

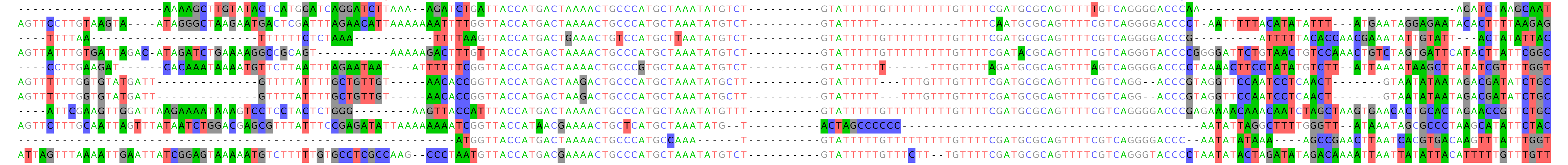
Start crop Point



End crop Point

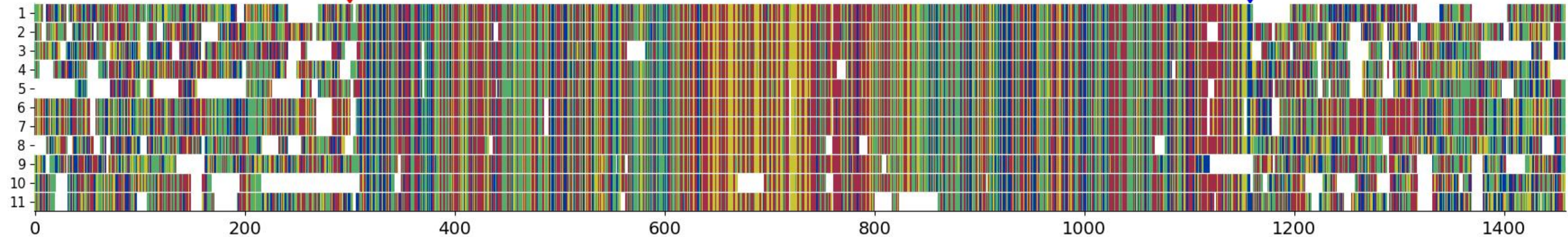


