

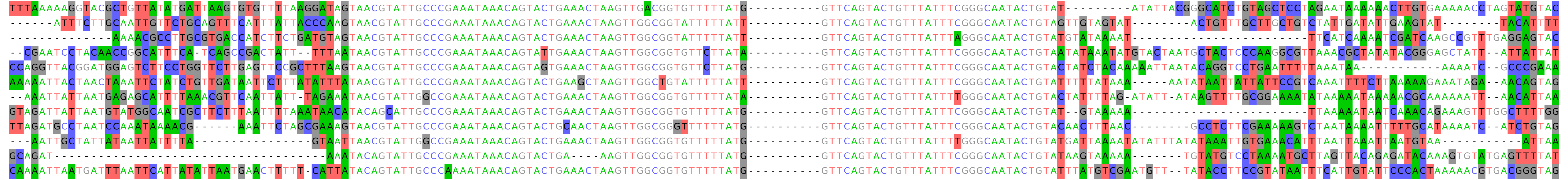
Start crop Point



End crop Point



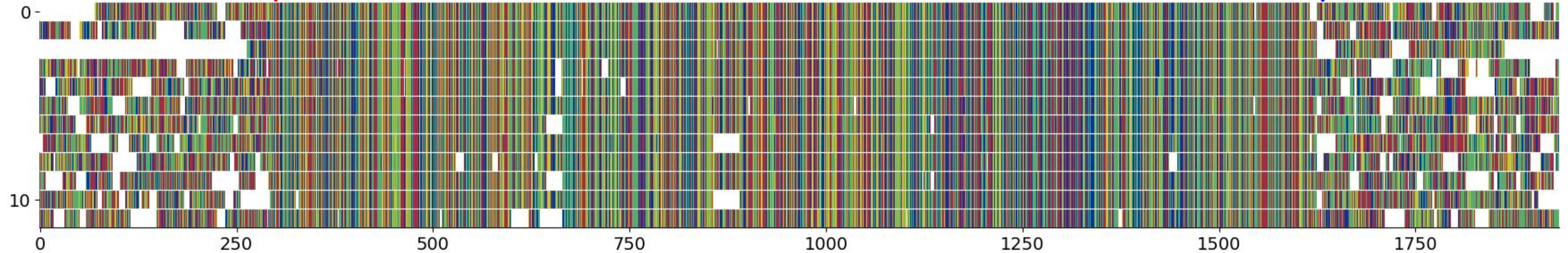
