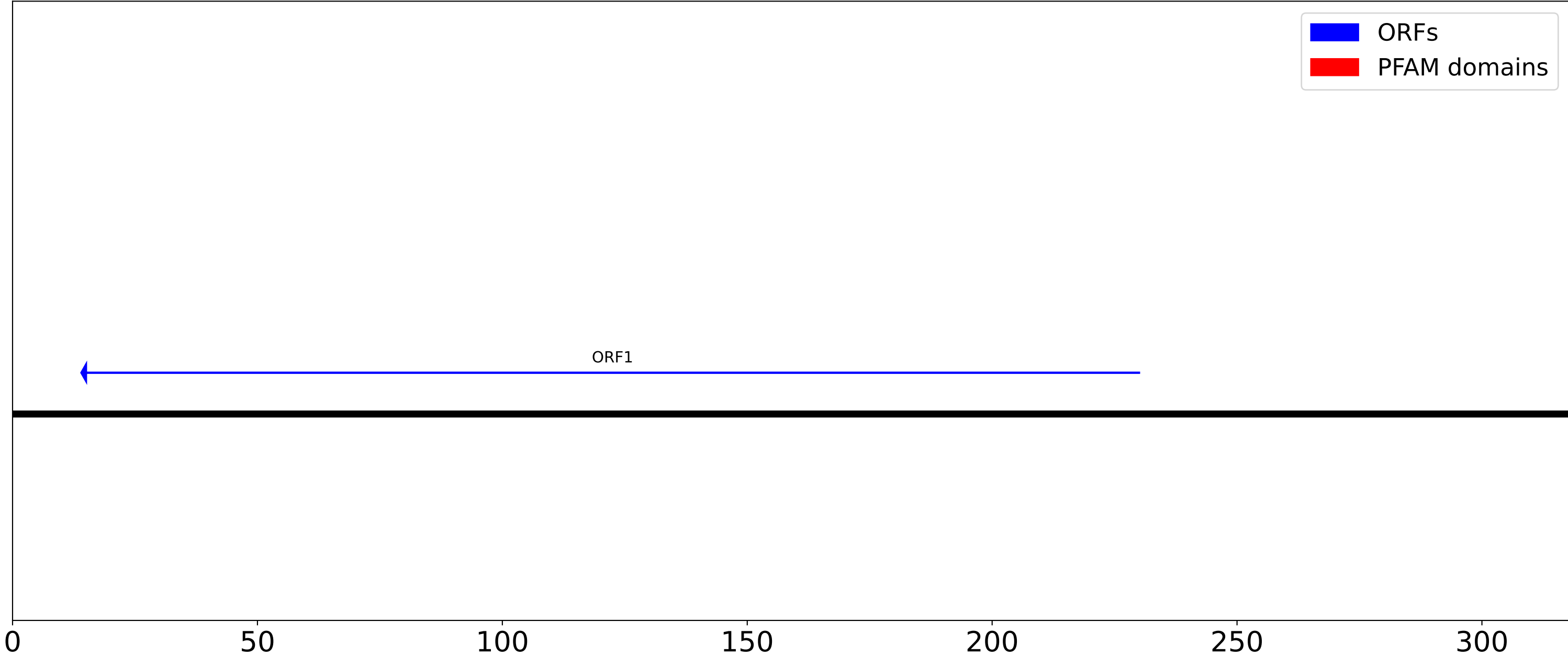
Before TEtrimmer 318 bp



After TEtrimmer ORF and PFAM domain plot



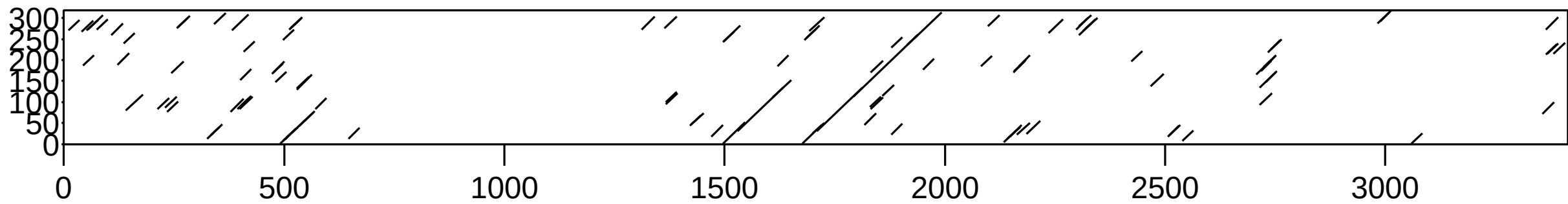
Before TEtrimmer ORF and PFAM domain plot



Dotplot

(windowsize = 25, threshold = 50.00 09/10/25)

ensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)