MSA length = 859



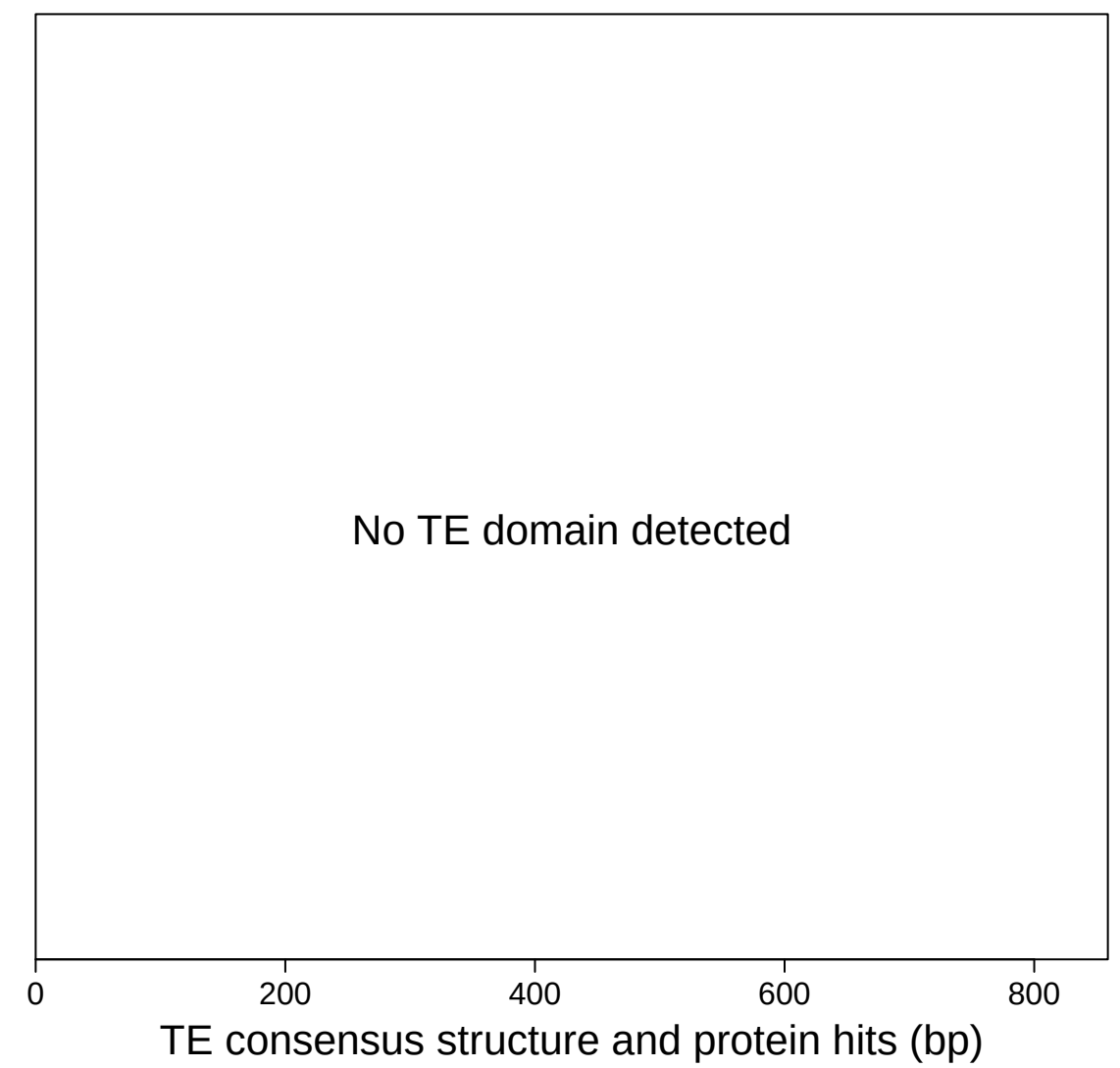
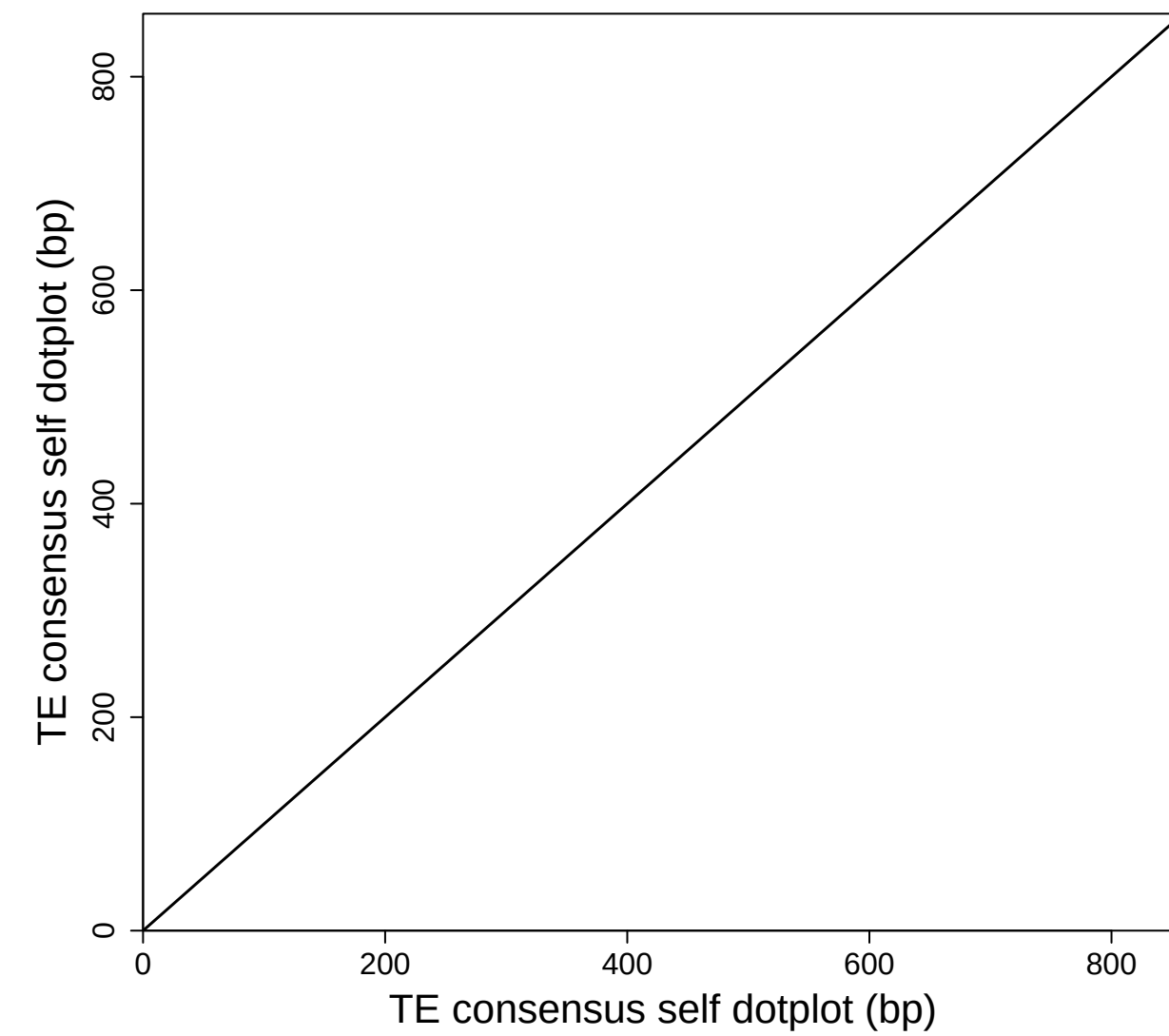