MSA length = 1333



Start crop Point

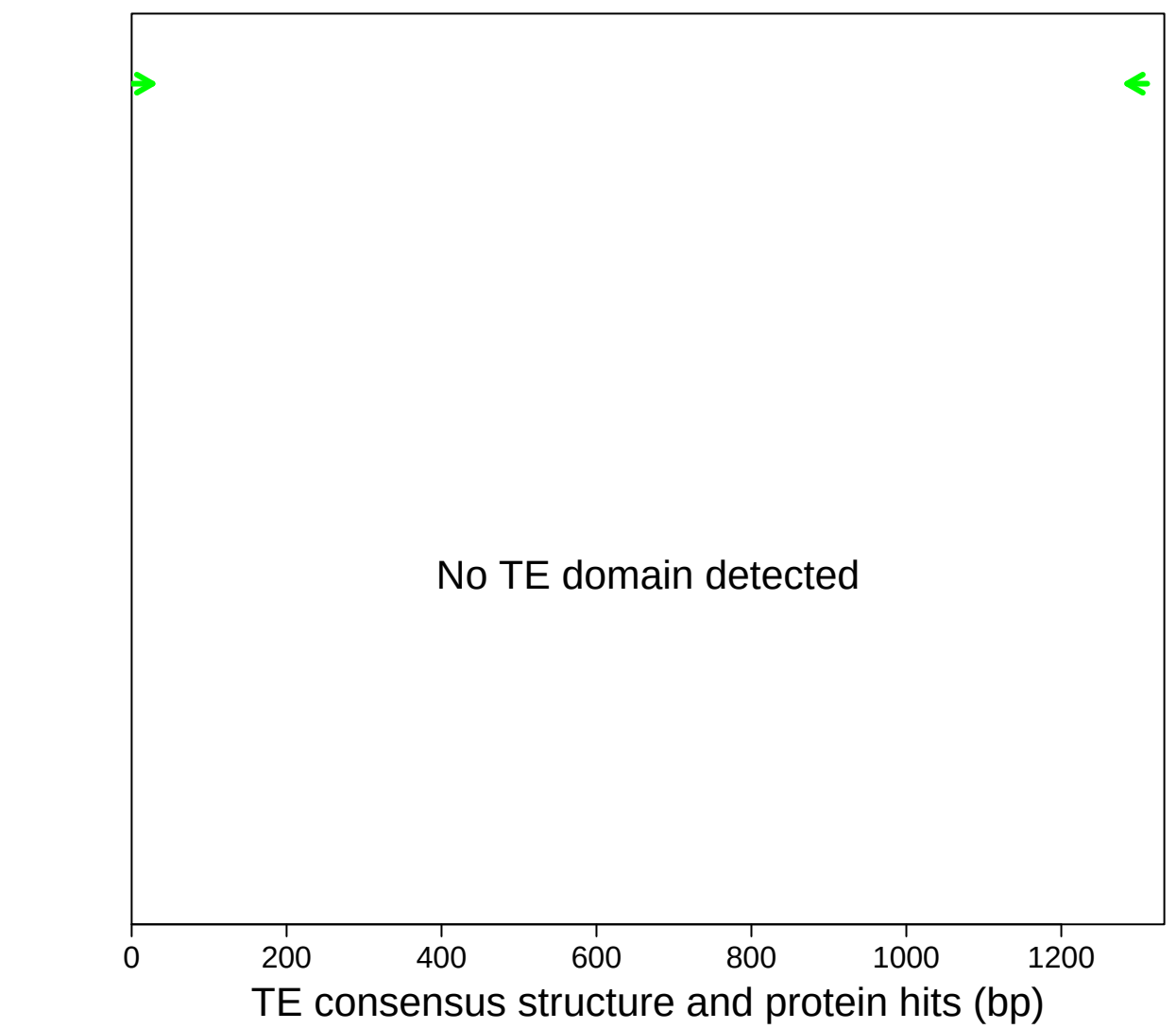
