

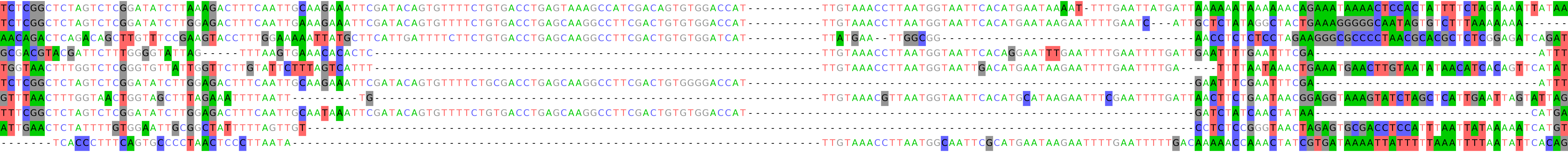
Start Crop Point



End Crop Point



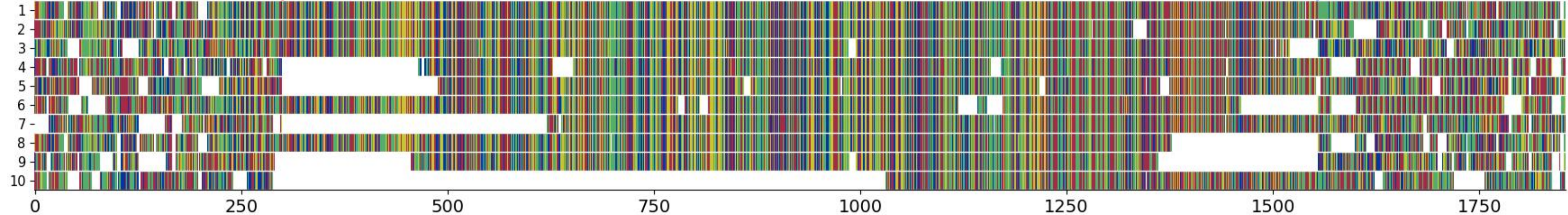