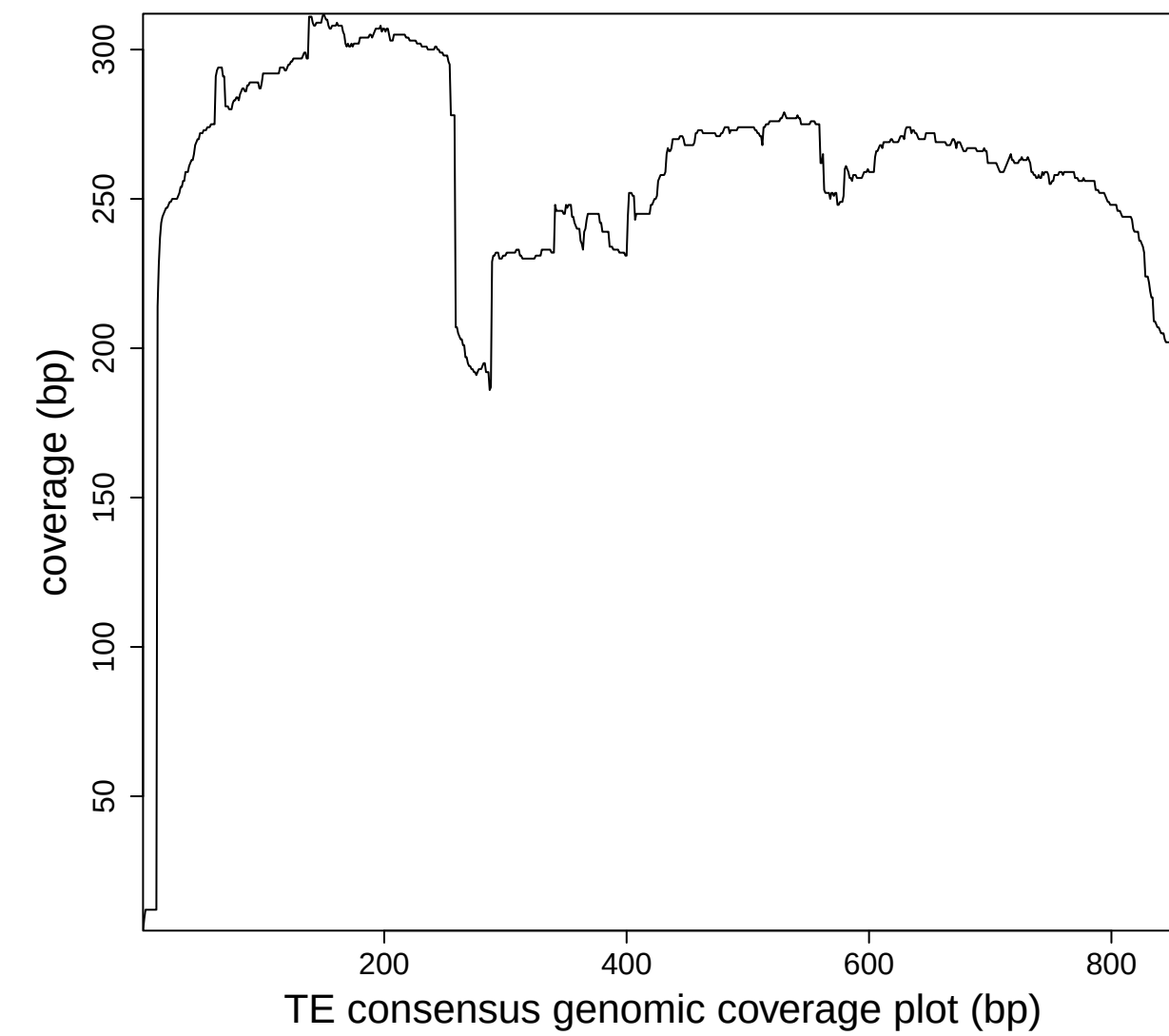
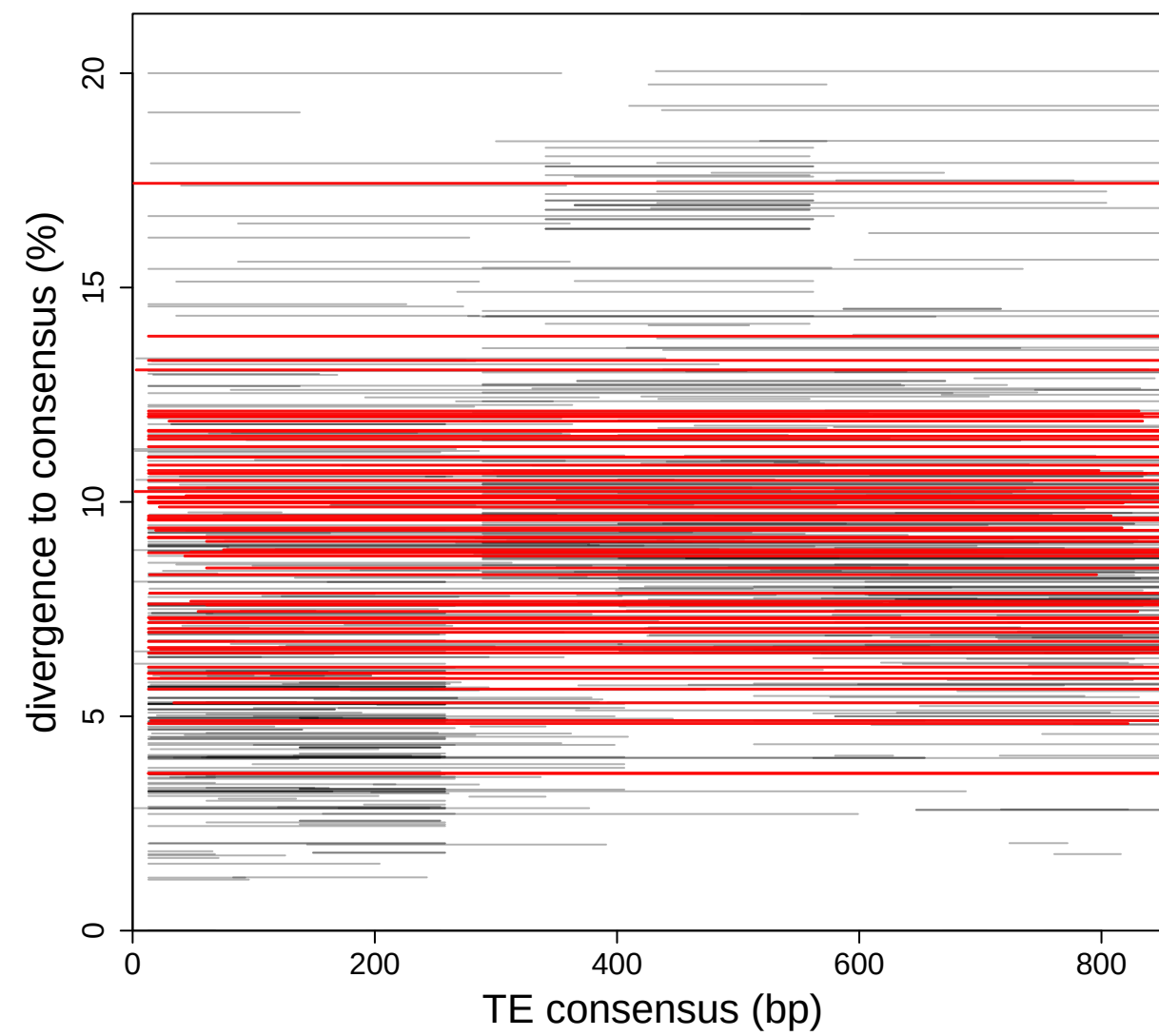
Start crop Point

