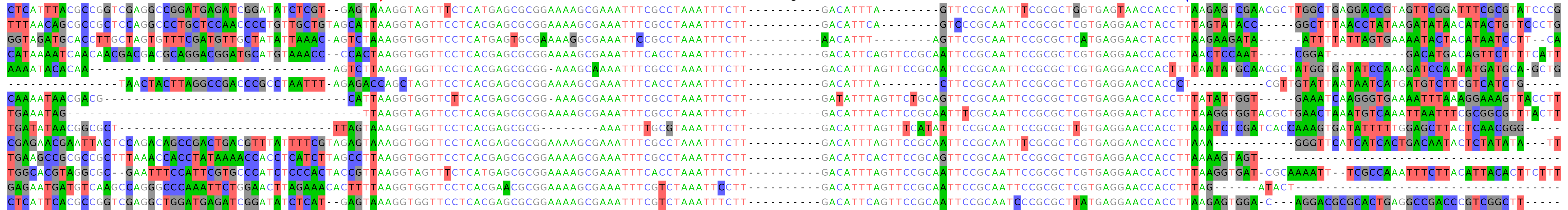
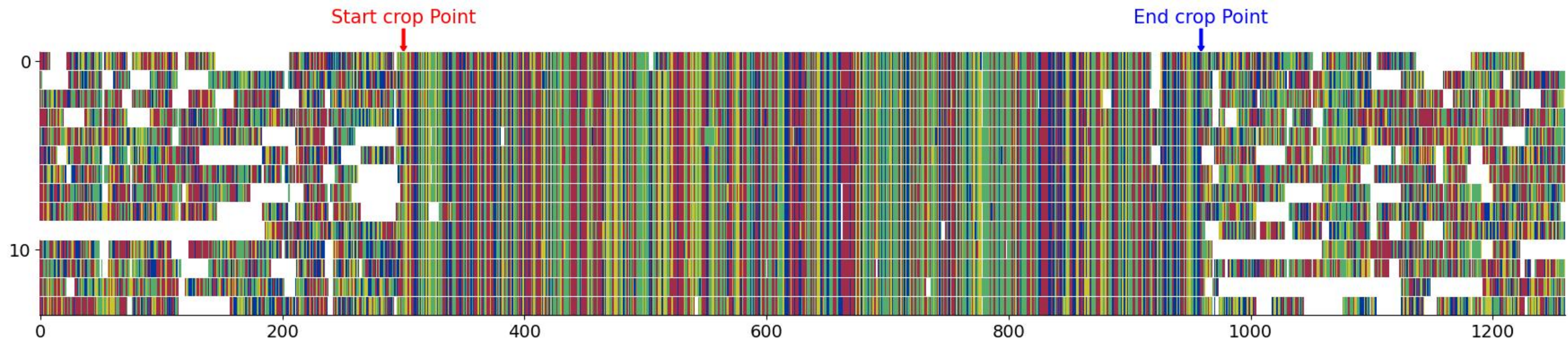


Start crop Point

End crop Point

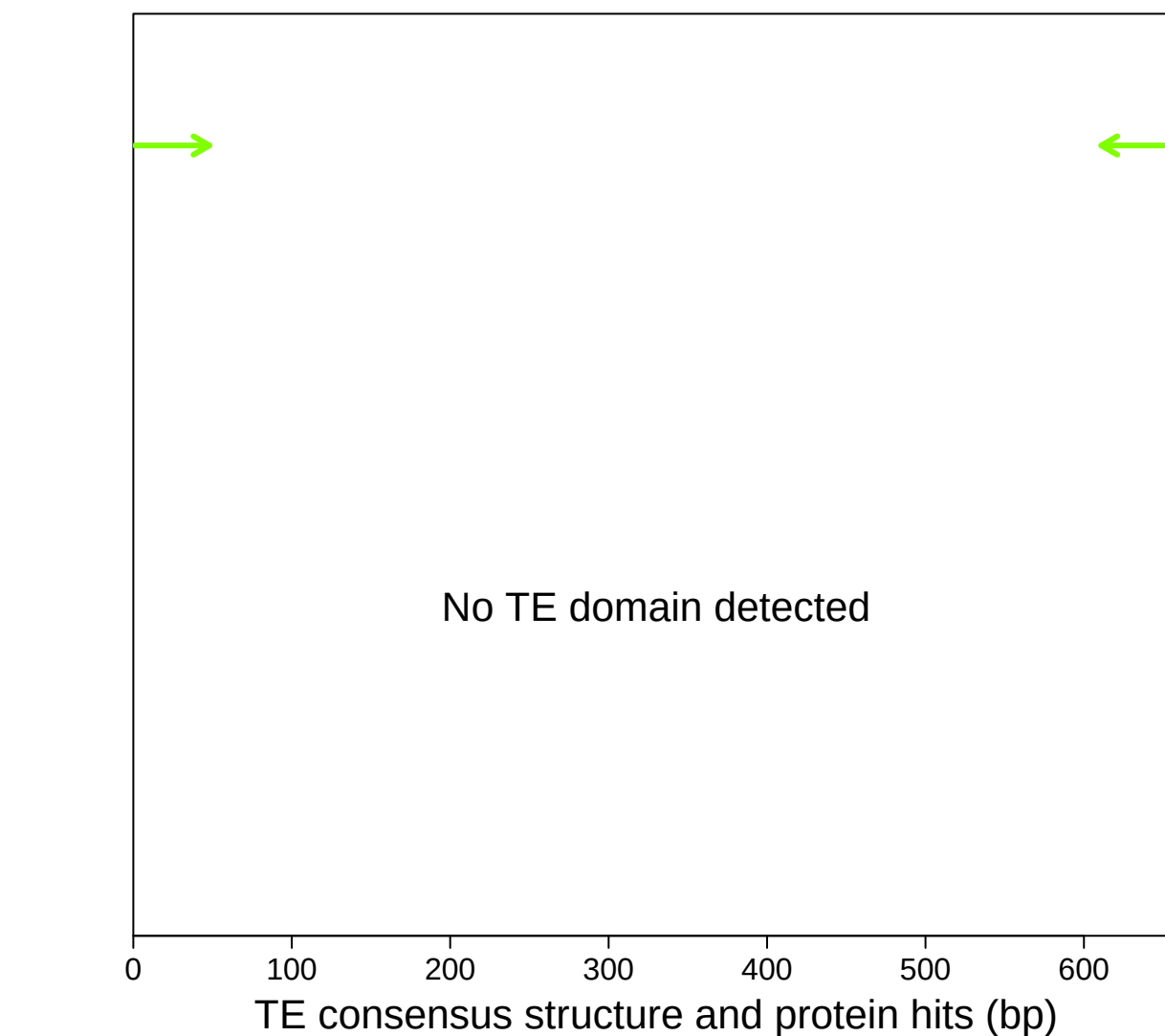