MSA length = 1255



Start crop Point

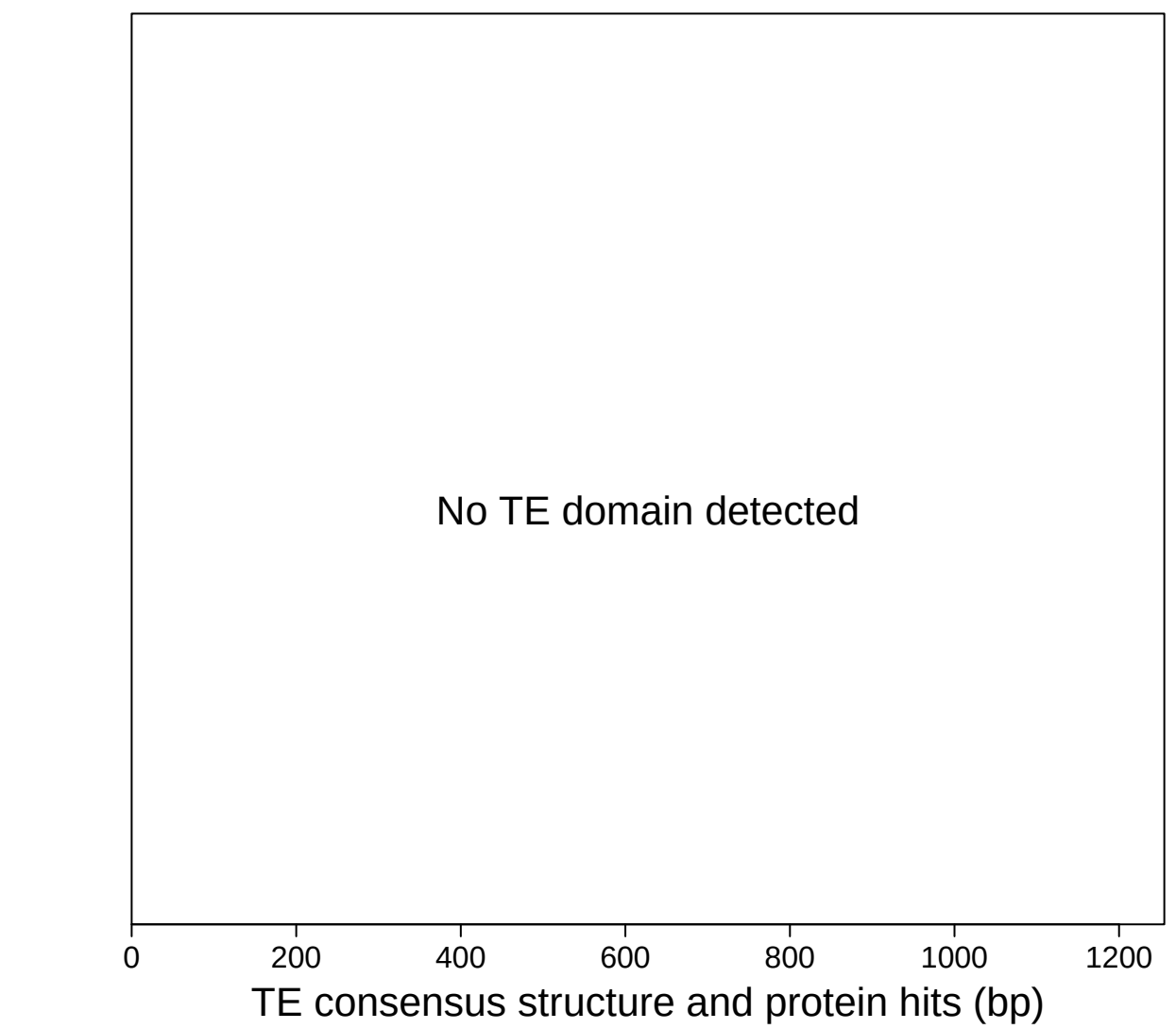