End crop Point



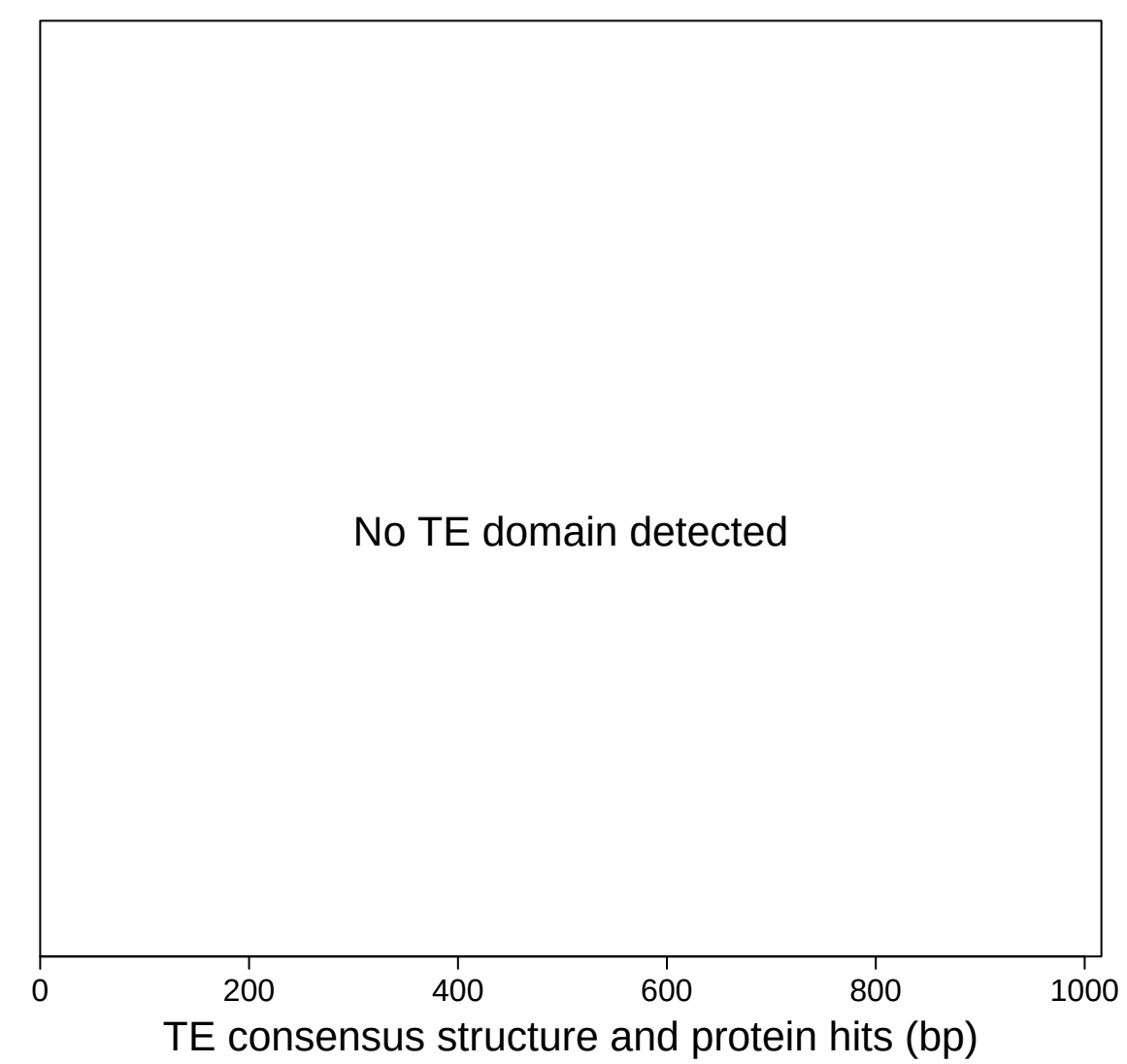
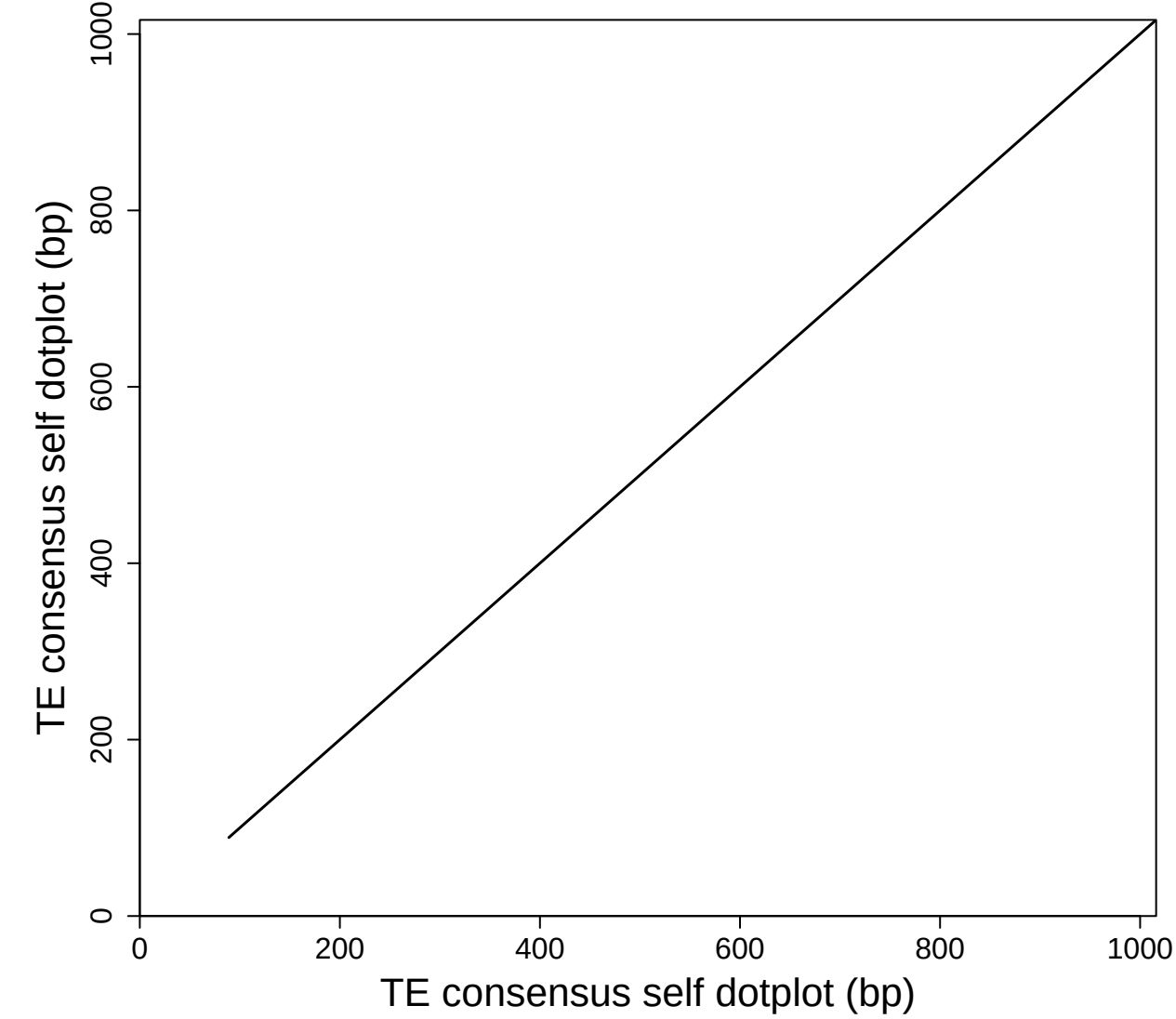
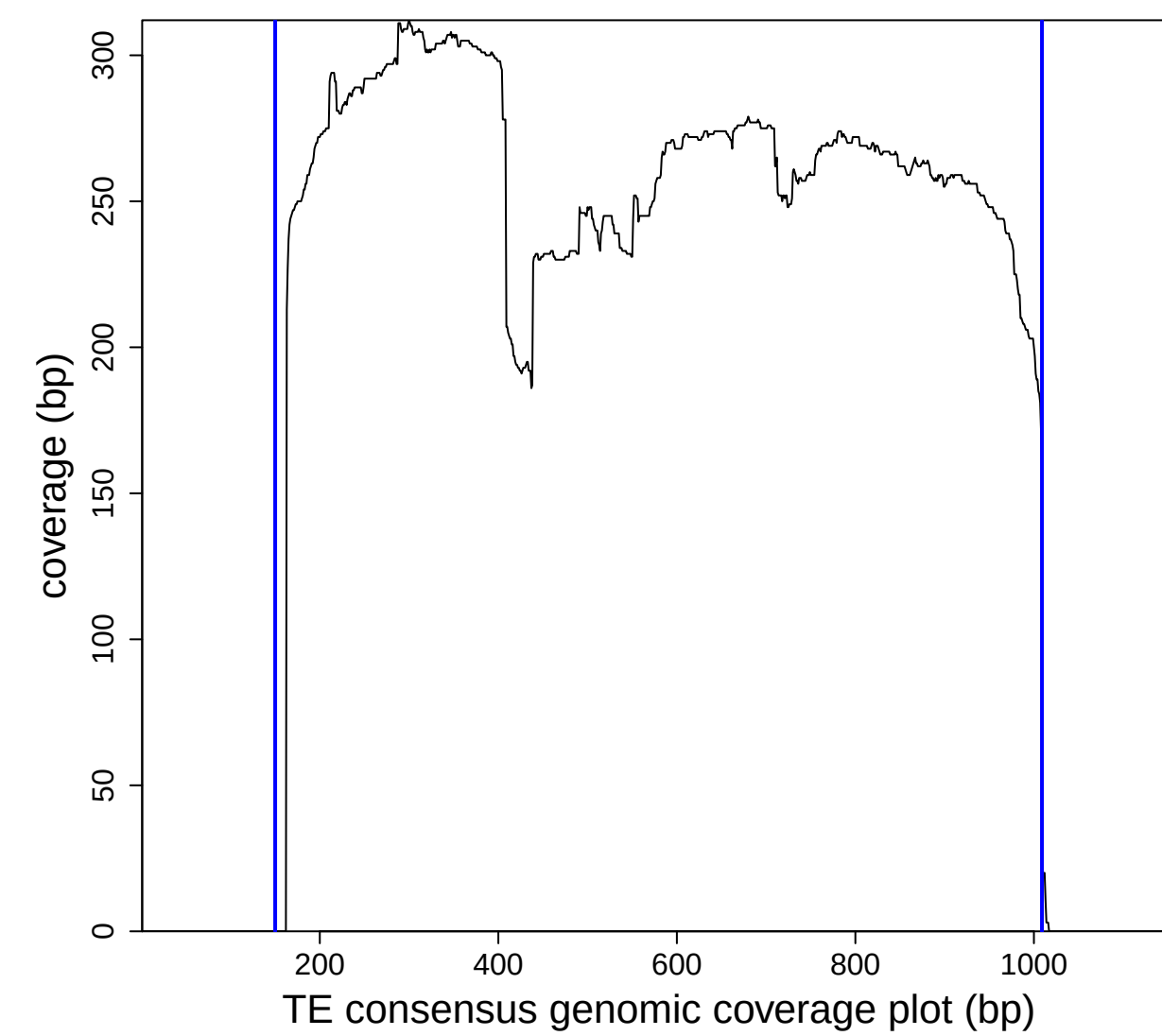