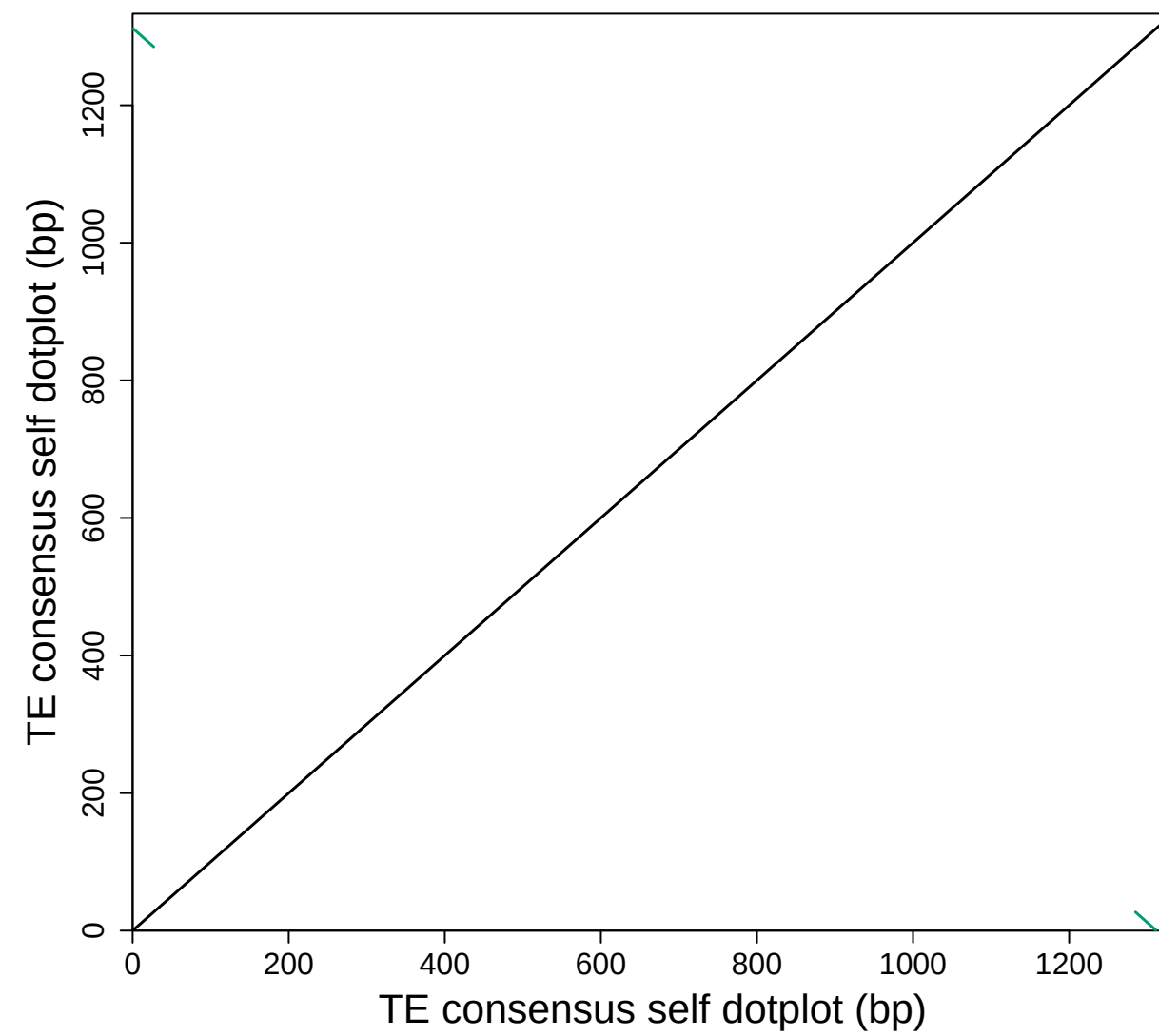
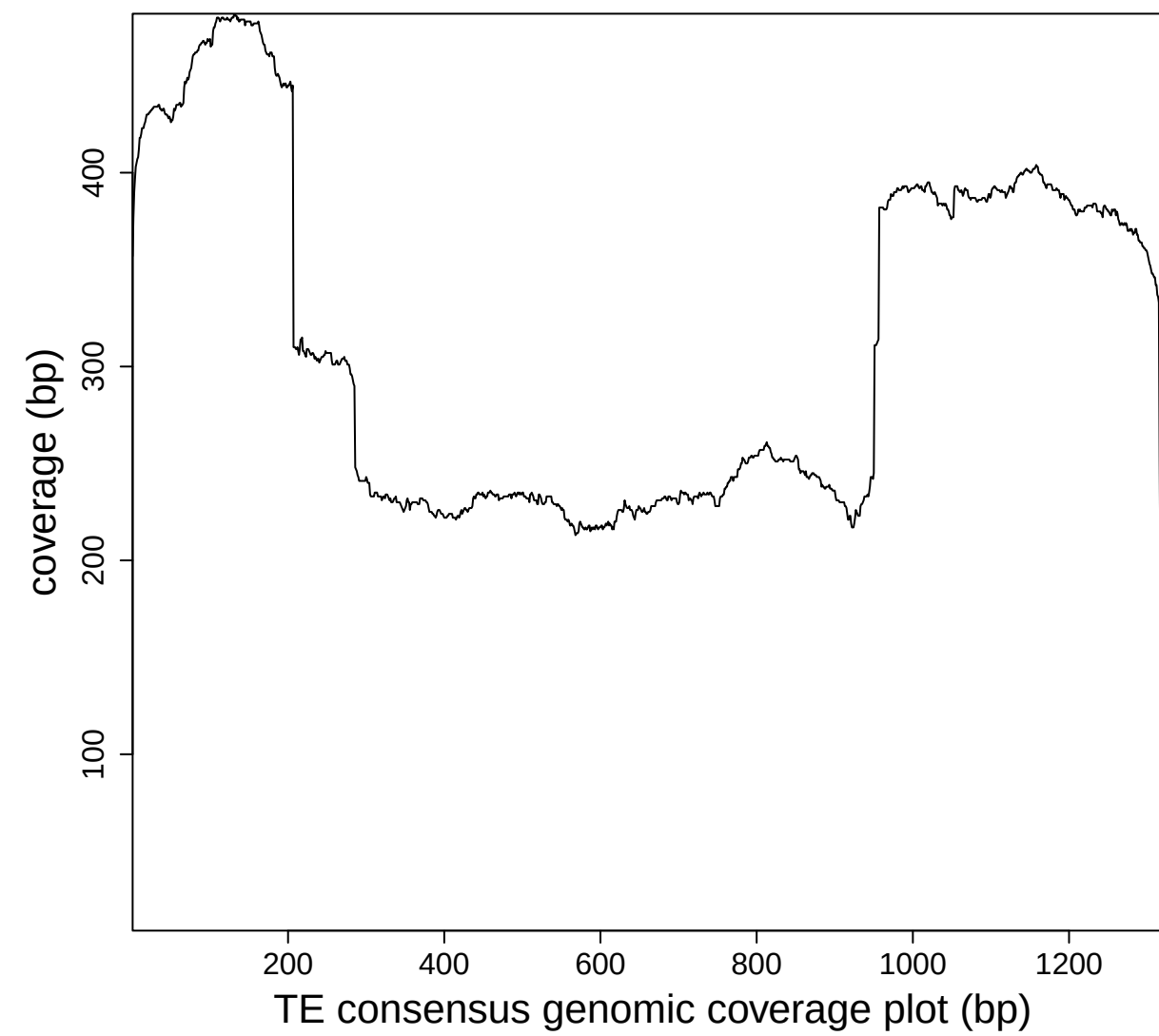
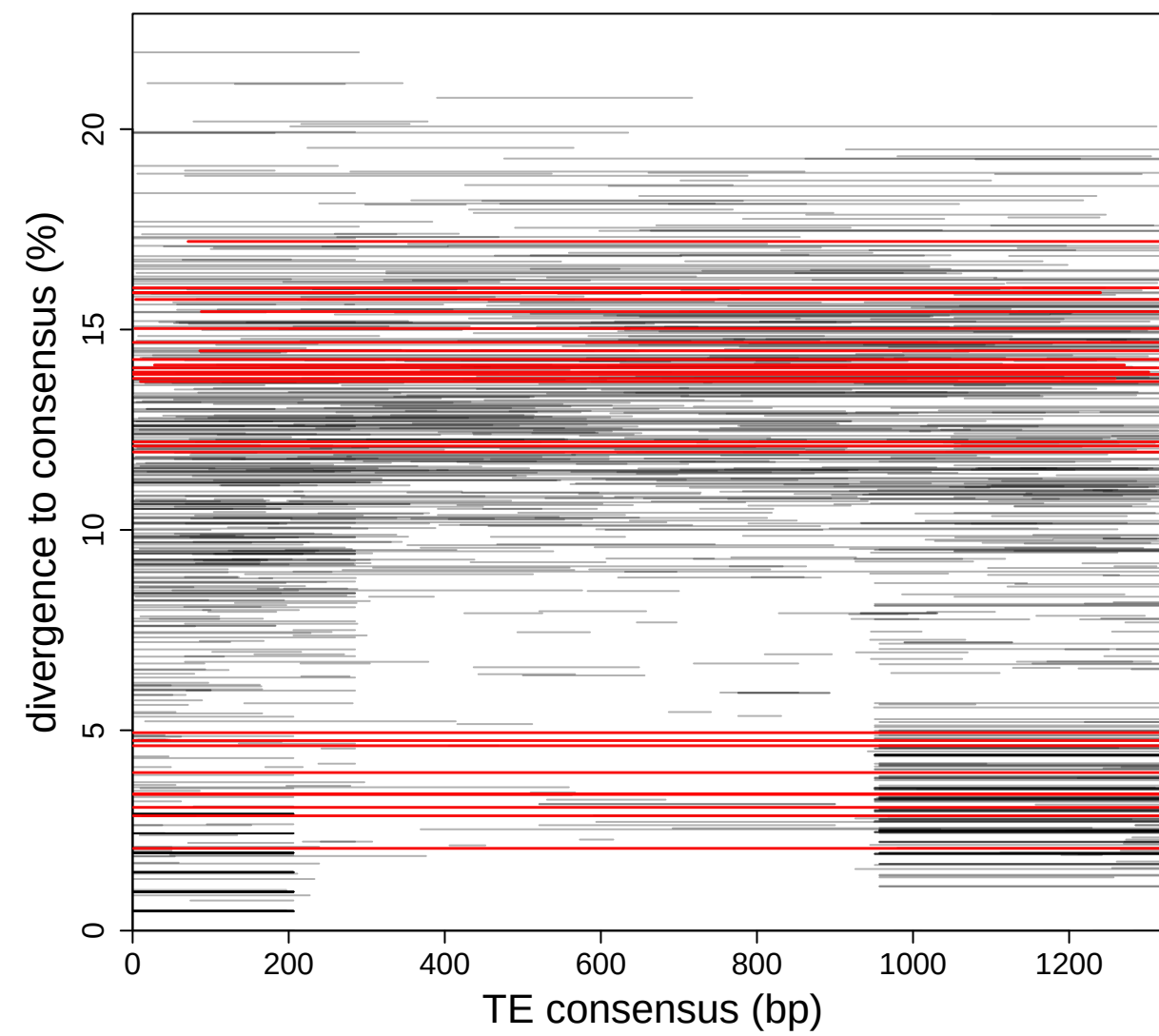