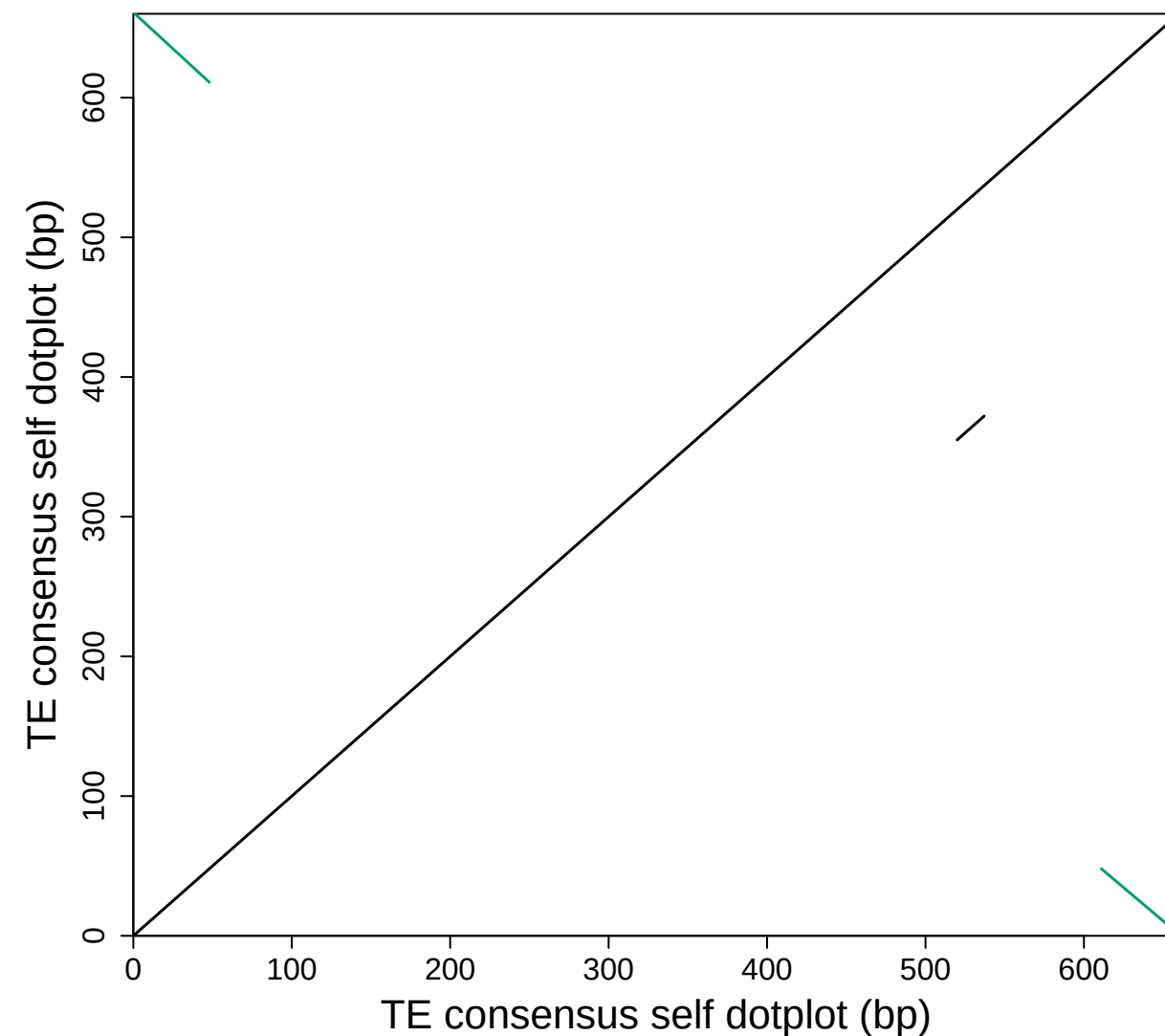
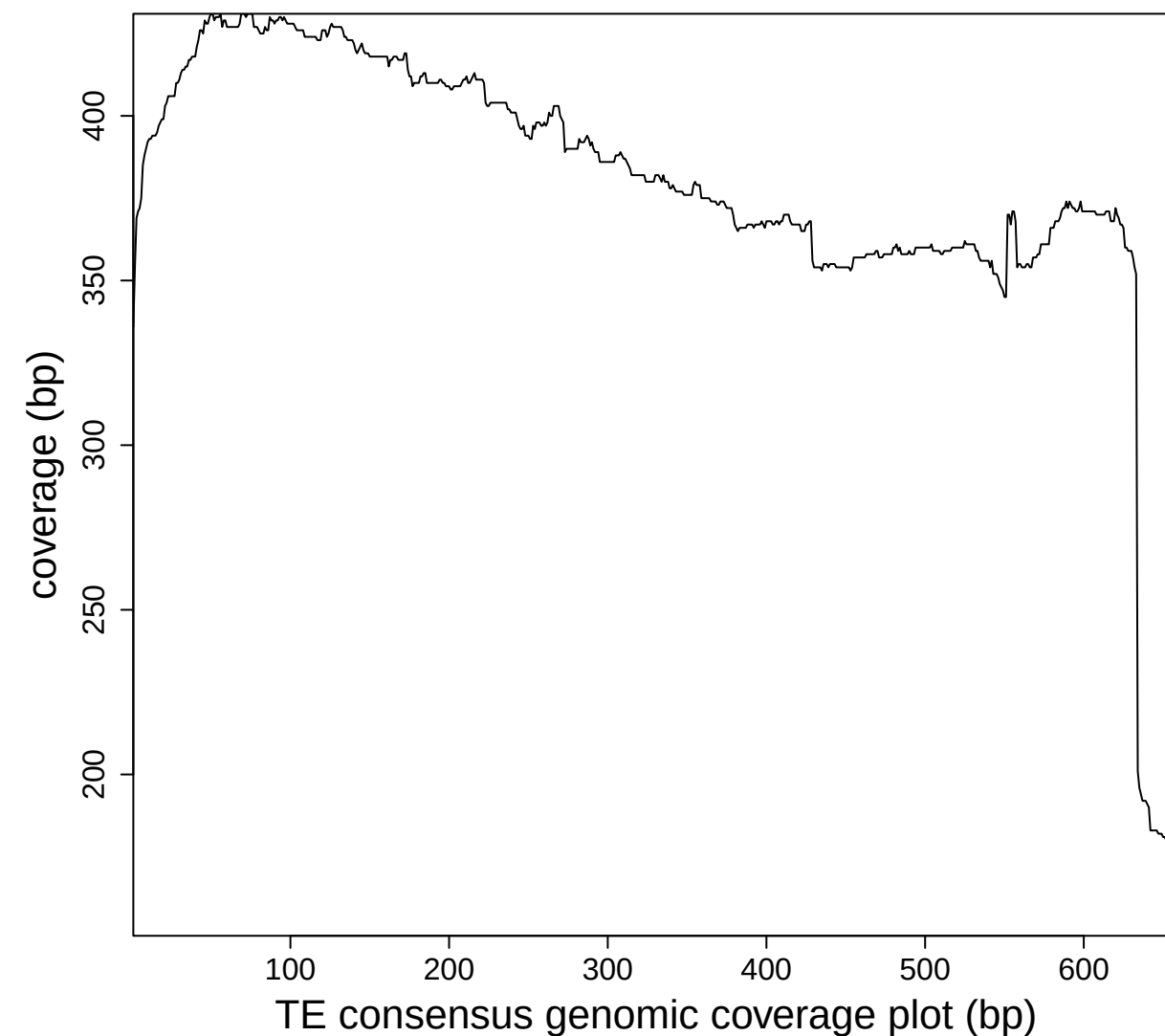
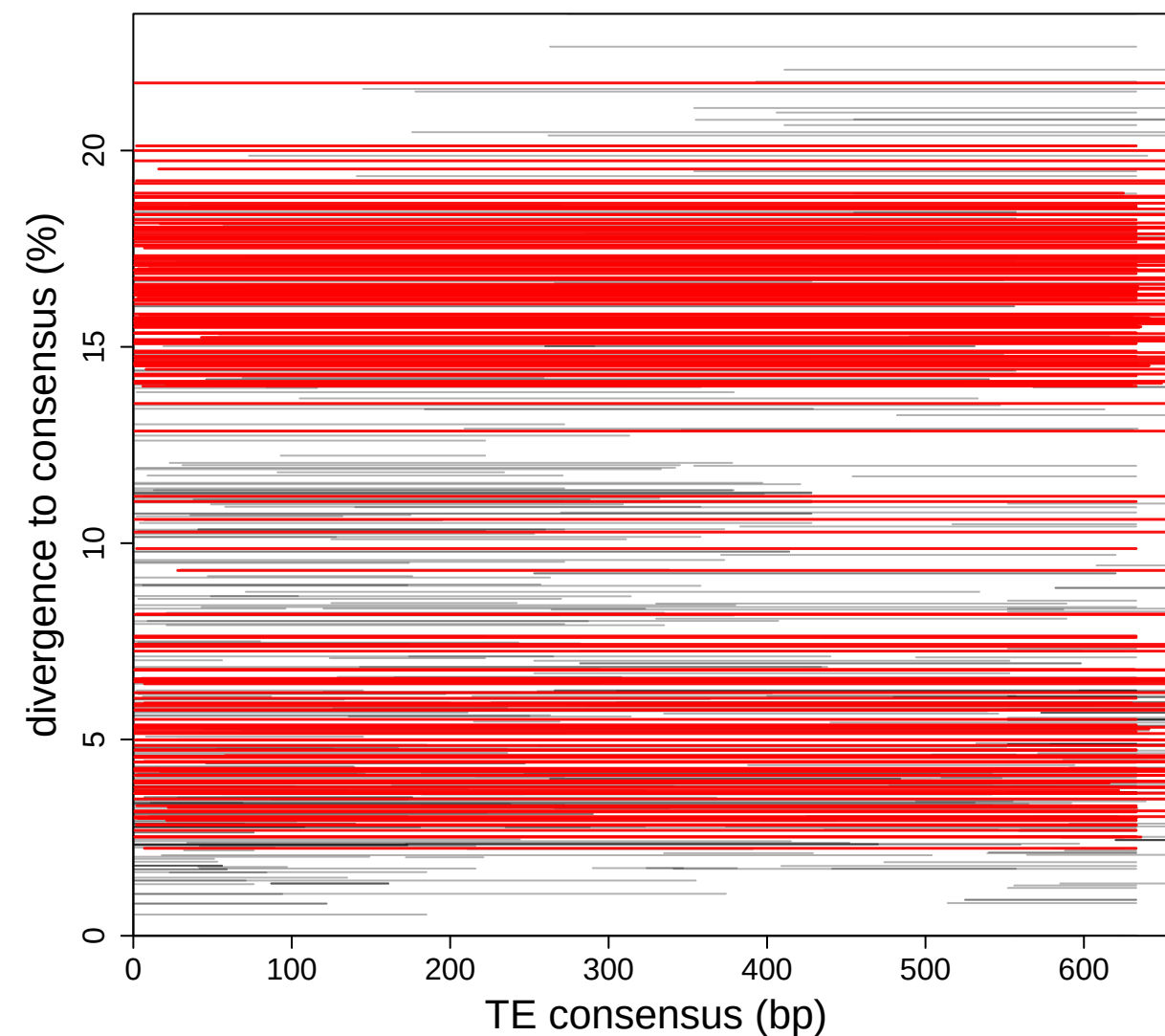
MSA length = 660