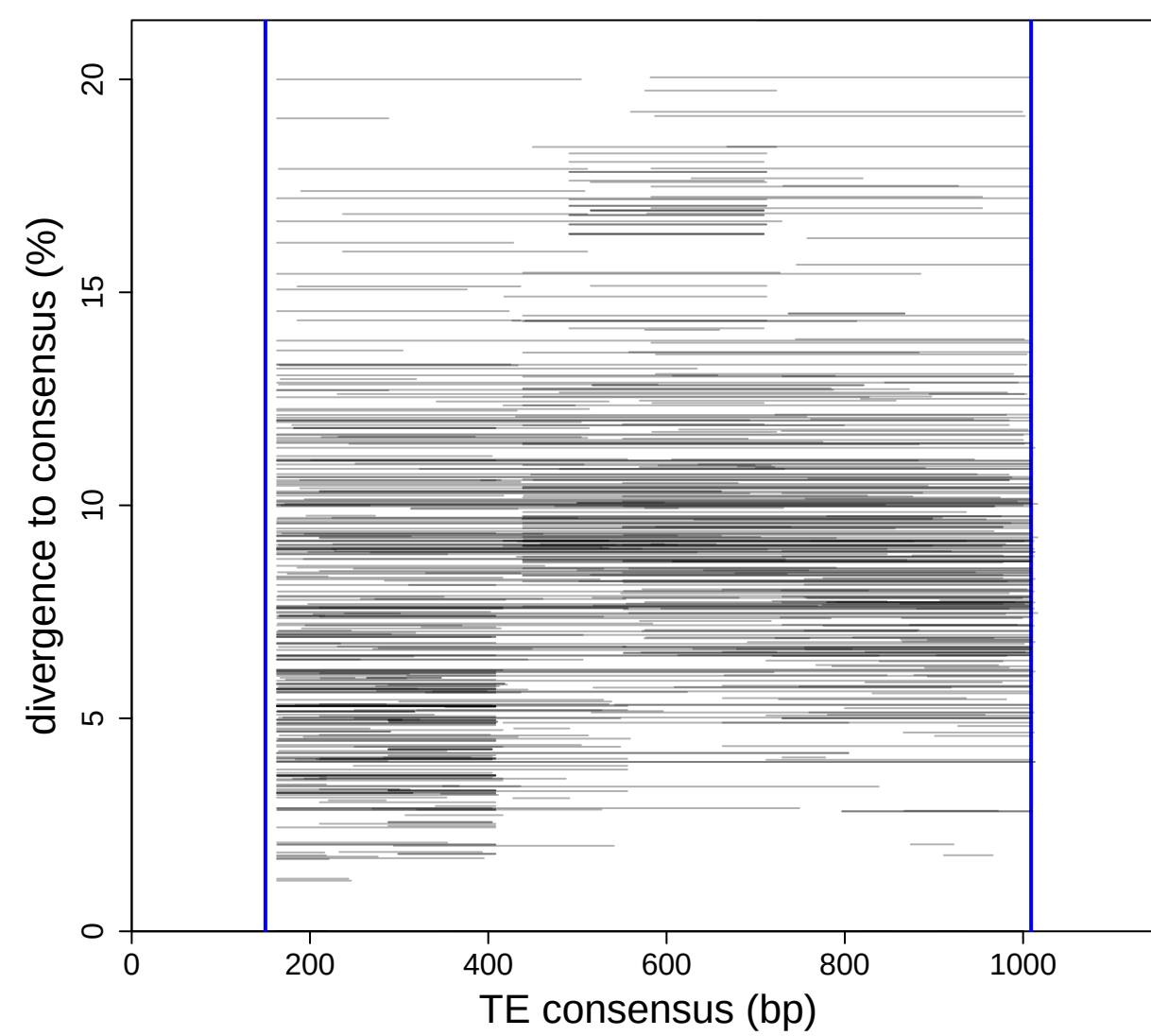
l.b.bed uf.bed g_3.bed fm_1.bed 0 0 bcIn.fa aln.fa cl.fa gs.fa
size: 859bp; fragments: 632; full length: 68 (≥ 773.1 bp)