End crop Point



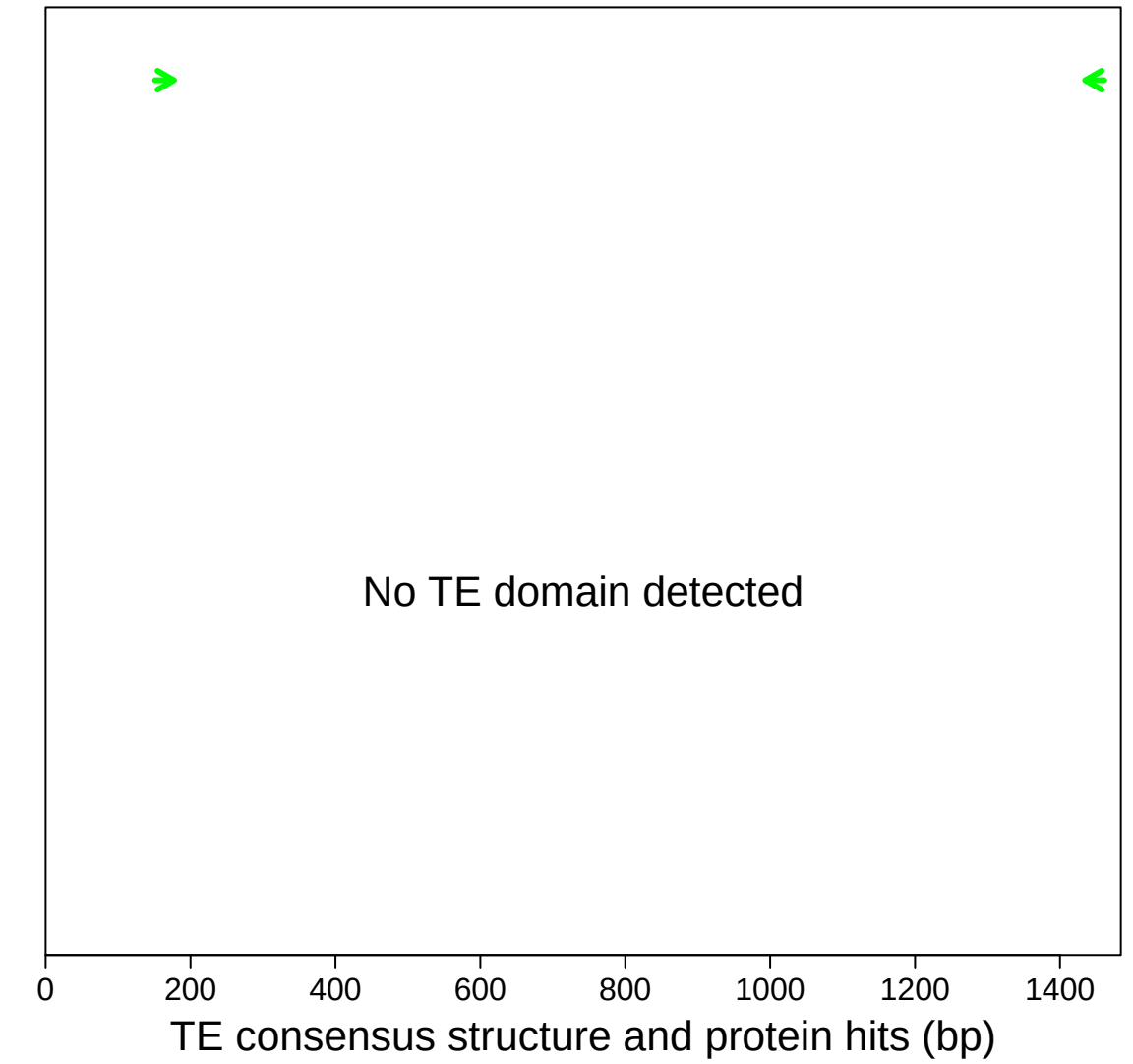
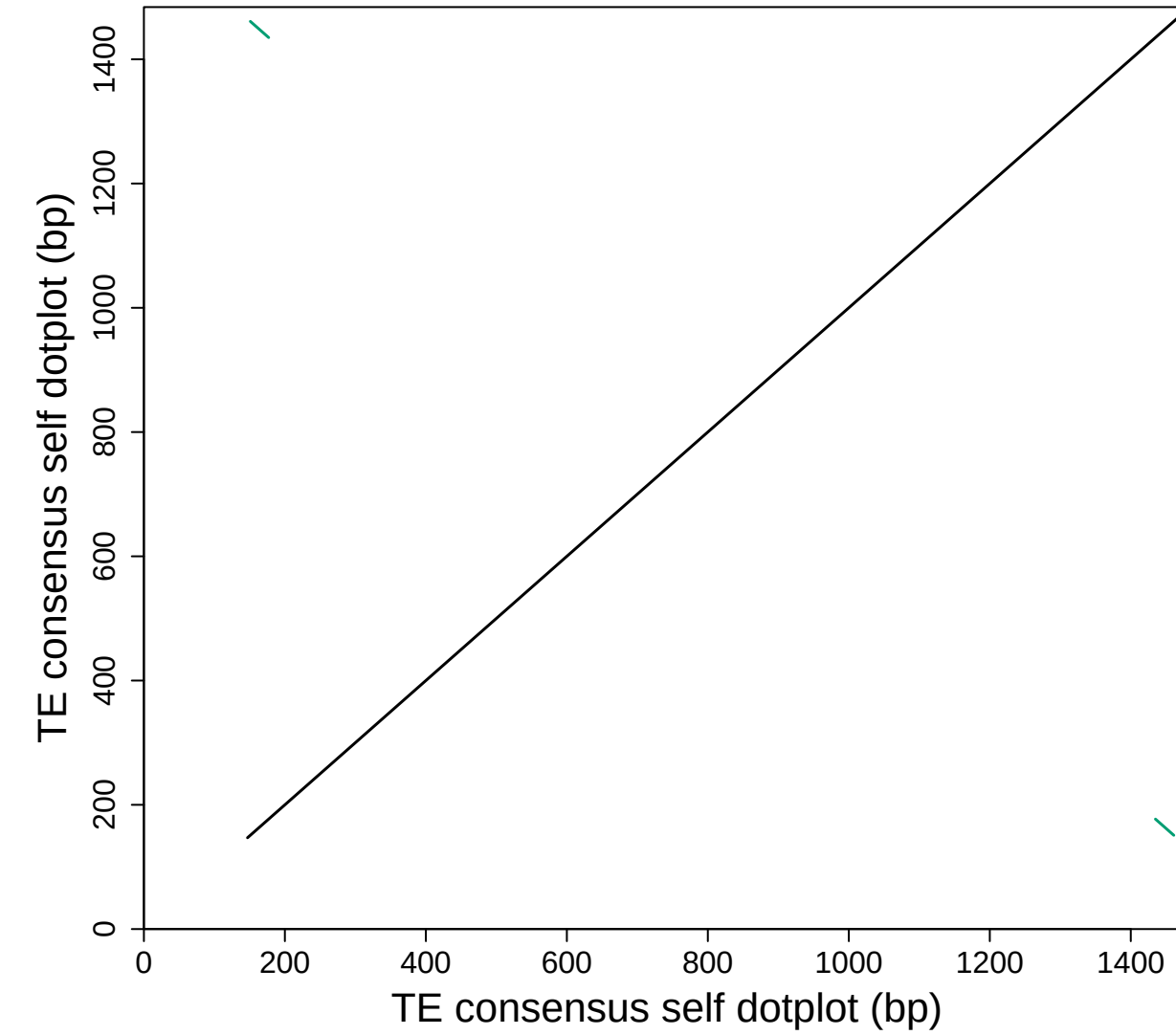
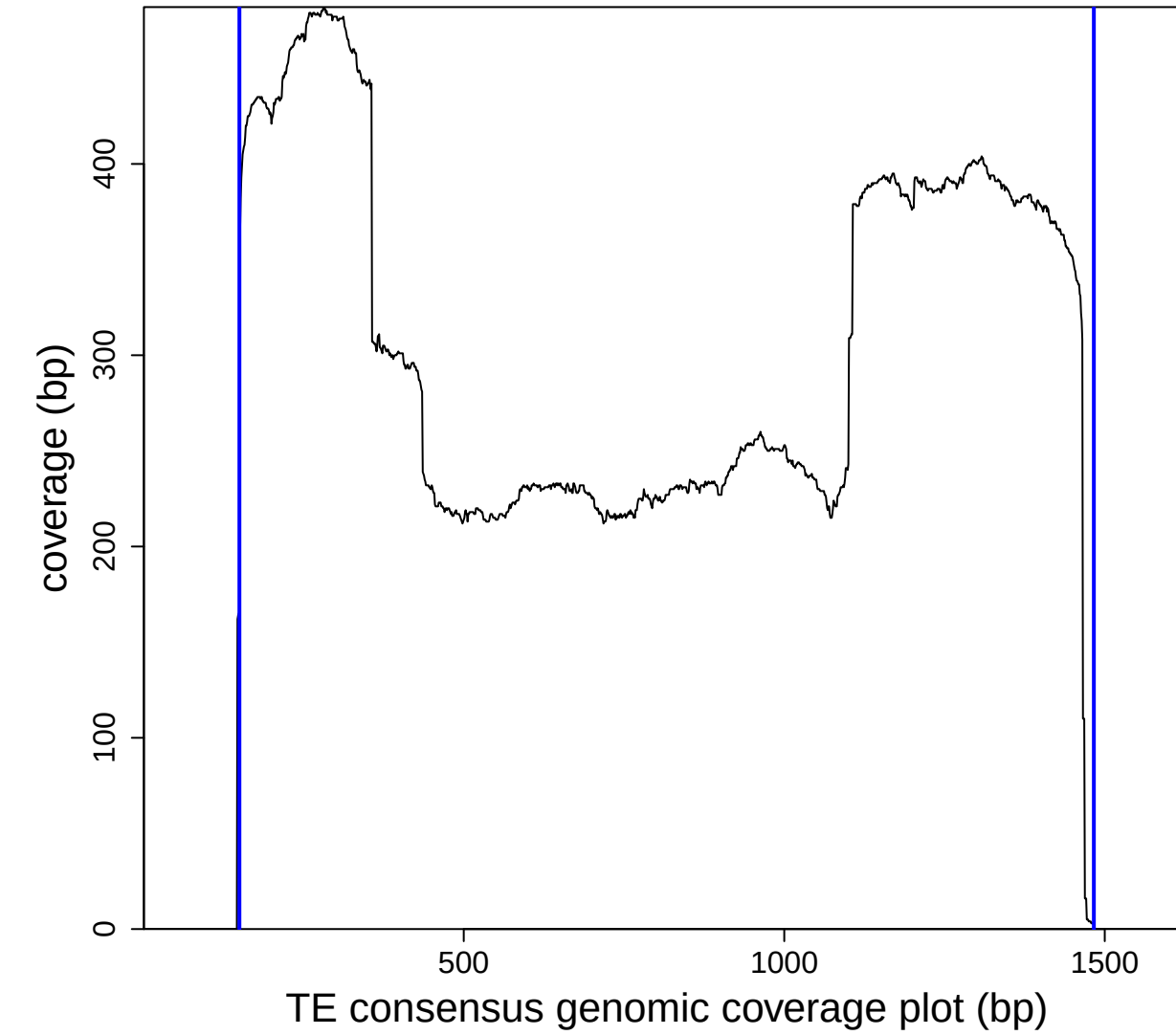