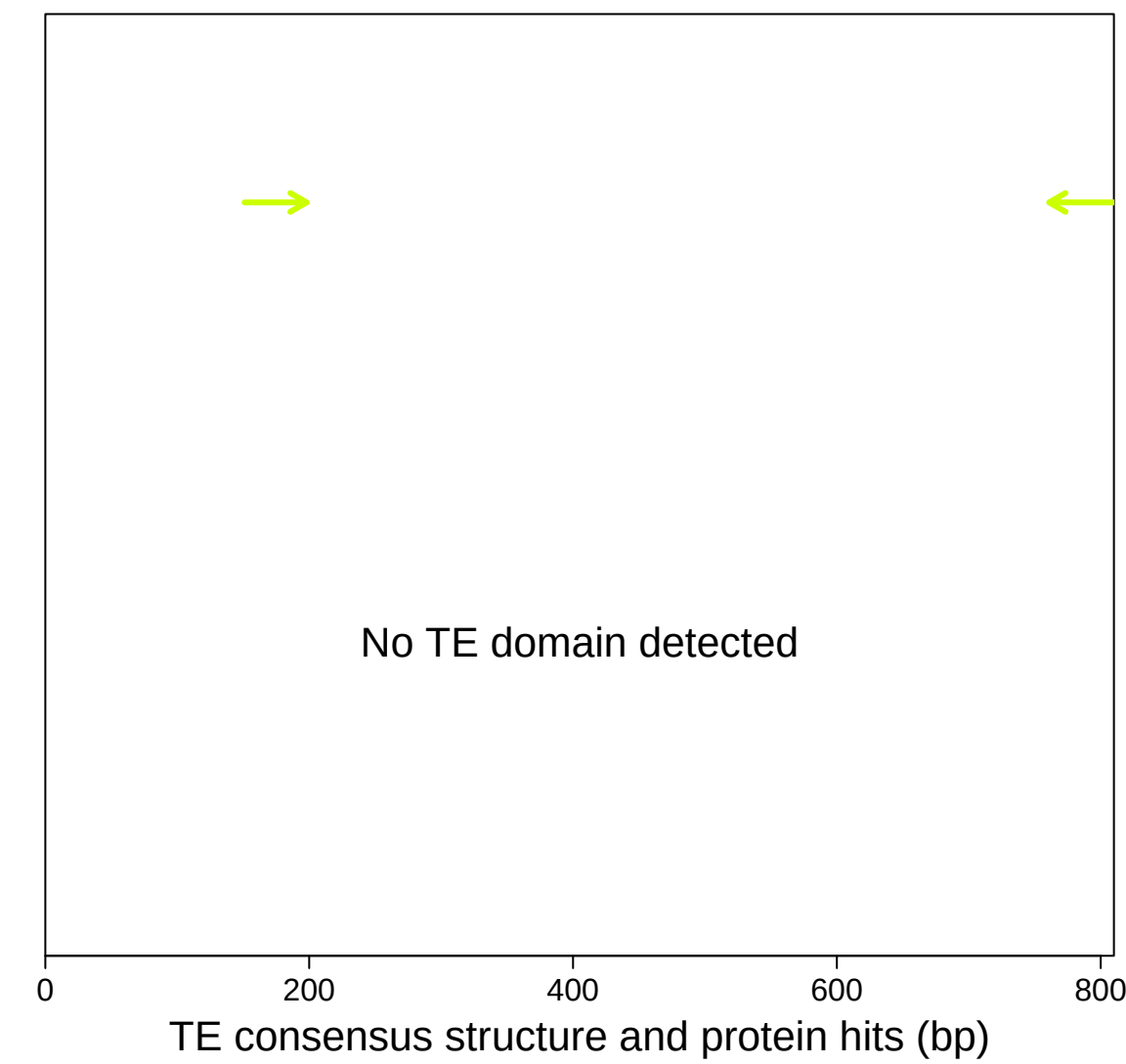
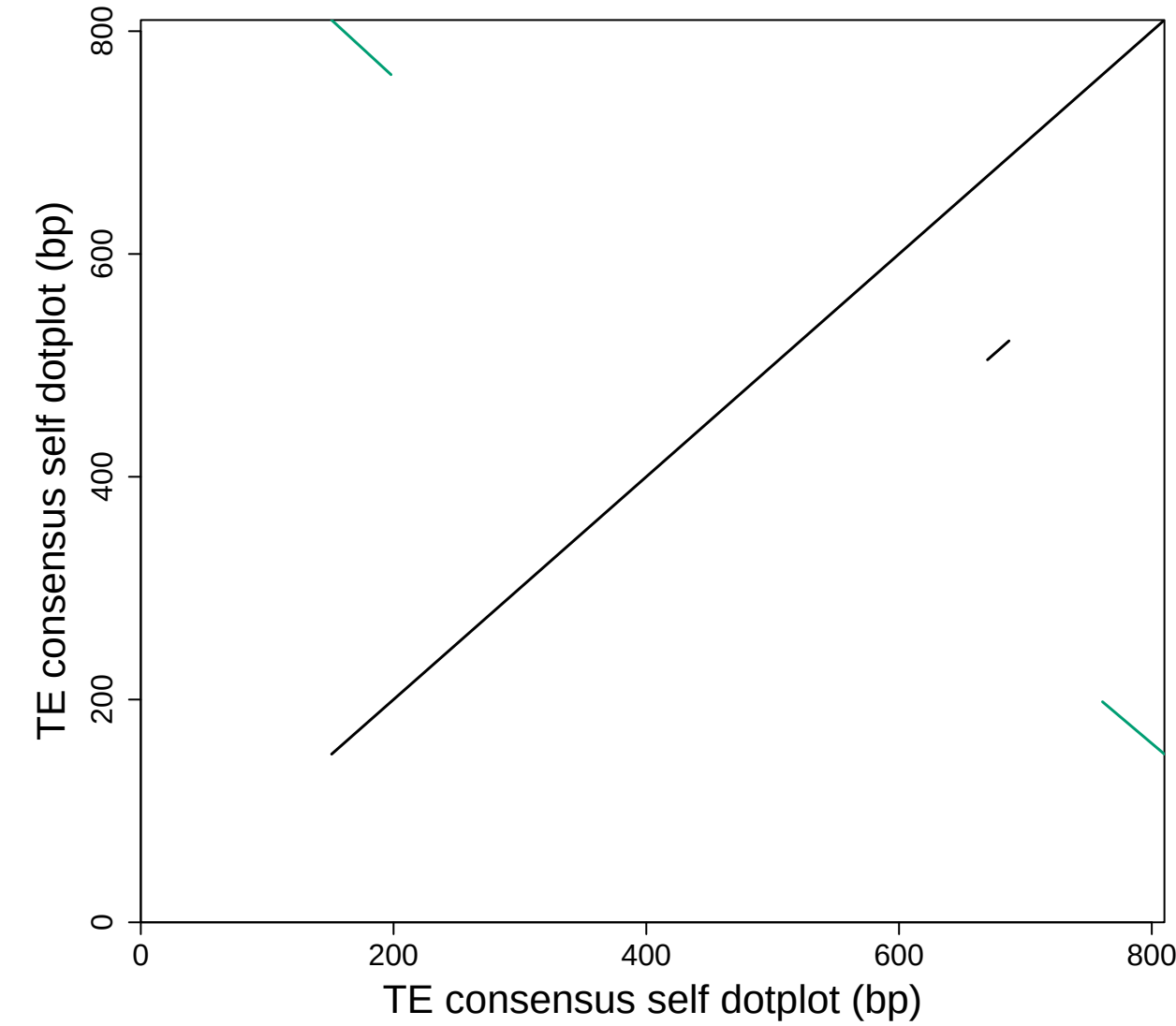
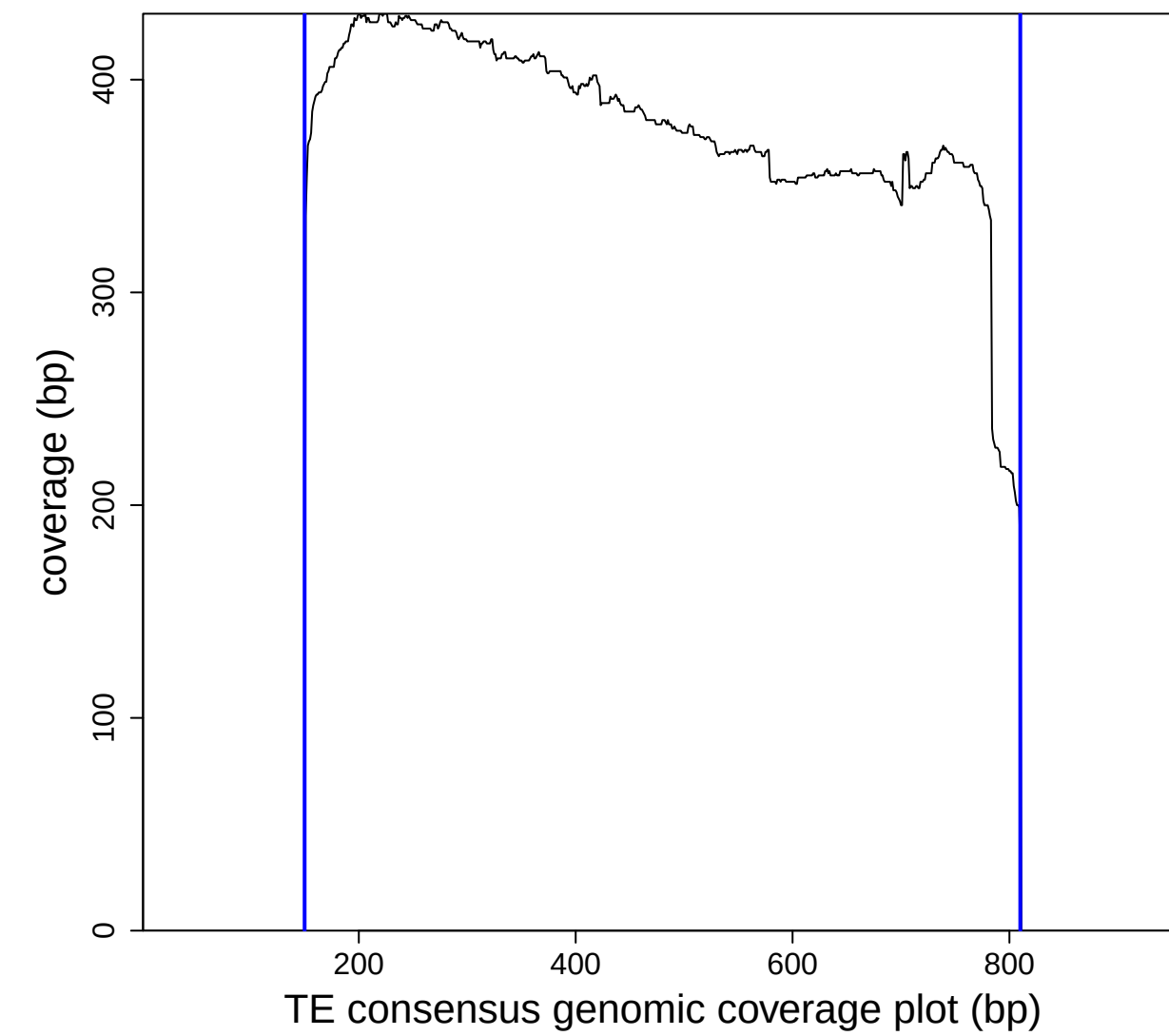
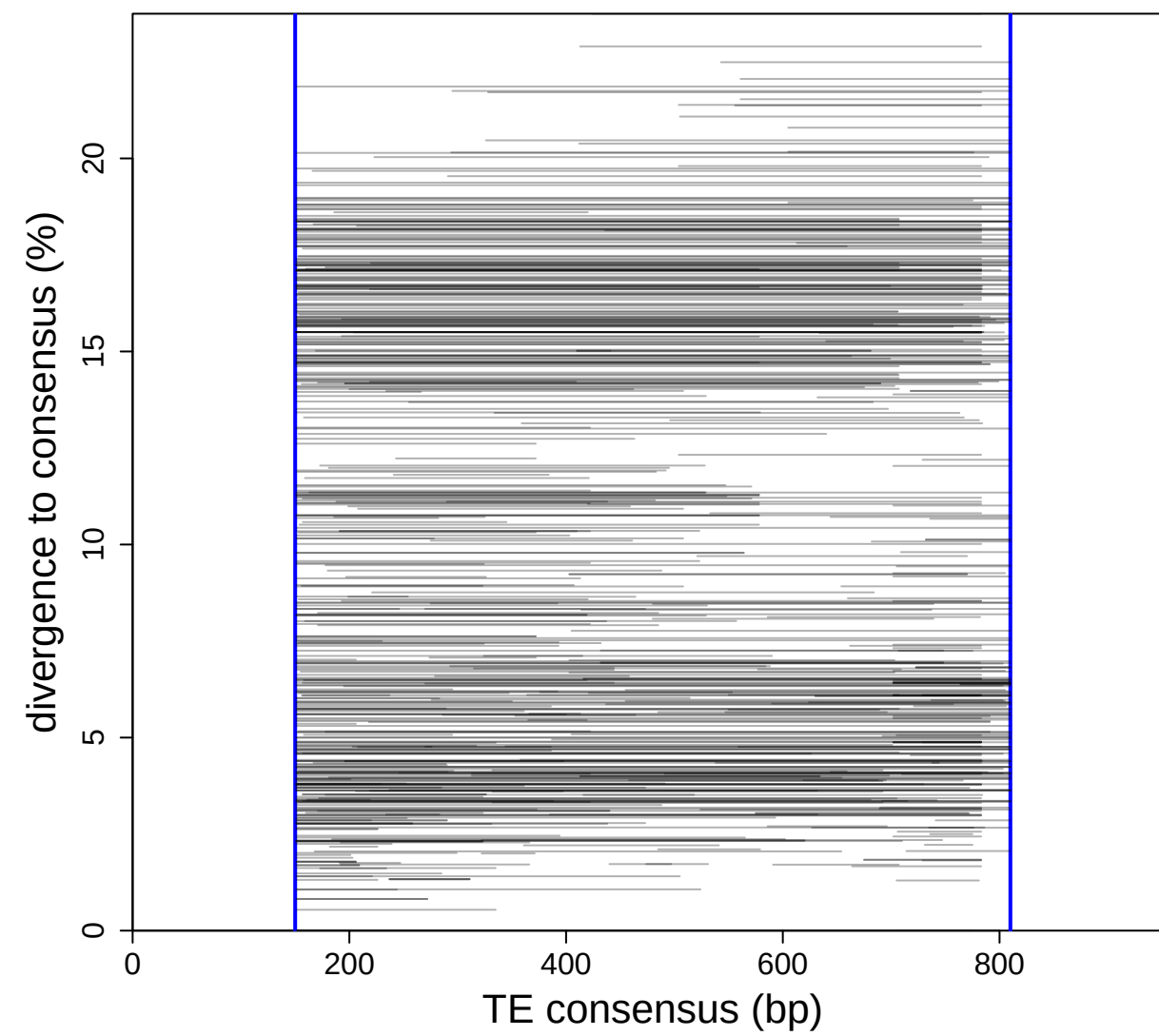