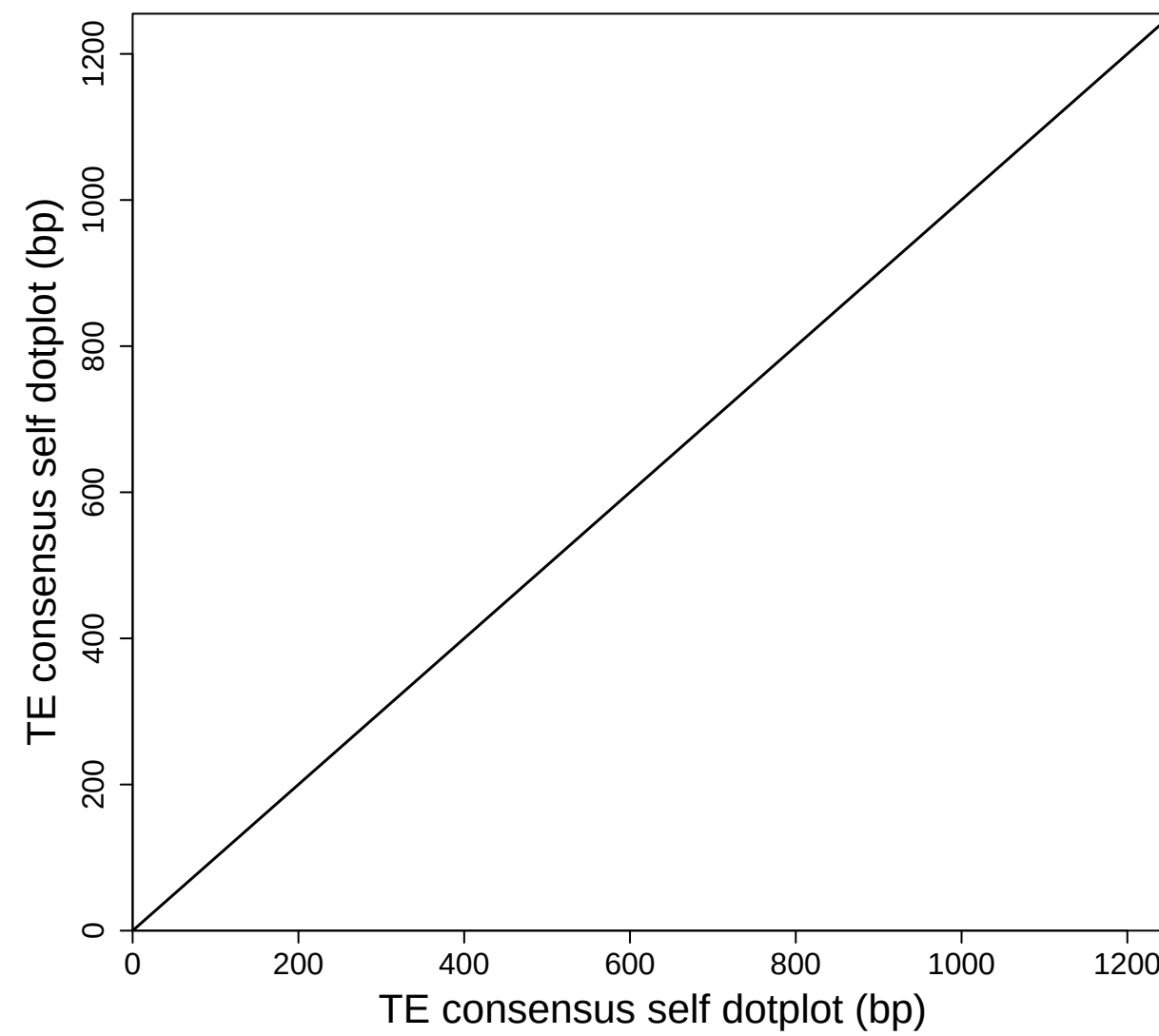
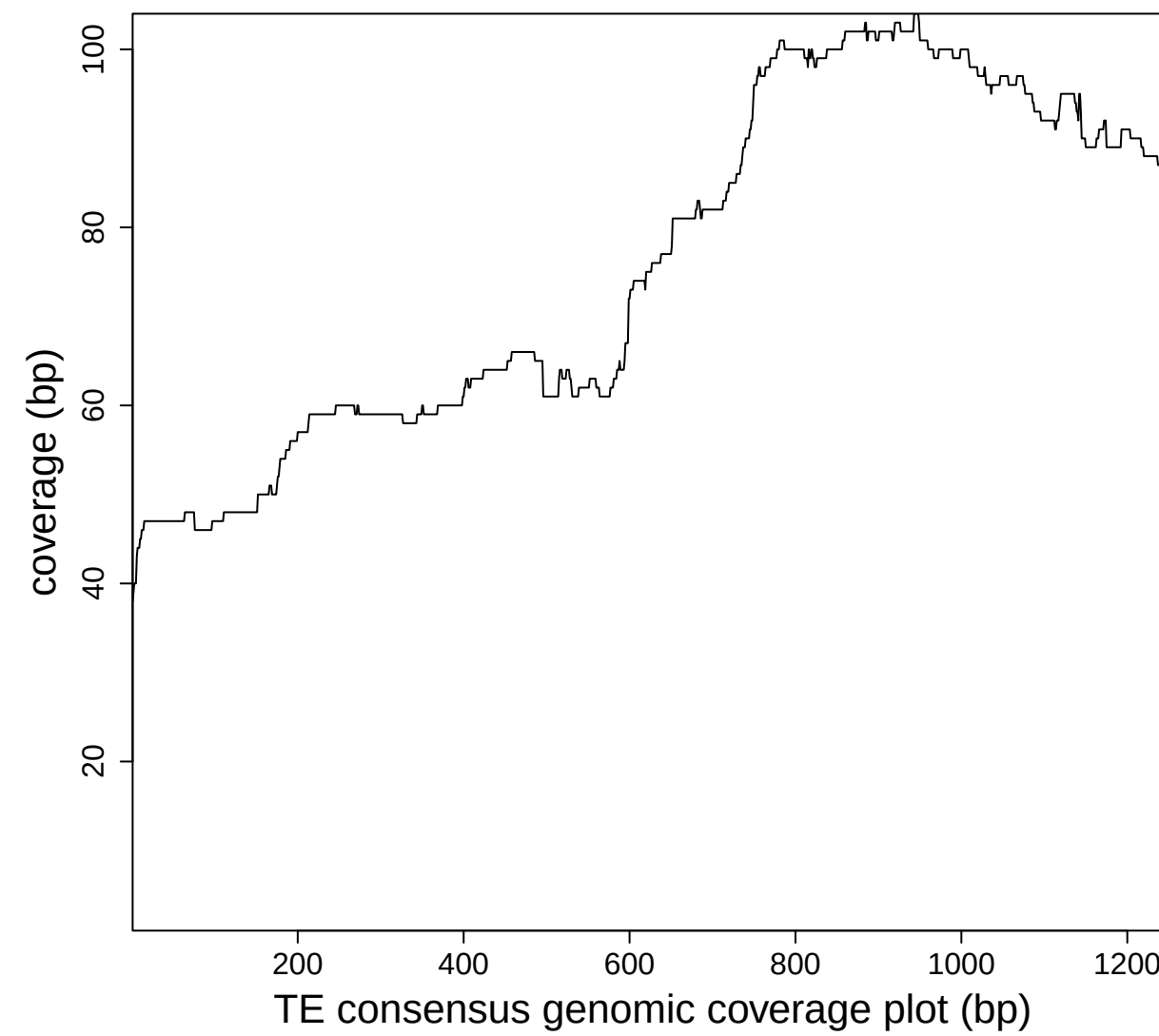
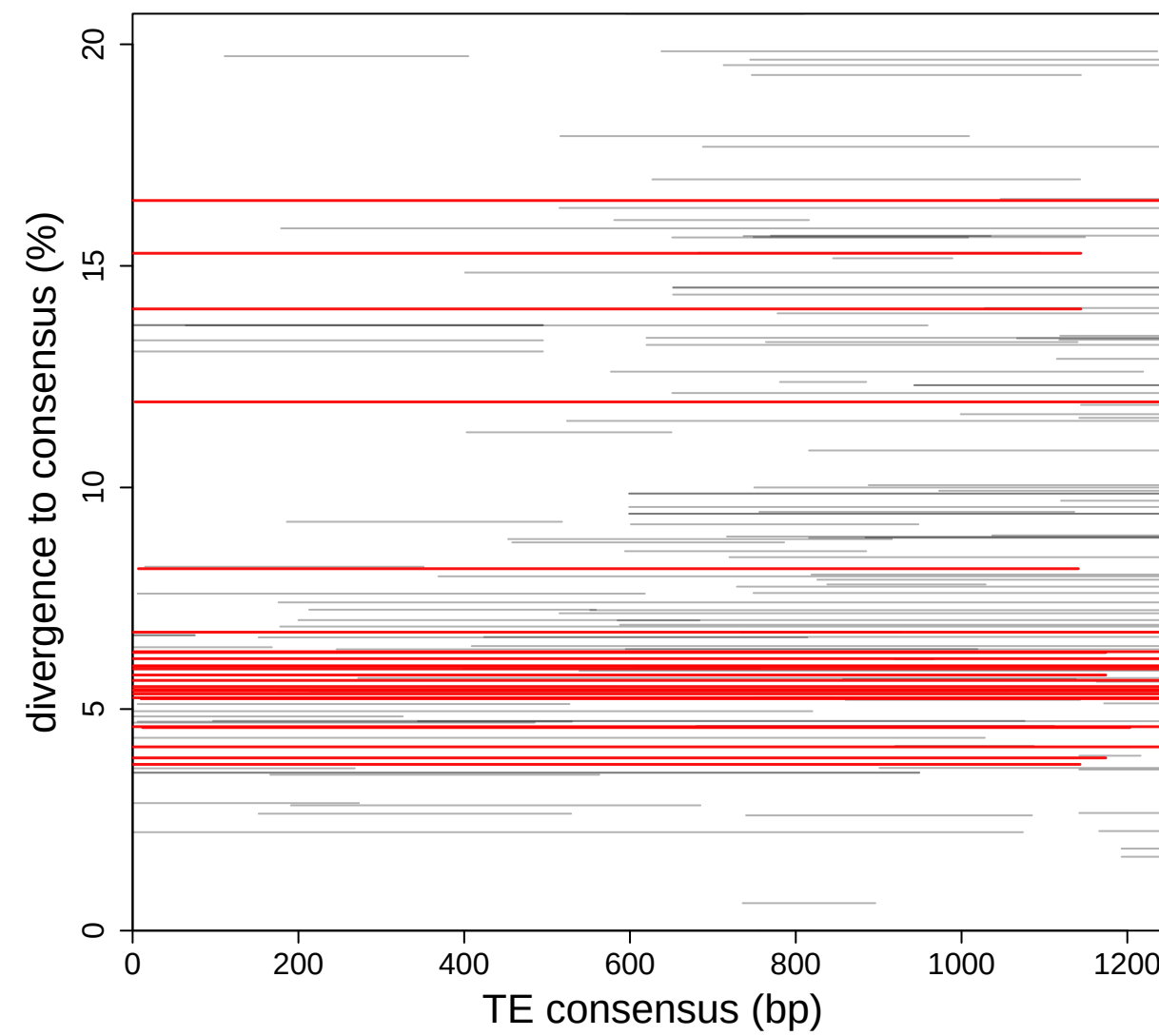