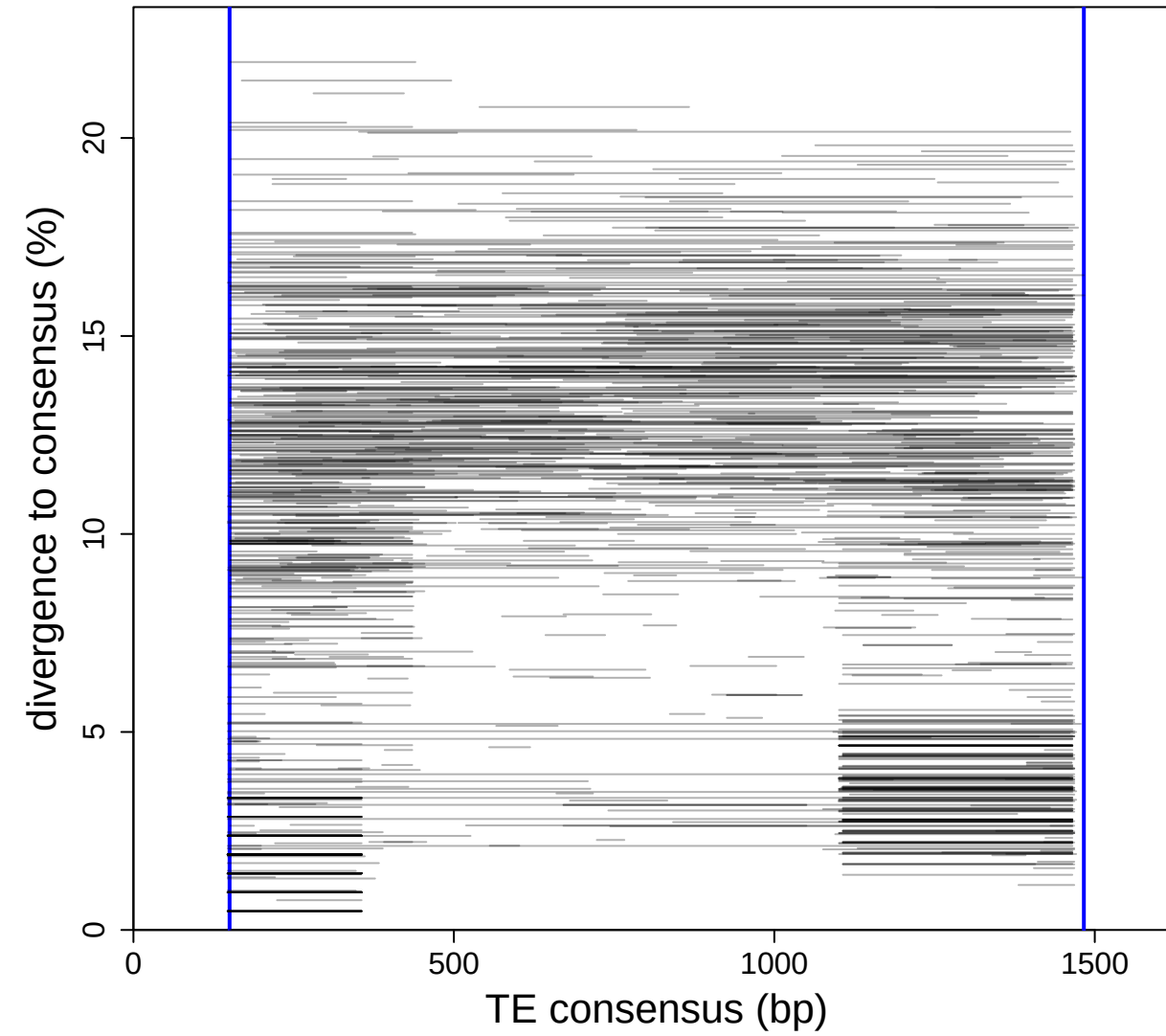
l.b.bed uf.bed g_1.bed fm_1.bed 0 0 bcln.fa aln.fa cl.fa gs.fa
size: 1333bp; fragments: 1211; full length: 28 (>=1199.7bp)