After TEtrimmer 859 bp

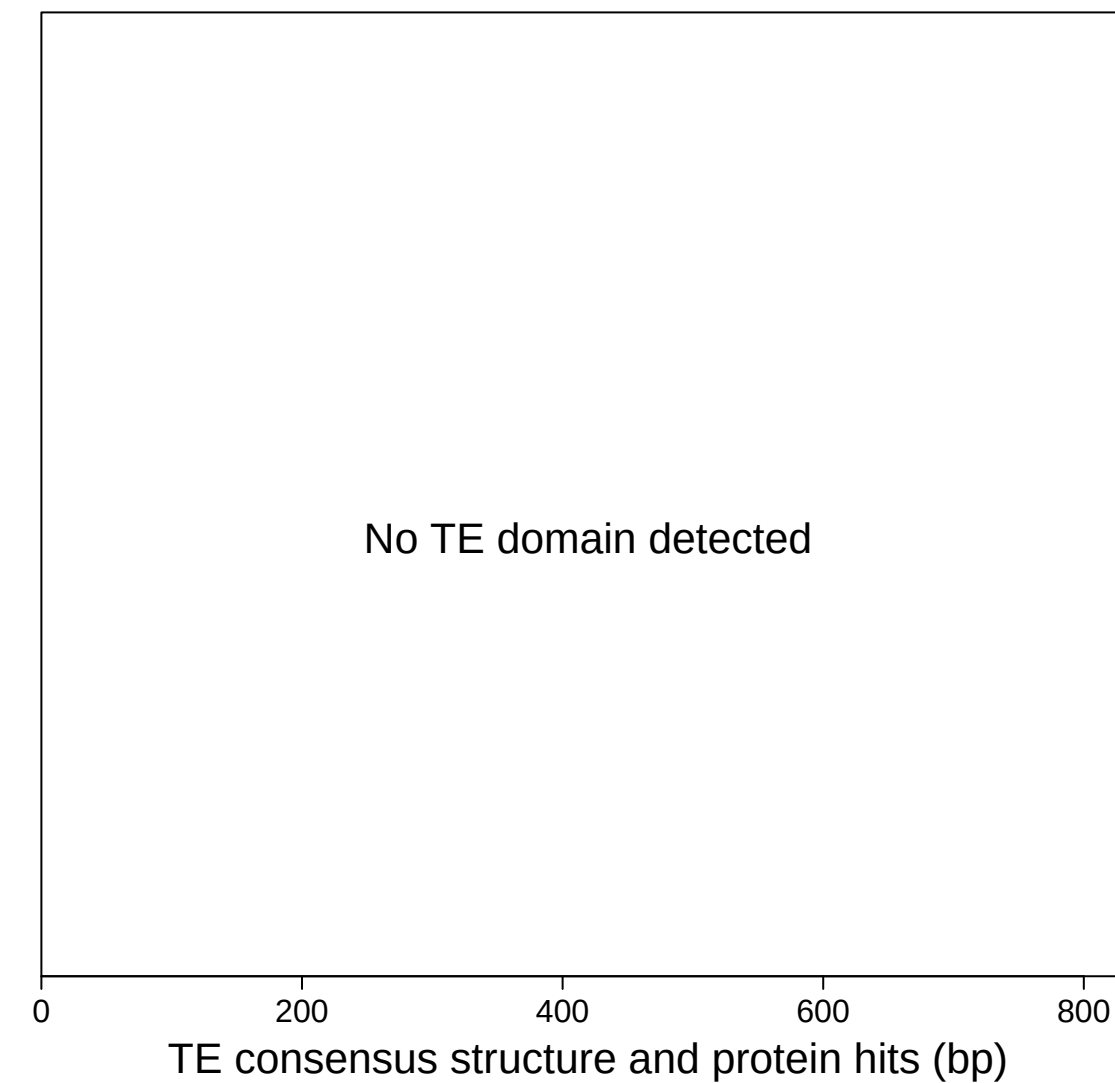
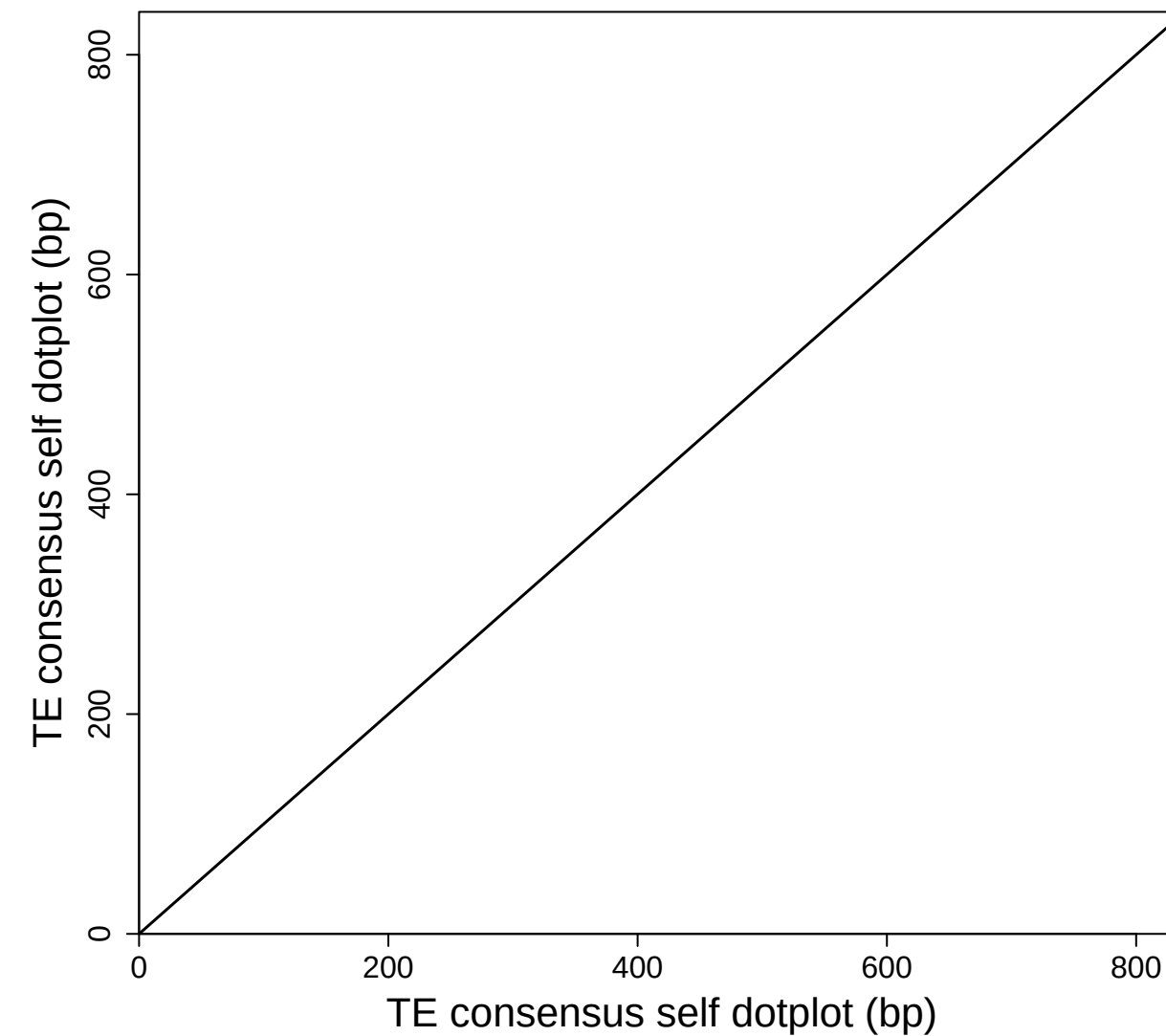