End crop Point



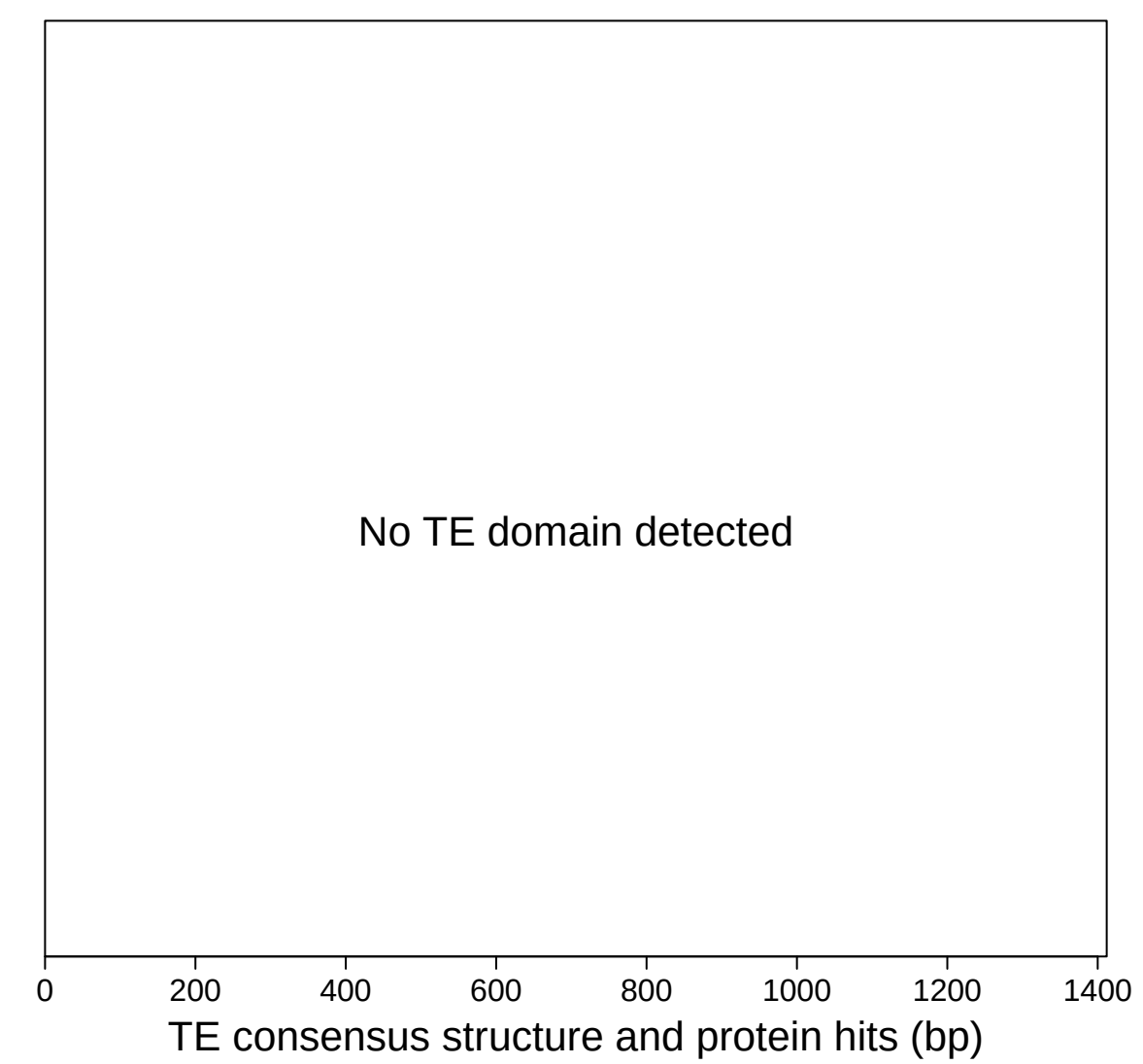
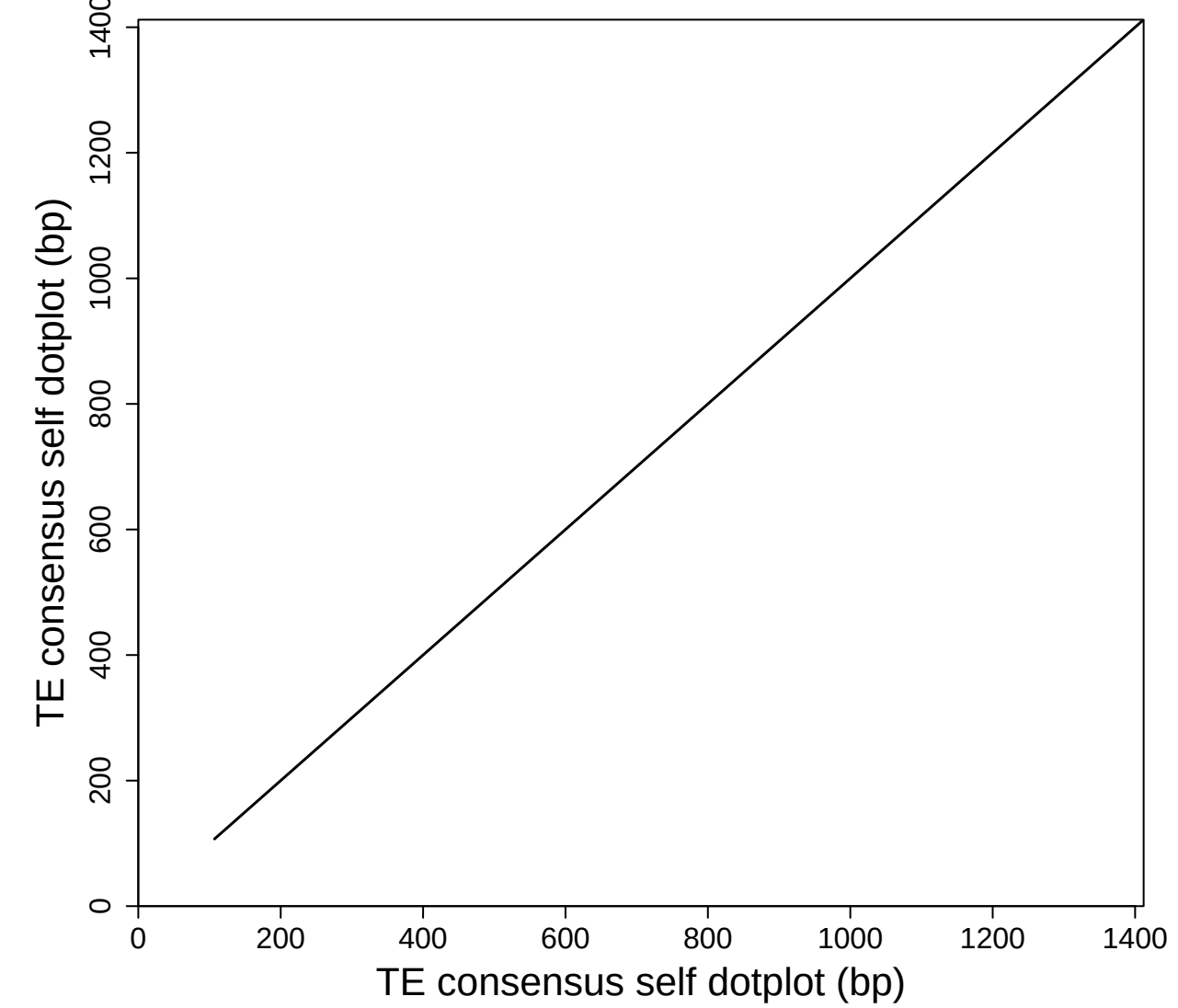
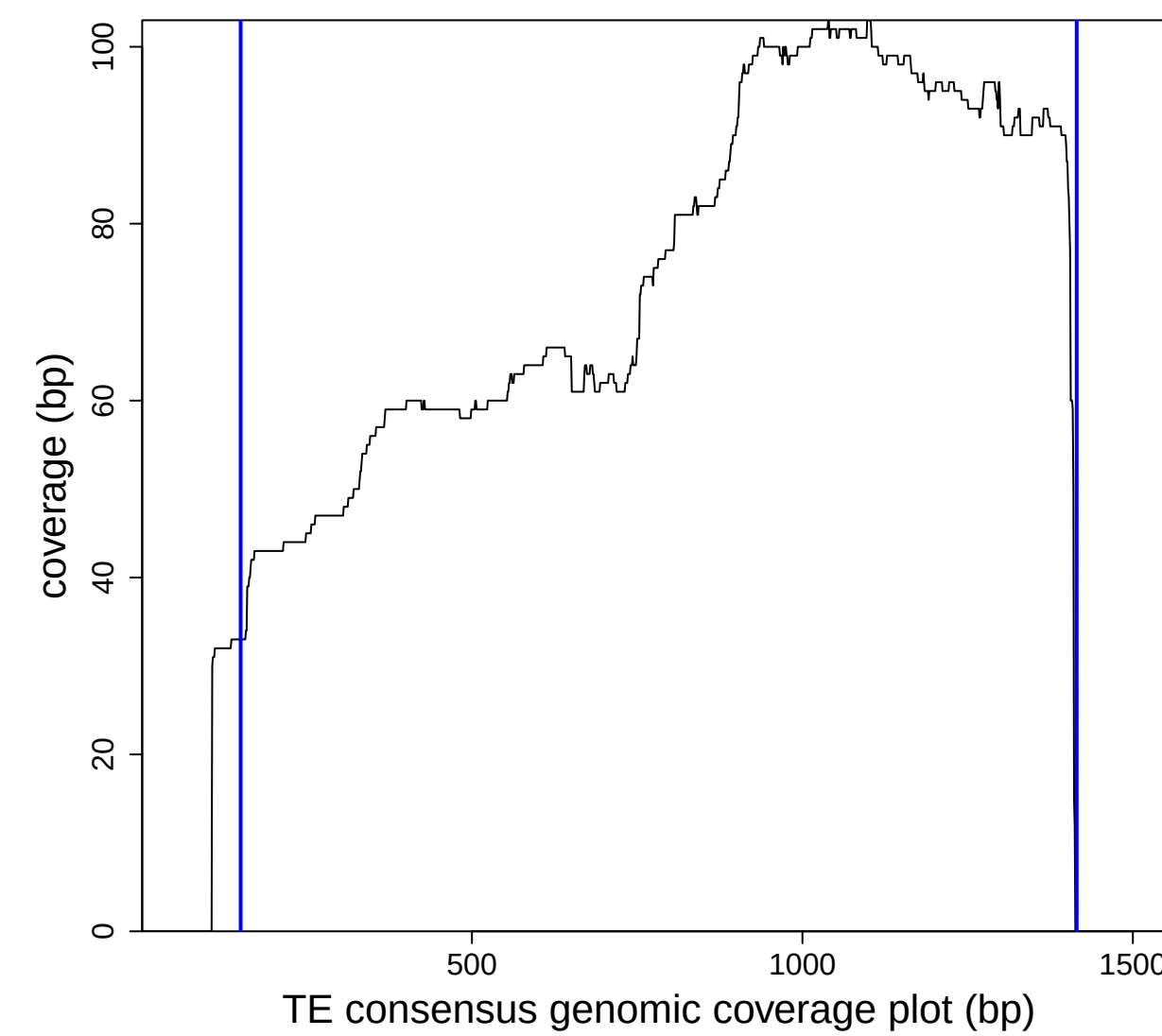