After TEtrimmer 1333 bp



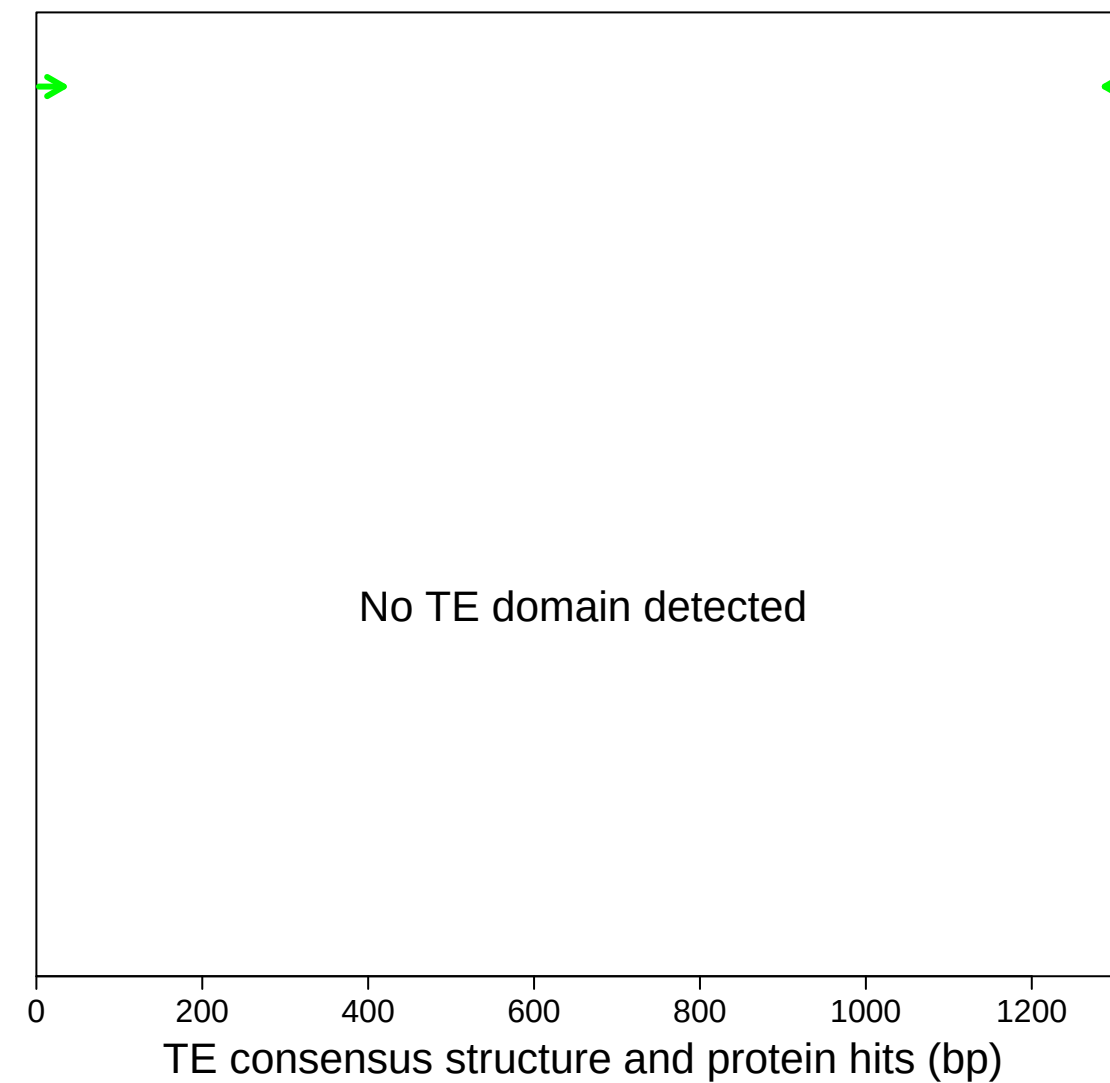
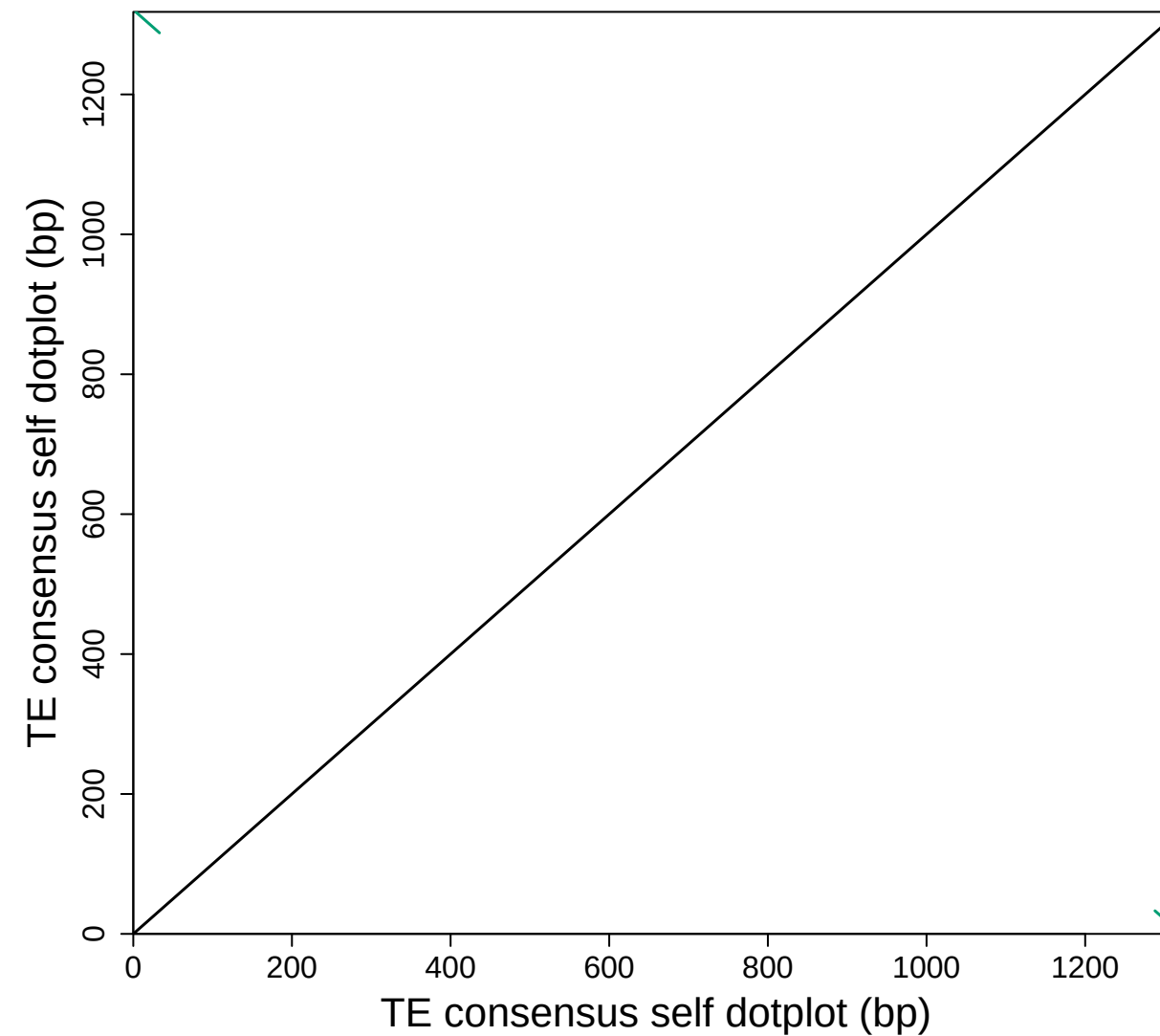
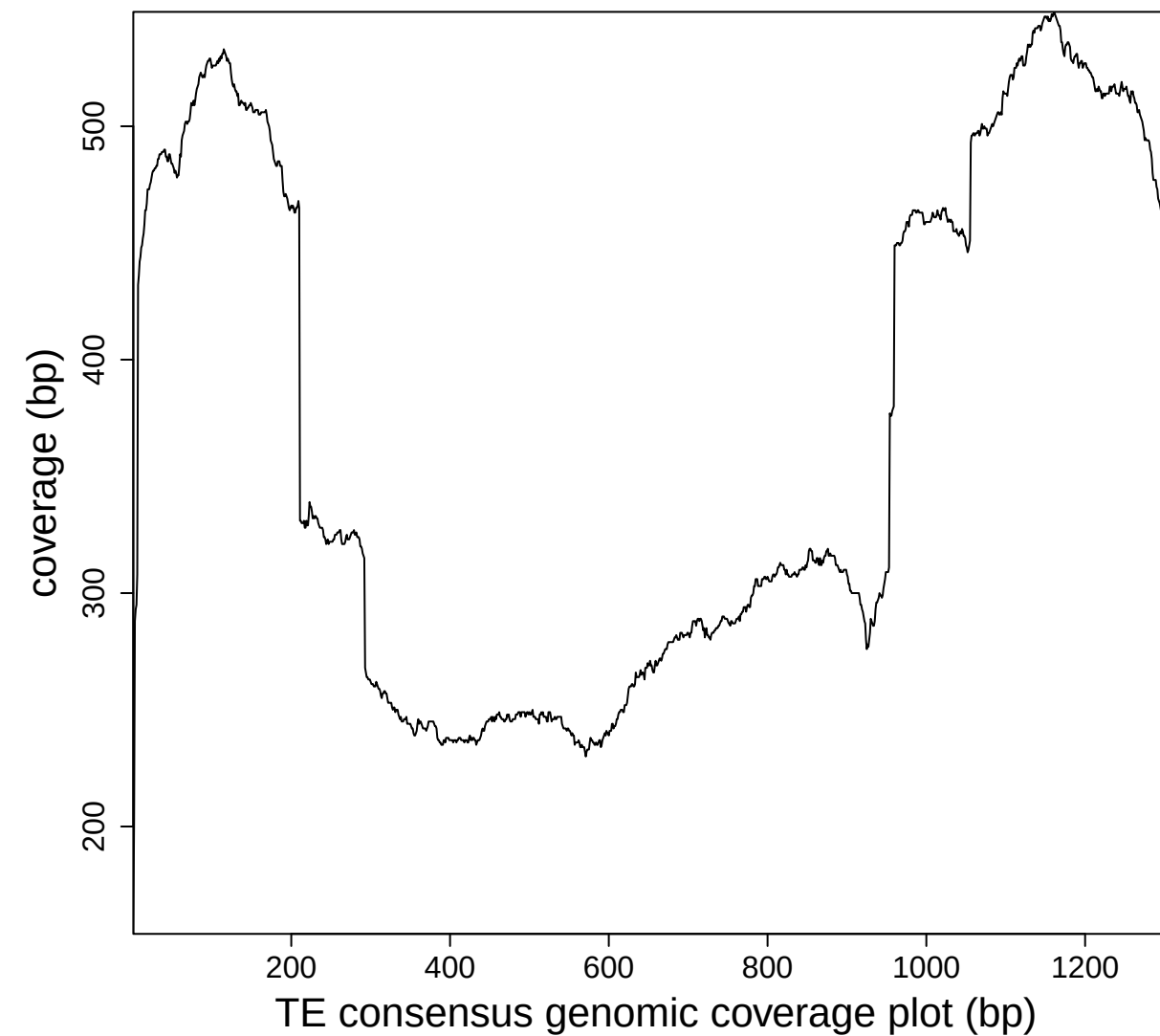