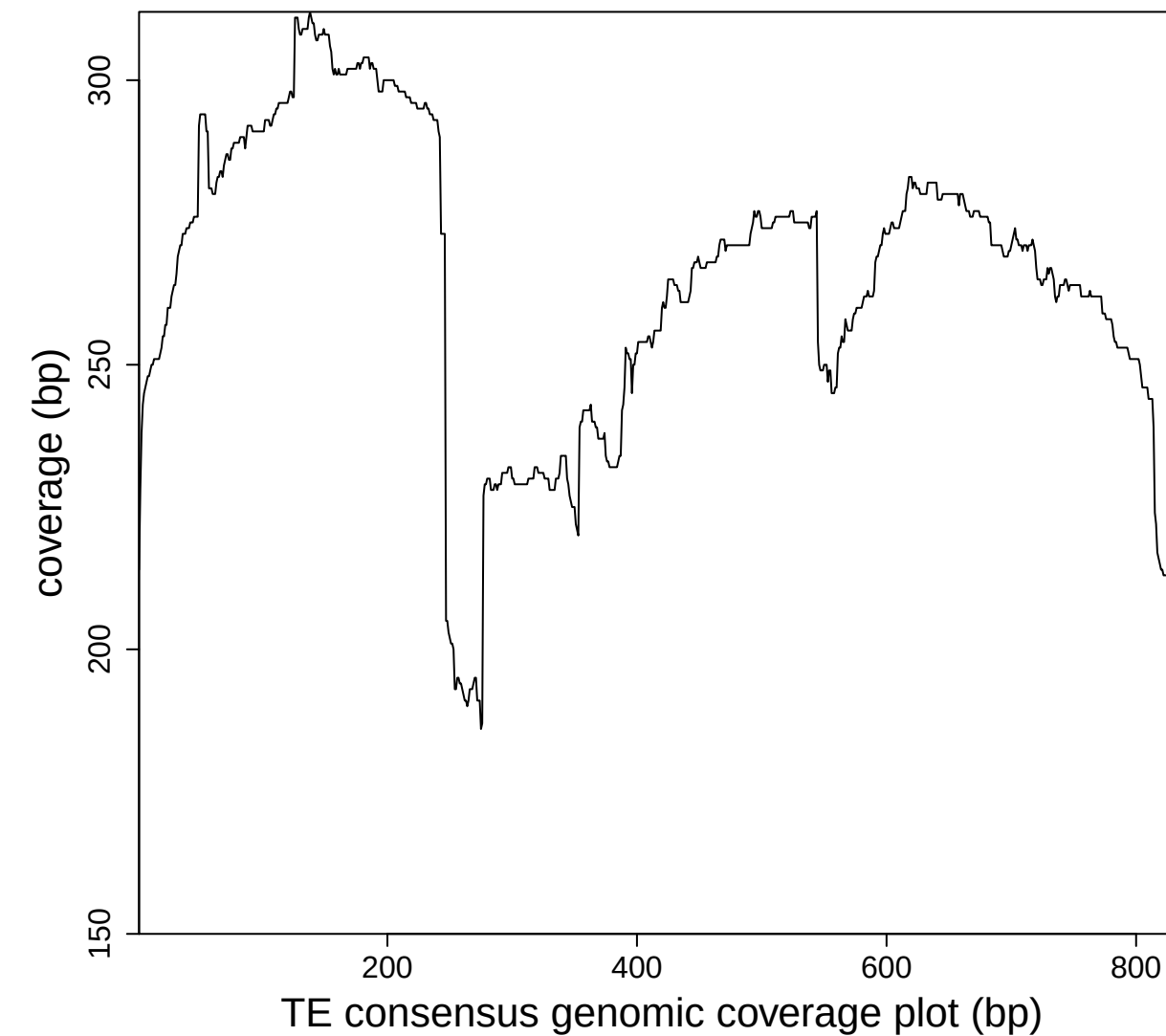
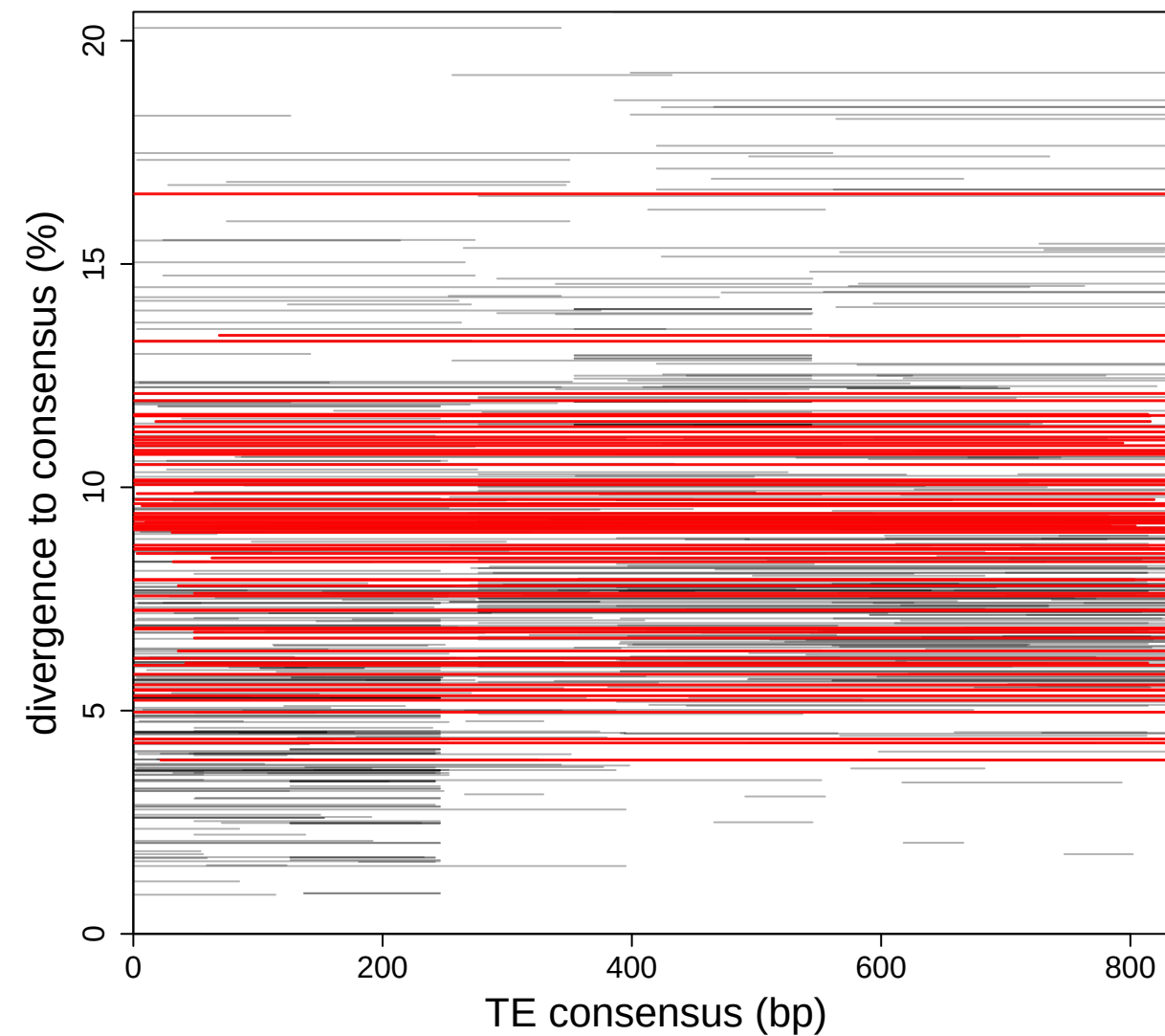