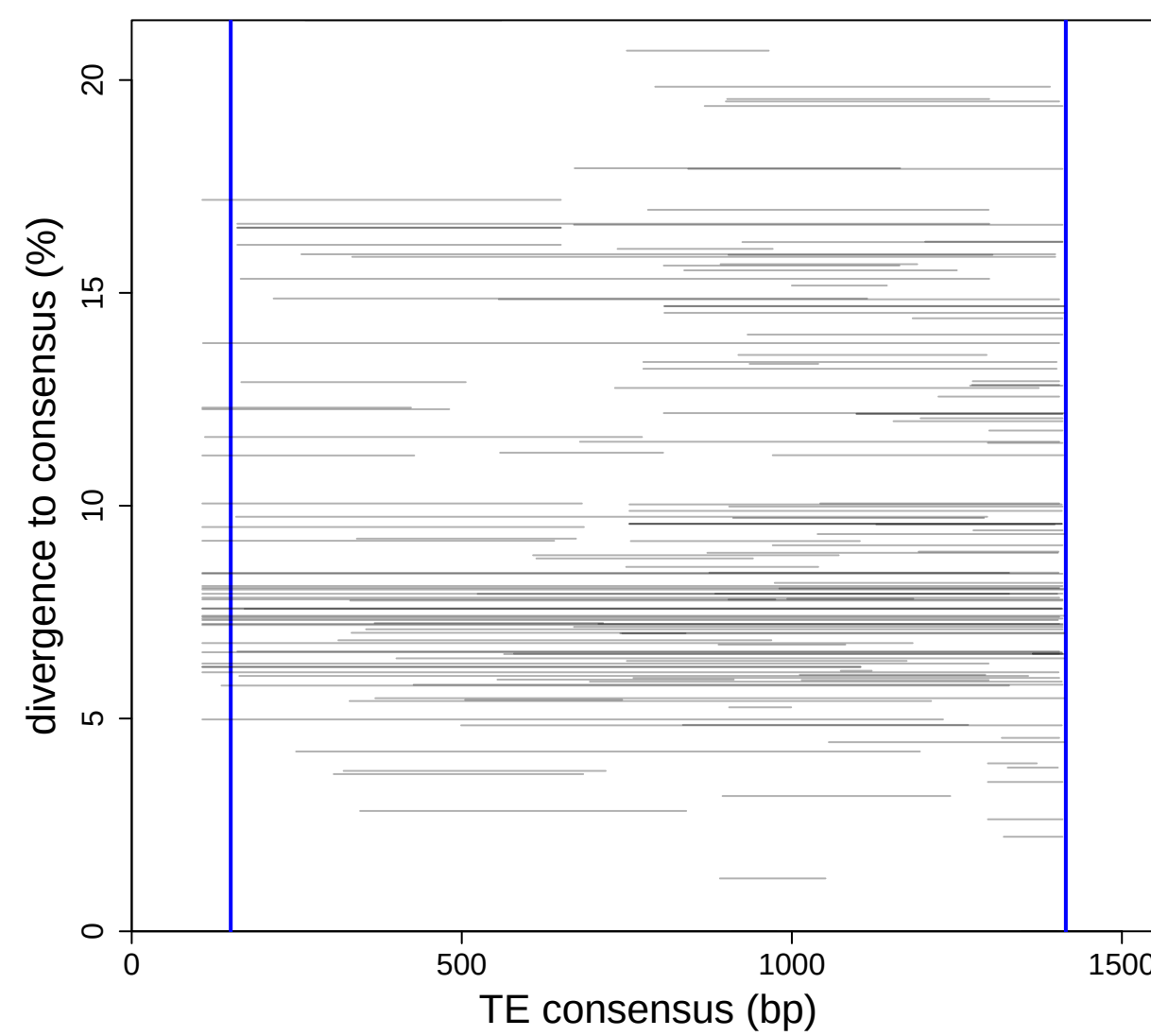
bed_uf.bed g_3.bed fm_1.bed 0_0_bcln.fa aln.fa cl.fa gs.fa c
size: 1255bp; fragments: 161; full length: 27 (≥ 1129.5 bp)