l.b.bed uf.bed g_3.bed fm_1.bed 0_0_bcln.fa aln.fa cl.fa gs.fa
size: 660bp; fragments: 653; full length: 215 (≥ 594 bp)

After TEtrimmer 660 bp

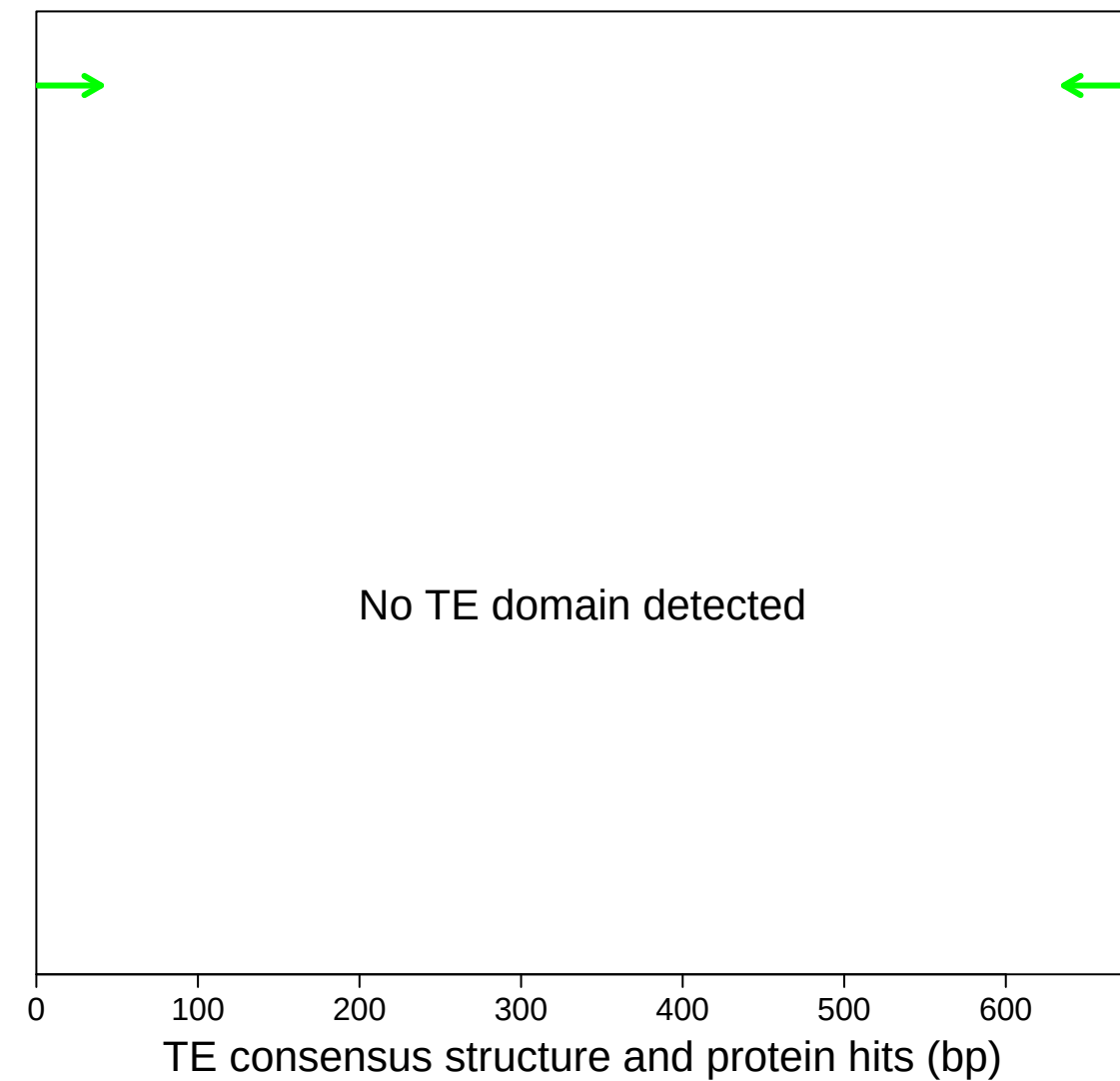
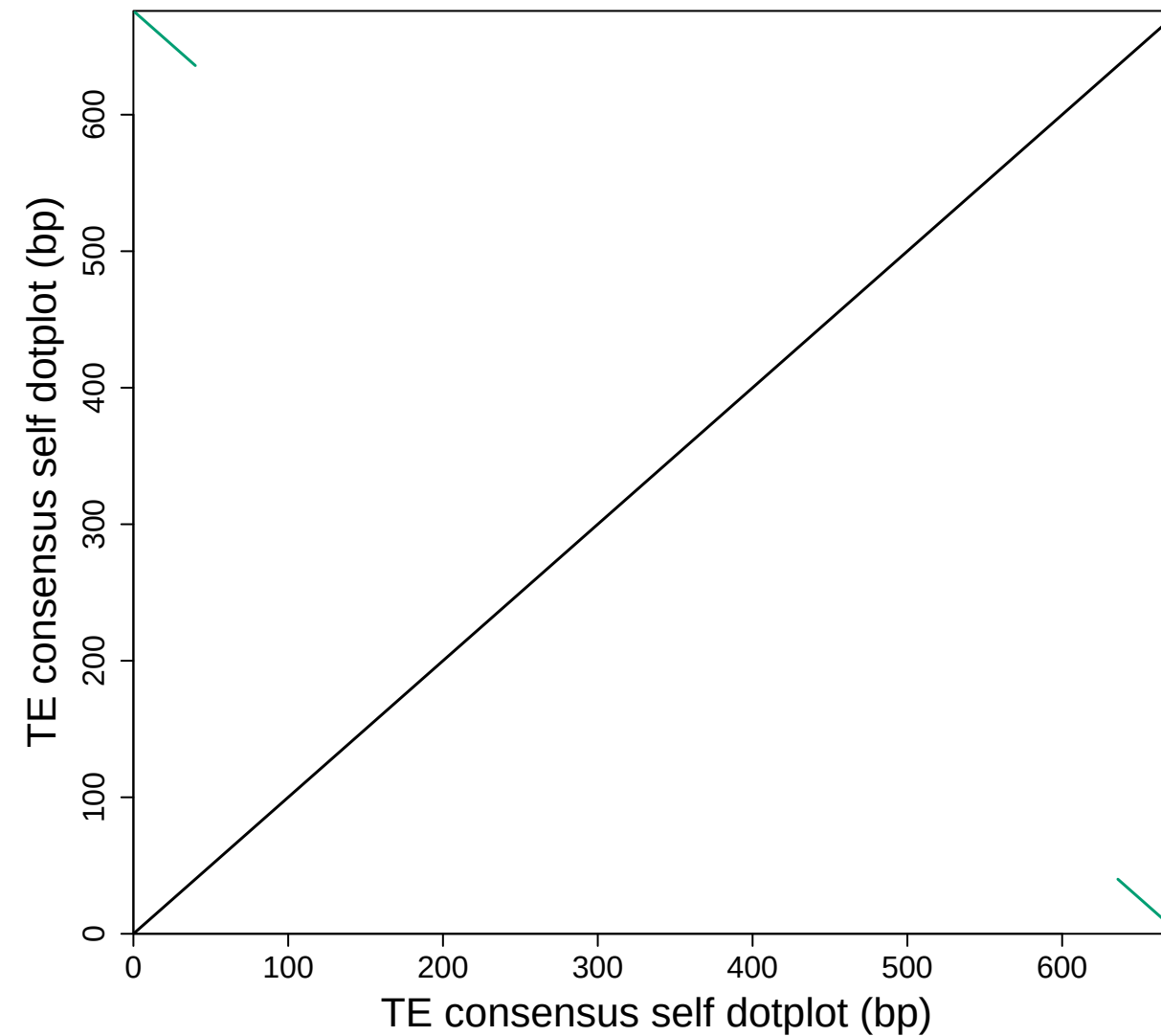
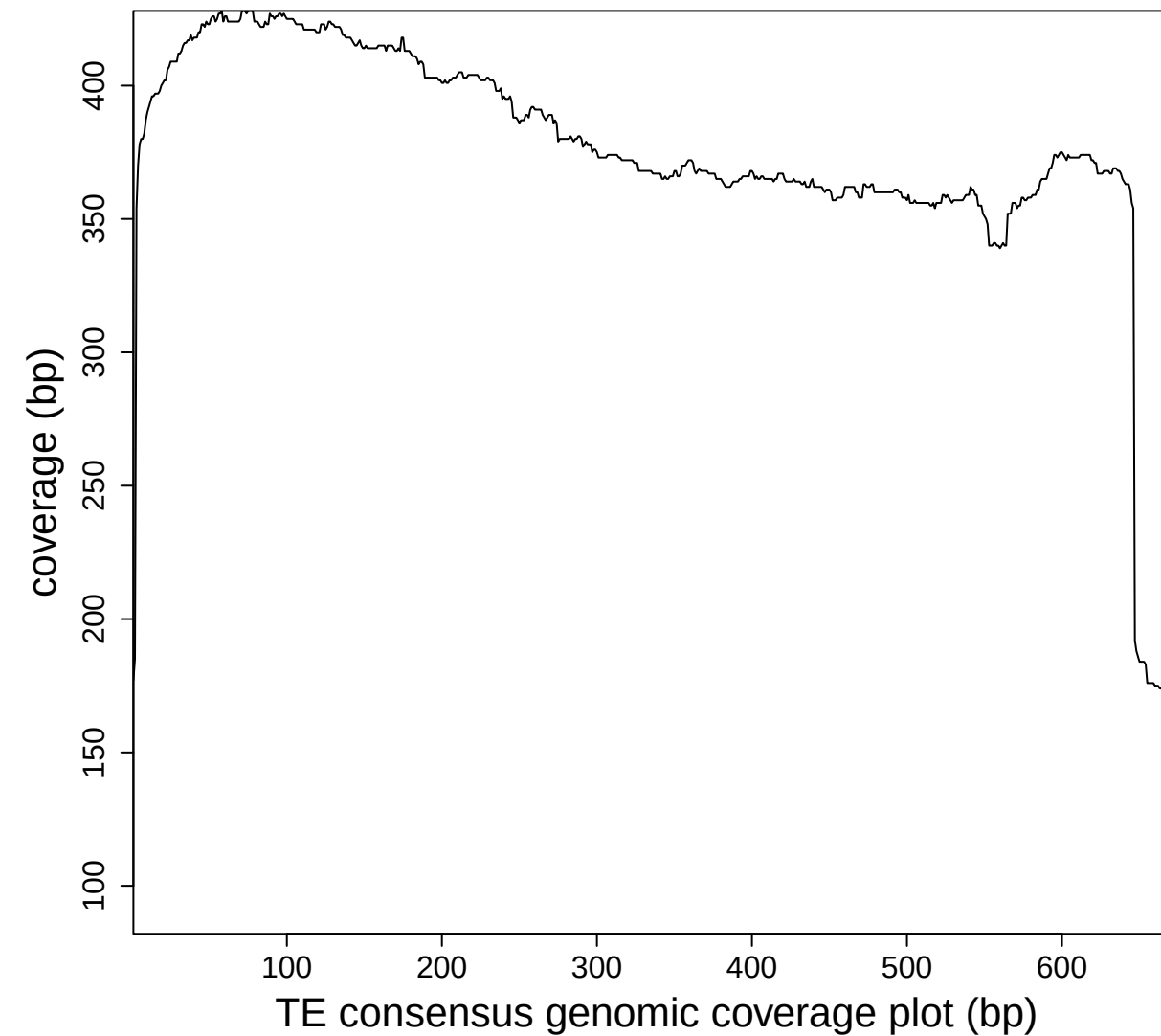
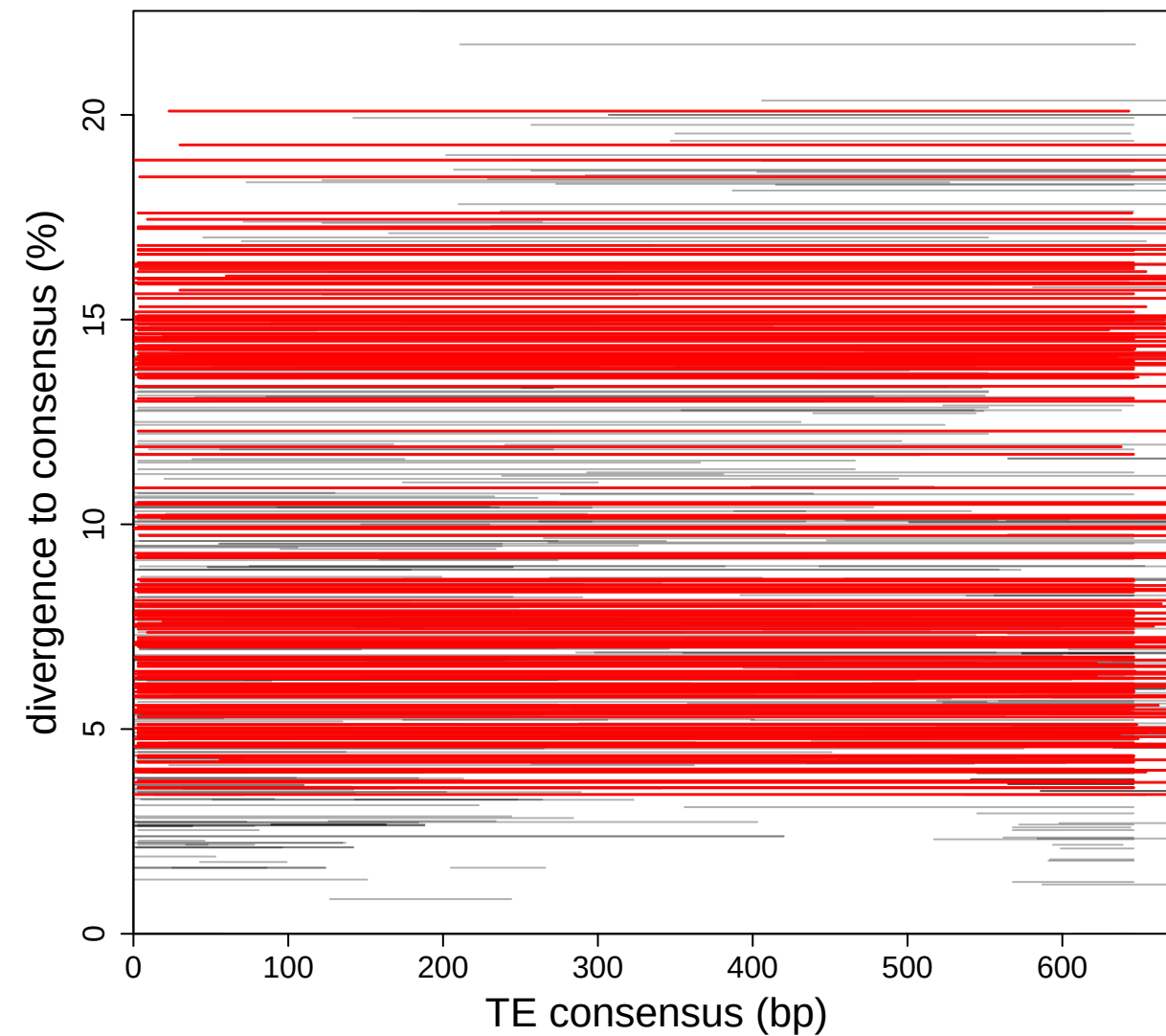