9.fasta.b.bed uf.bed g_3.bed fm_1.bed_0_0 bcln.fa aln.fa cl.fa
size: 1159bp; fragments: 632; full length: 0 (>=1043.1bp)

After TEtrimmer Extended plot Blue lines are boundaries

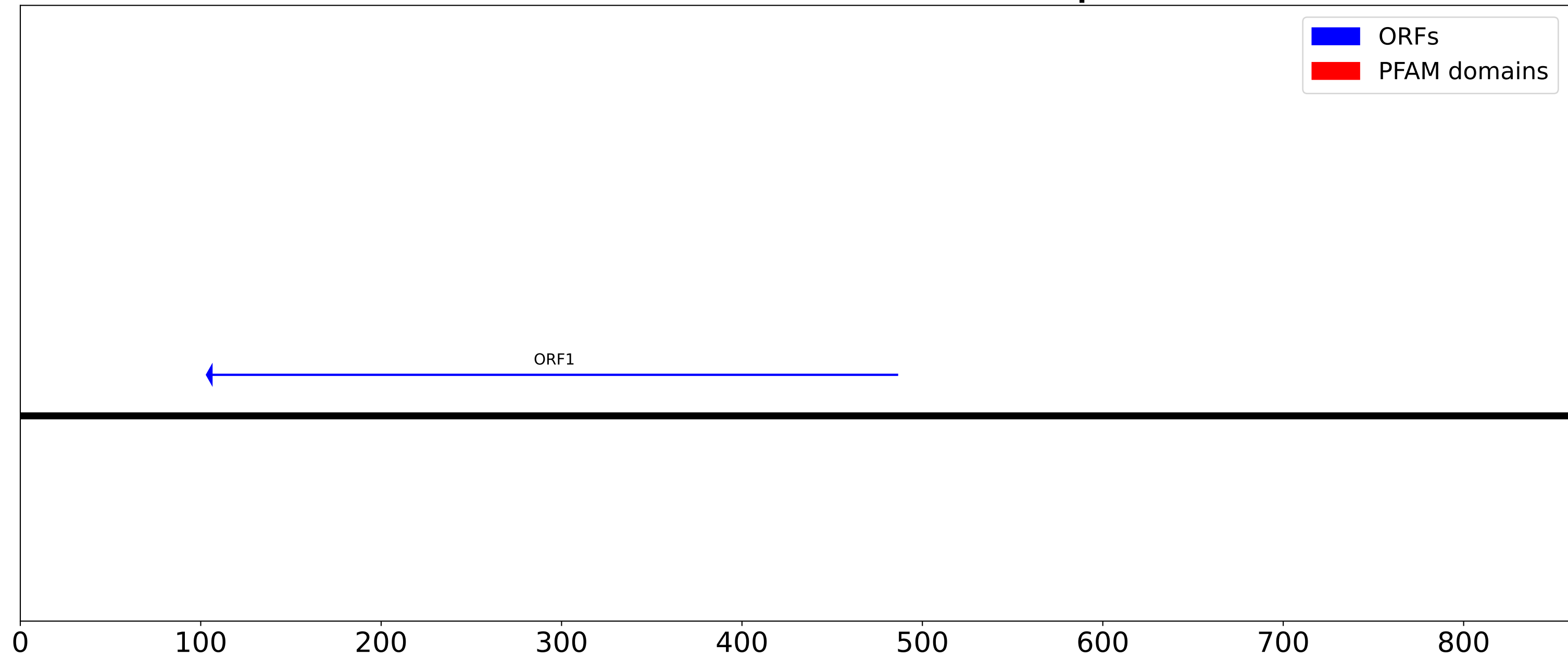


TE: rnd_4_family_619
size: 839bp; fragments: 628; full length: 70 (≥ 755.1 bp)

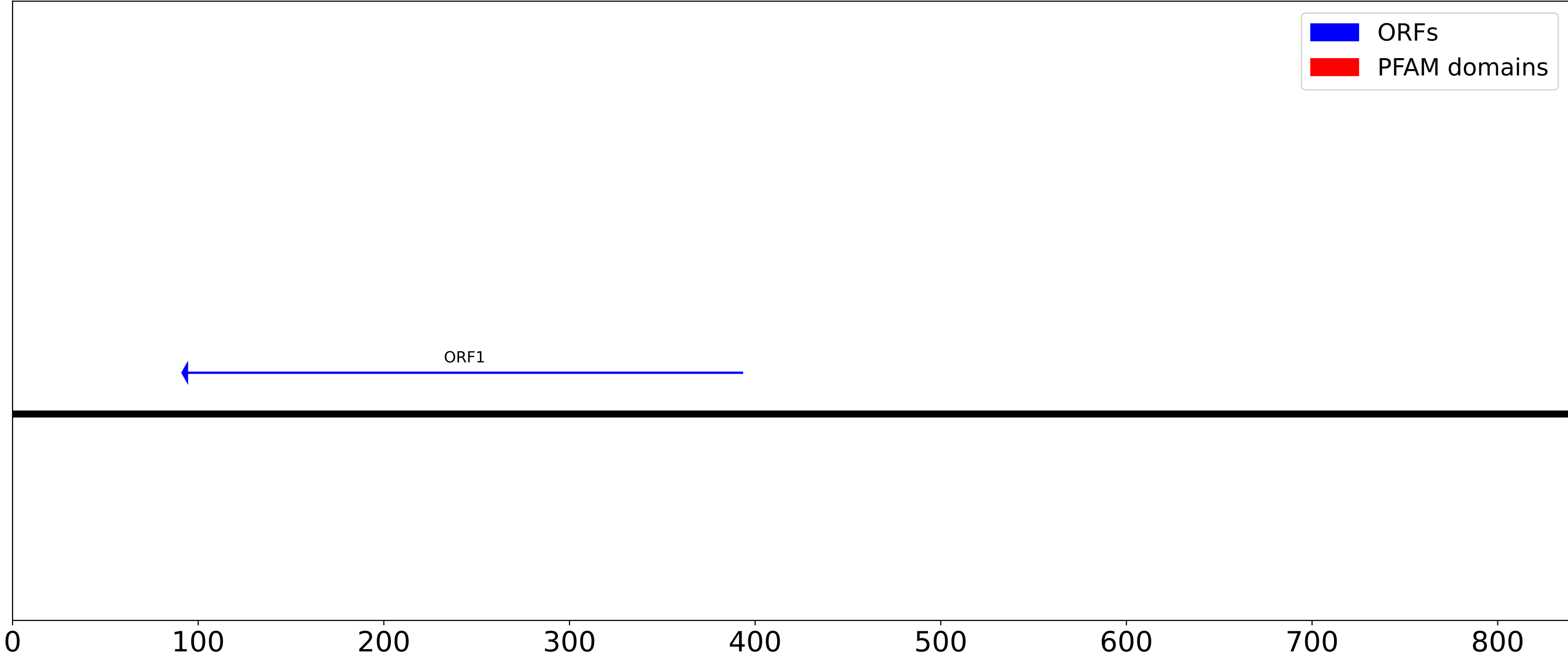


Before TEtrimmer 839 bp

After TEtrimmer ORF and PFAM domain plot



Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