After TEtrimmer 1255 bp

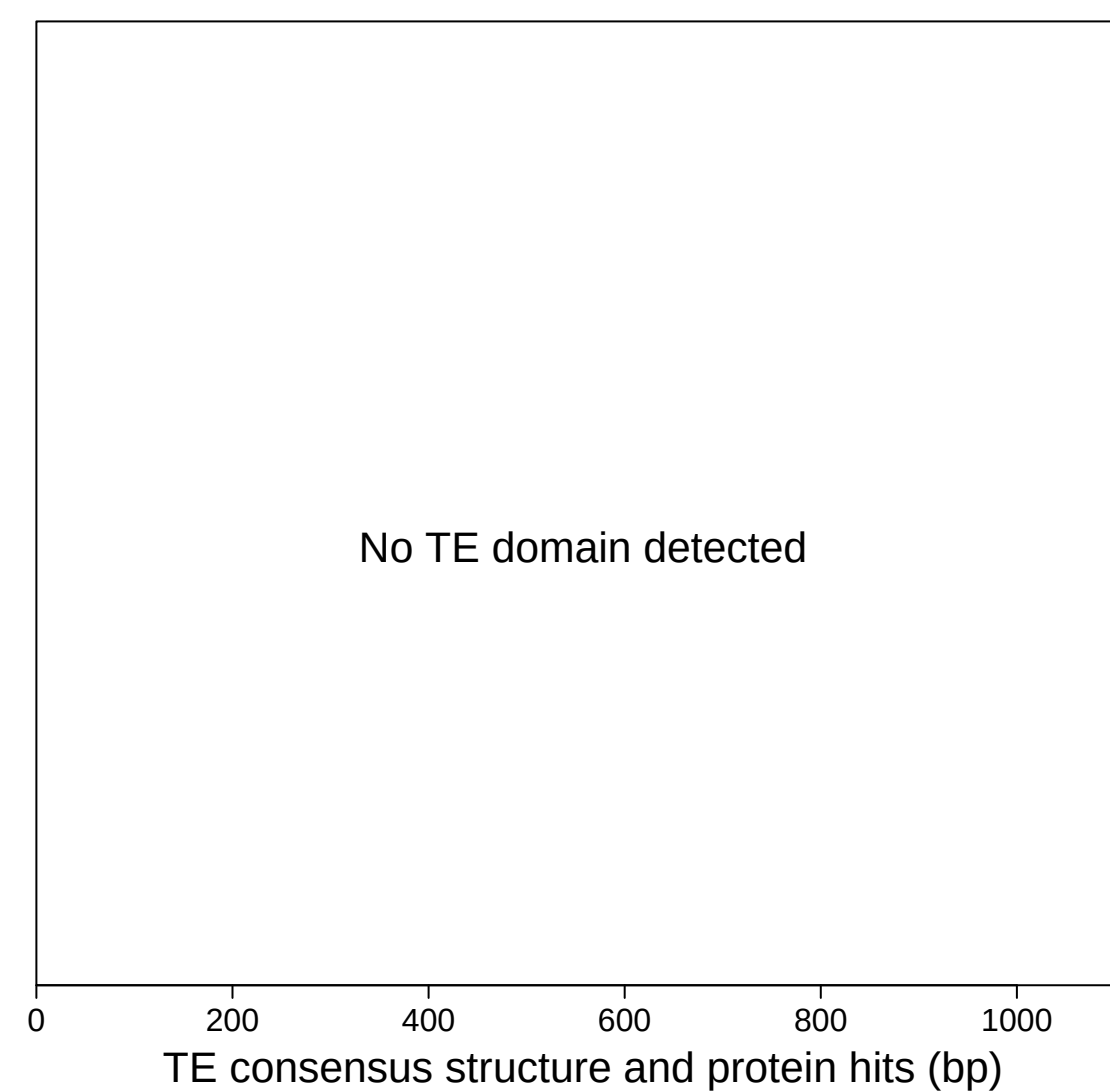
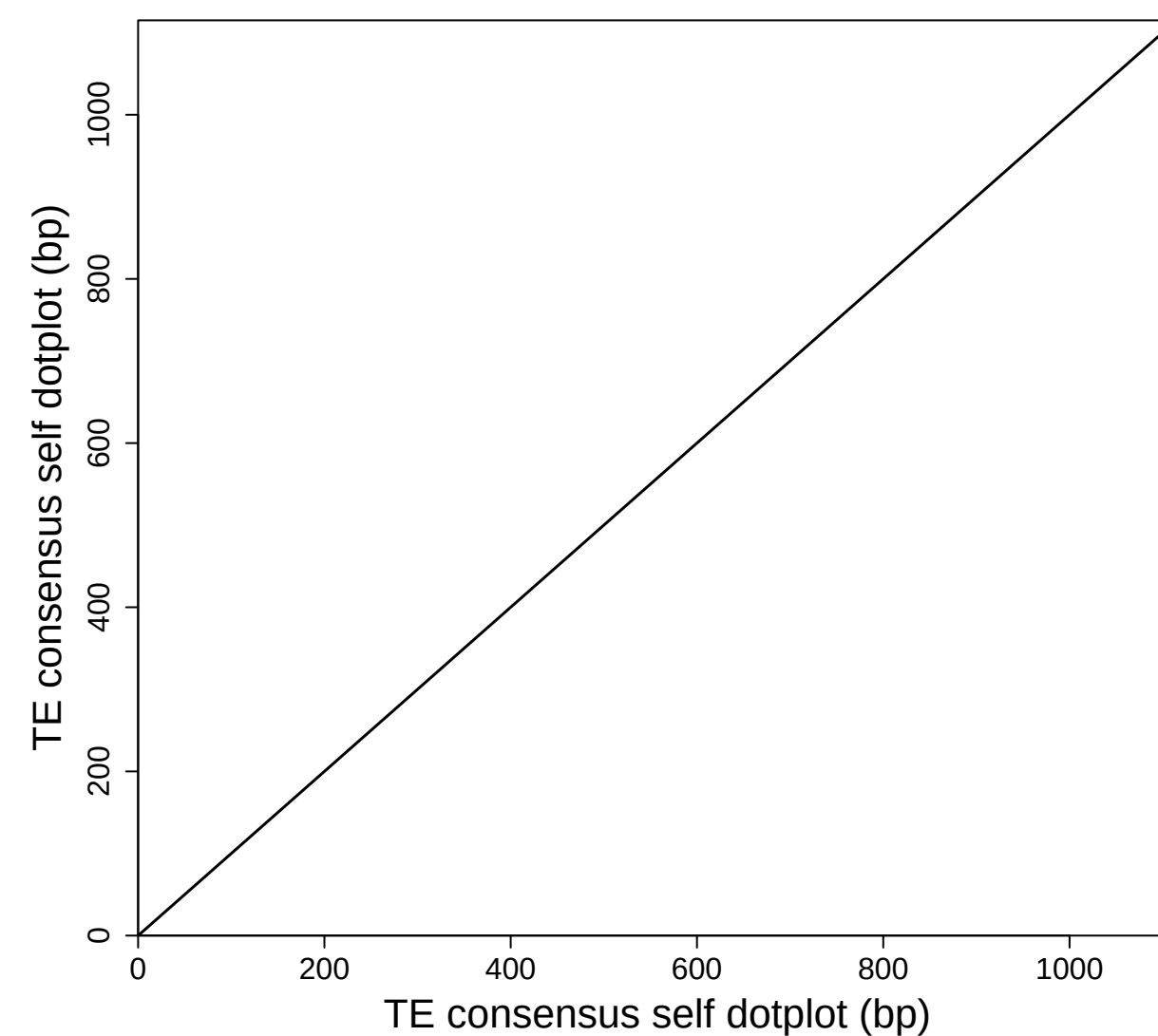