5.fasta.b.bed_uf.bed_g_3.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa
size: 960bp; fragments: 638; full length: 0 (>=864bp)

After TEtrimmer Extended plot Blue lines are boundaries

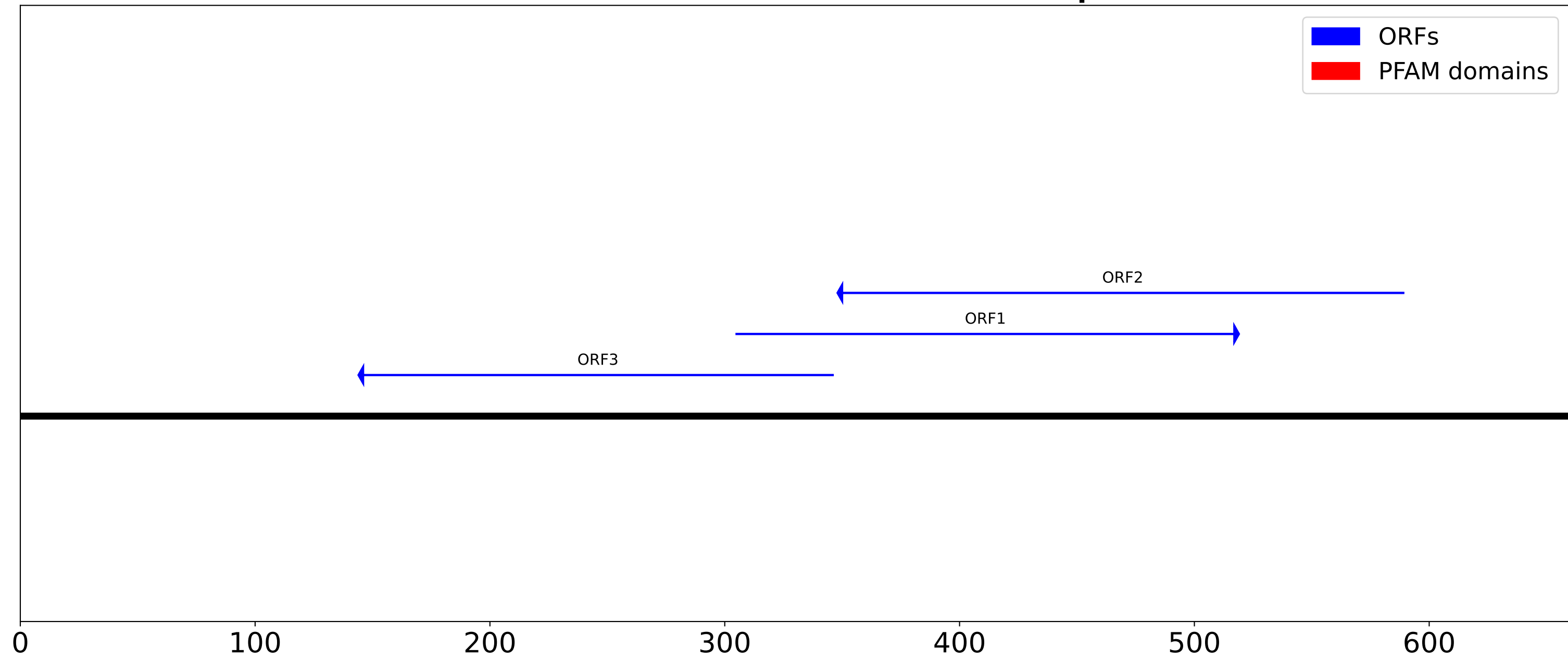


TE: rnd_4_family_205
size: 676bp; fragments: 643; full length: 214 (≥ 608.4 bp)