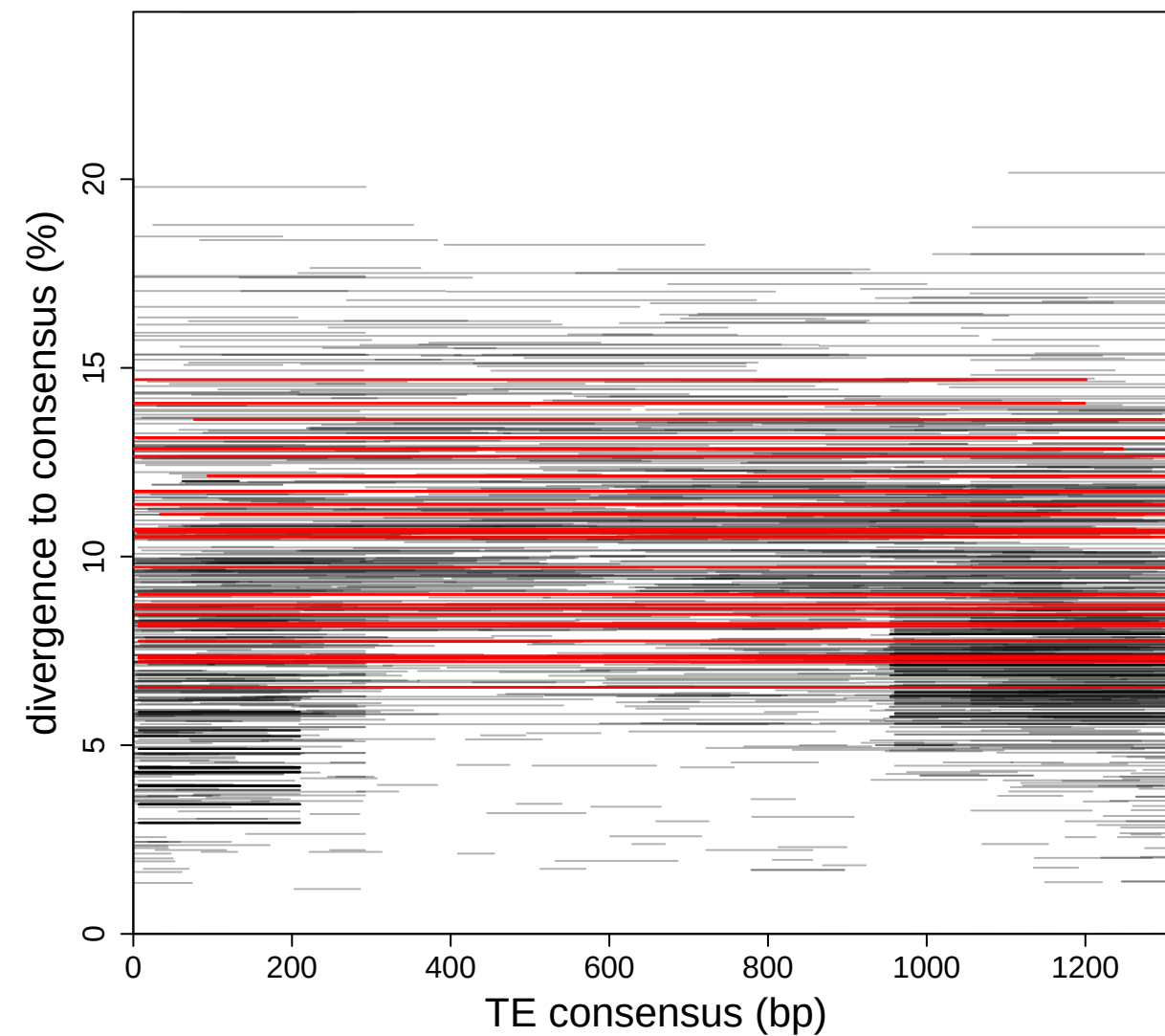
0.fasta.b.bed uf.bed g 1.bed fm 1.bed 0 0 bcln.fa aln.fa cl.fa
size: 1633bp; fragments: 1192; full length: 0 (>=1469.7bp)