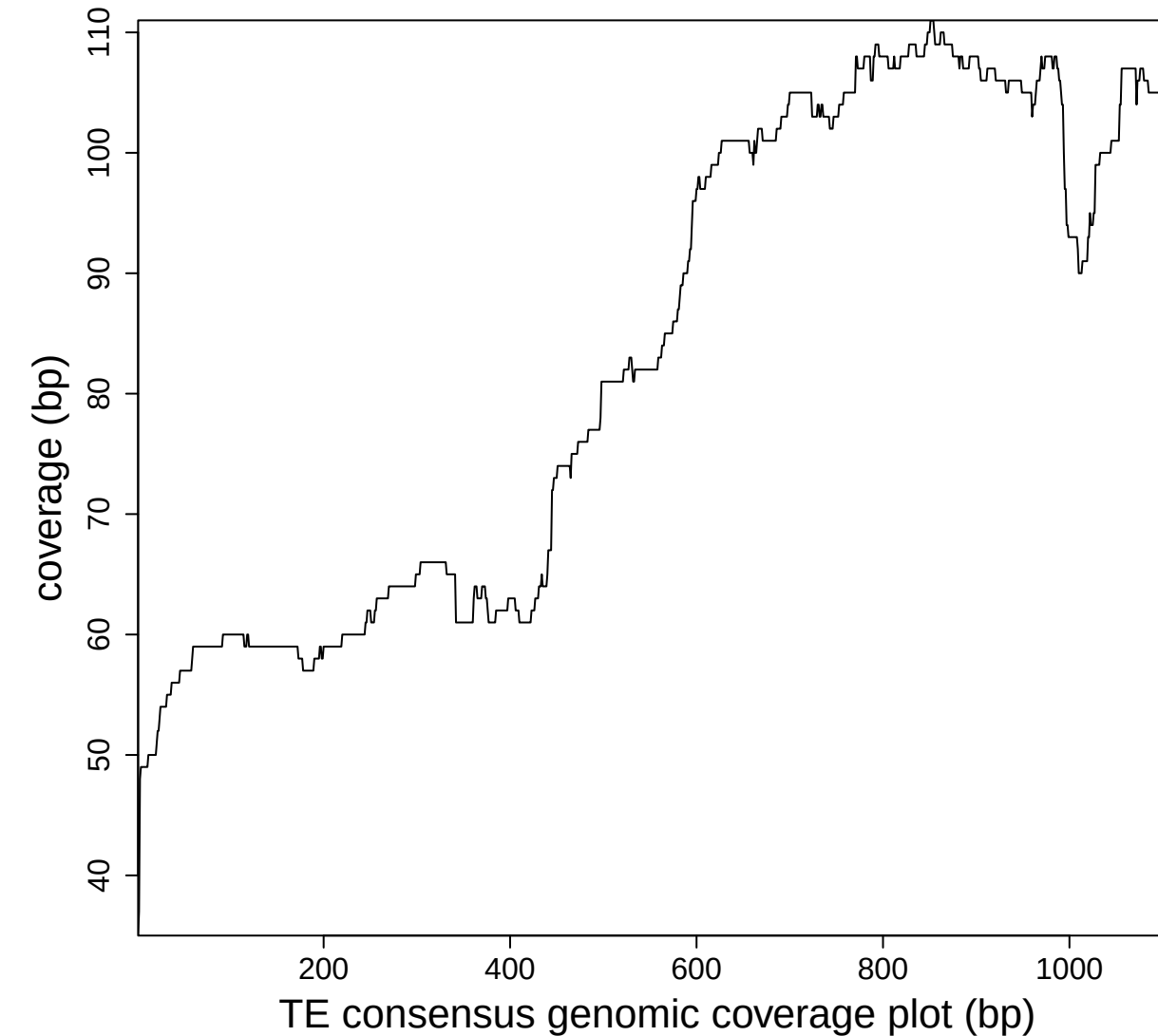
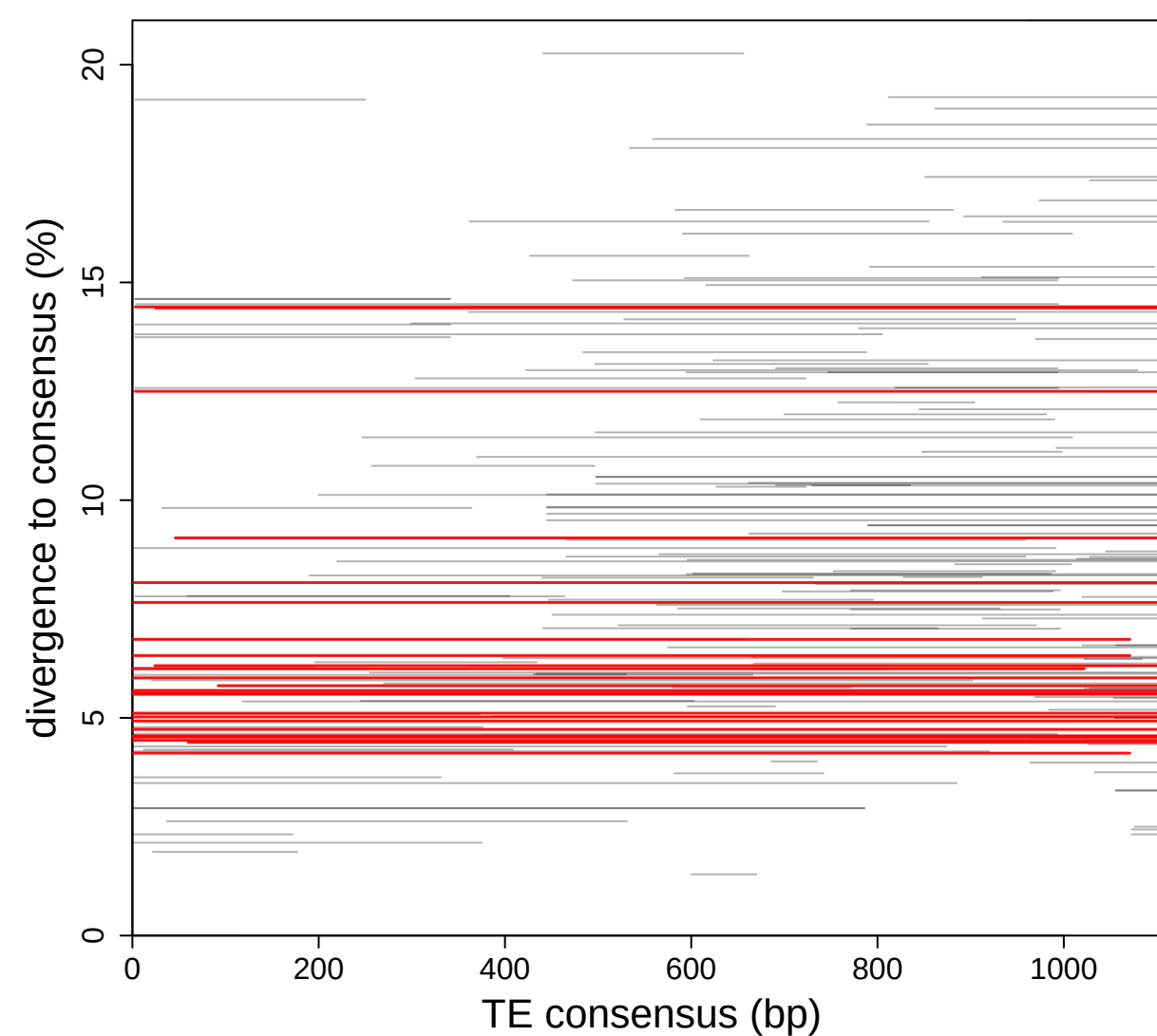