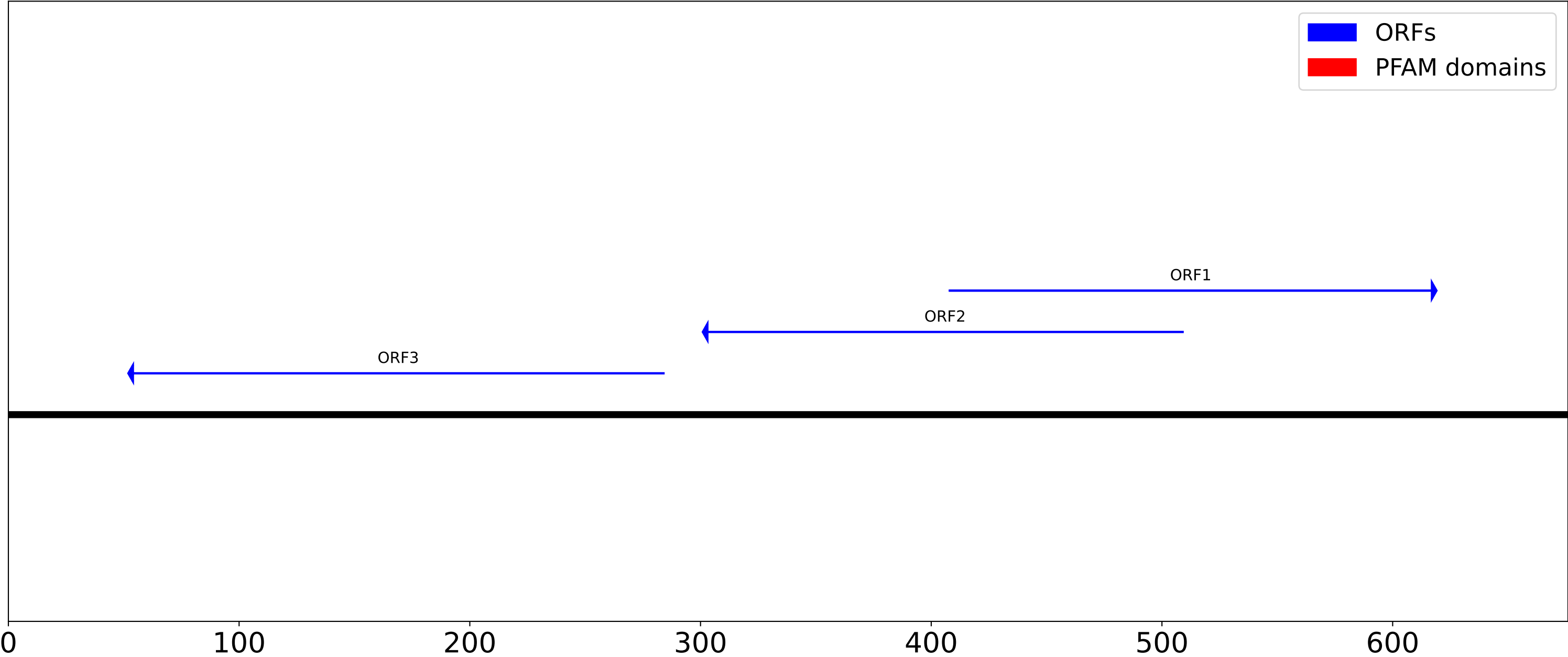


Before TEtrimmer 676 bp

After TEtrimmer ORF and PFAM domain plot



Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