After TEtrimmer Extended plot Blue lines are boundaries

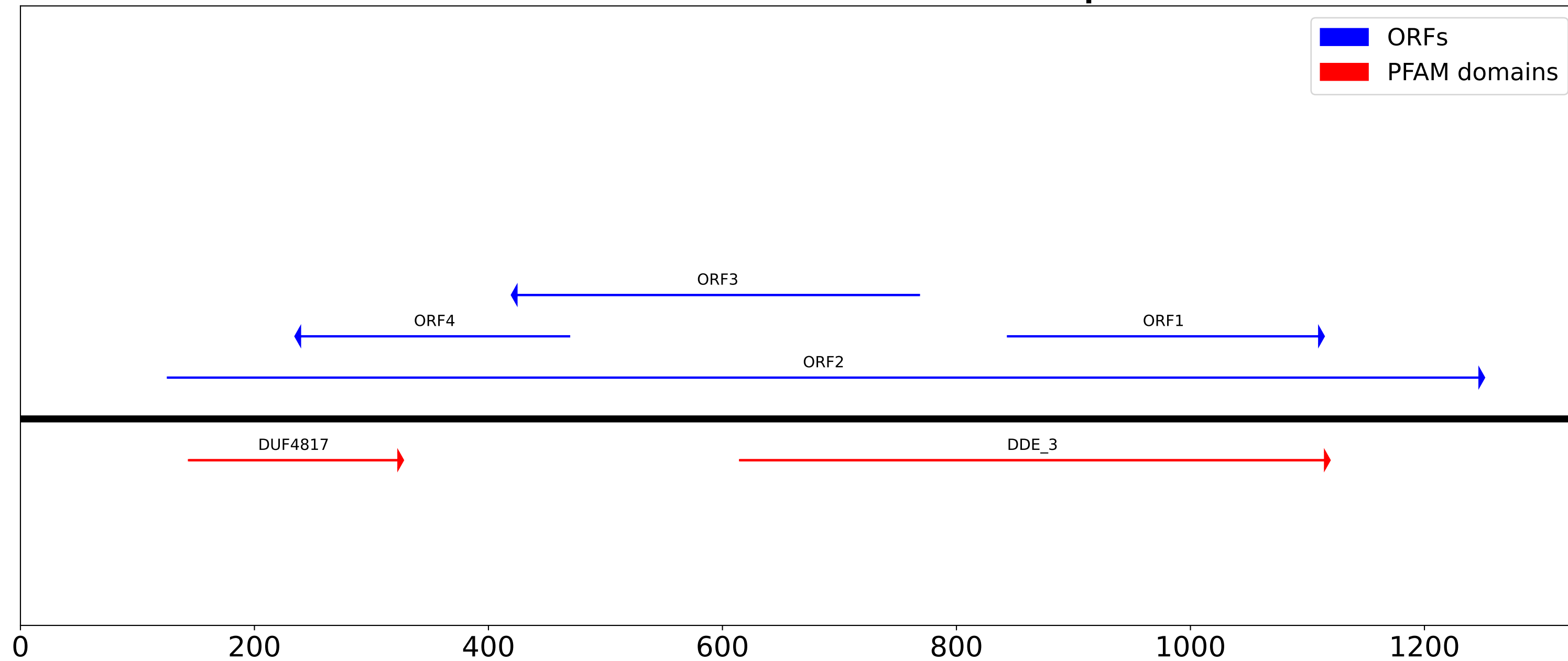


TE: rnd_5_family_970
size: 1318bp; fragments: 1554; full length: 32 (≥ 1186.2 bp)

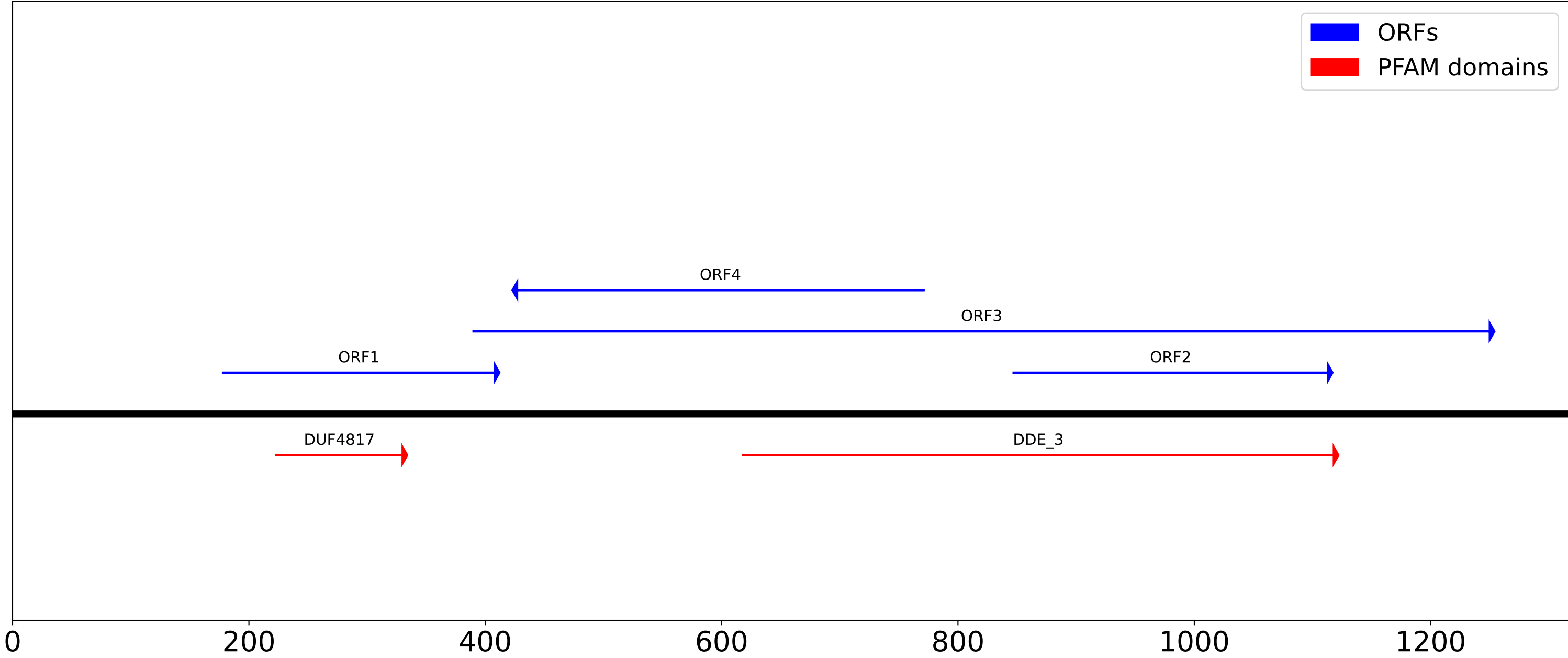
Before TEtrimmer 1318 bp



After TEtrimmer ORF and PFAM domain plot



Before TEtrimmer ORF and PFAM domain plot



Dotplot

(window size = 25, threshold = 50.00 10/10/25)