7.fasta.b.bed uf.bed g_3.bed fm_1.bed 0 0 bcln.fa aln.fa cl.f
size: 1565bp; fragments: 160; full length: 0 (>=1408.5bp)

After TEtrimmer Extended plot Blue lines are boundaries

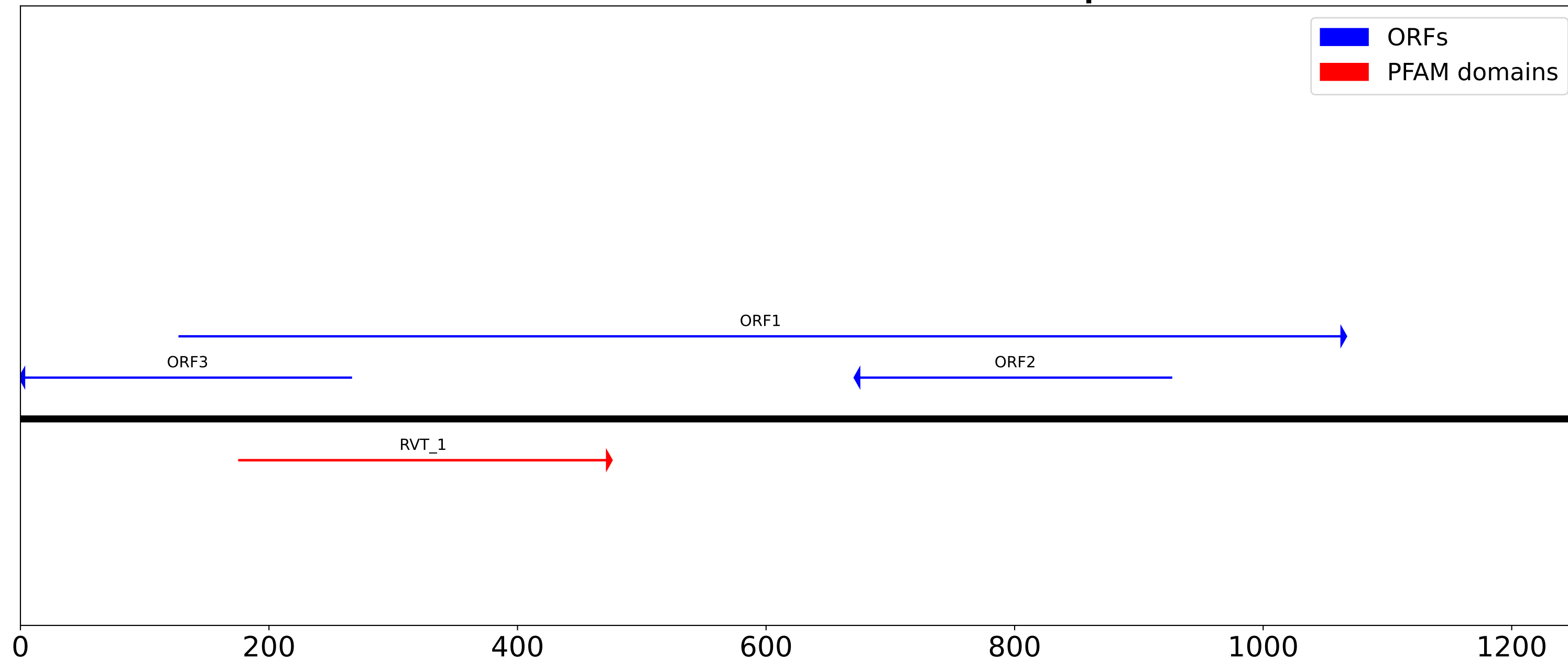


TE: rnd_5_family_3867
size: 1115bp; fragments: 188; full length: 28 (≥ 1003.5 bp)

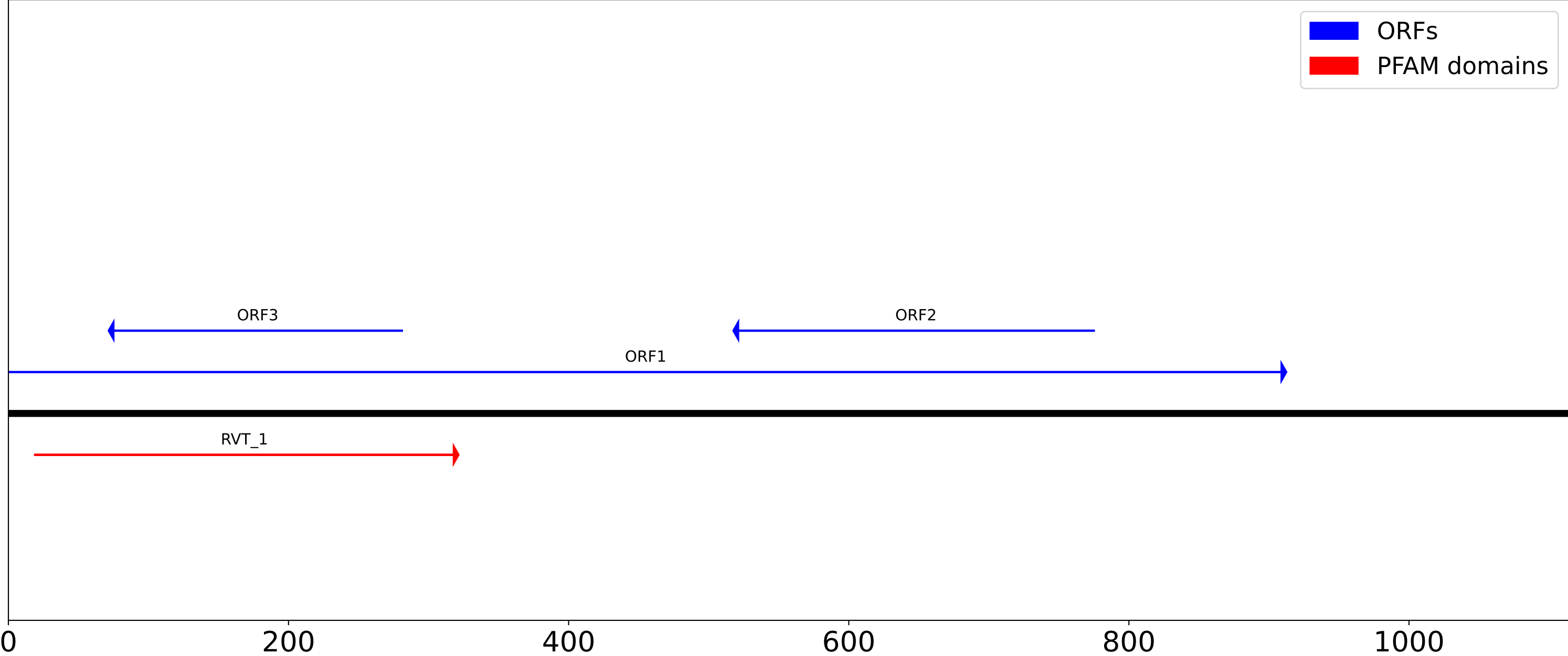


Before TEtrimmer 1115 bp

After TEtrimmer ORF and PFAM domain plot



Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

