

Start crop Point

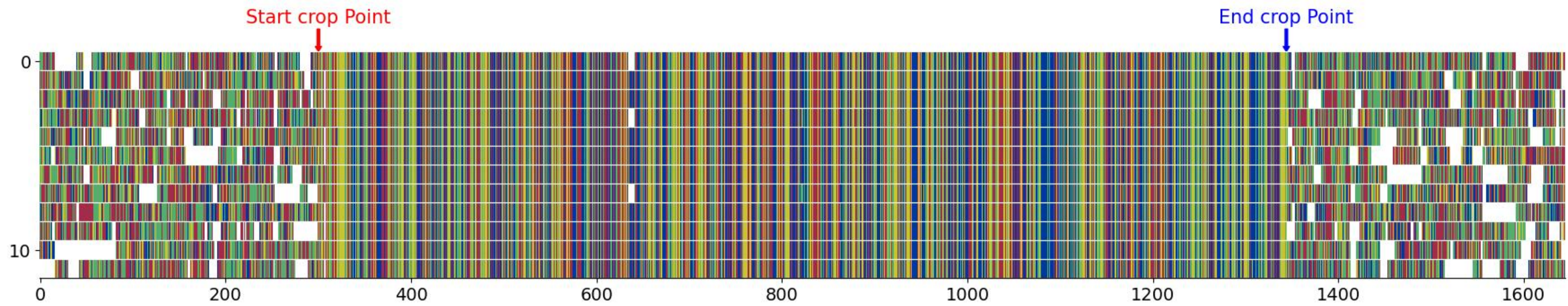


End crop Point



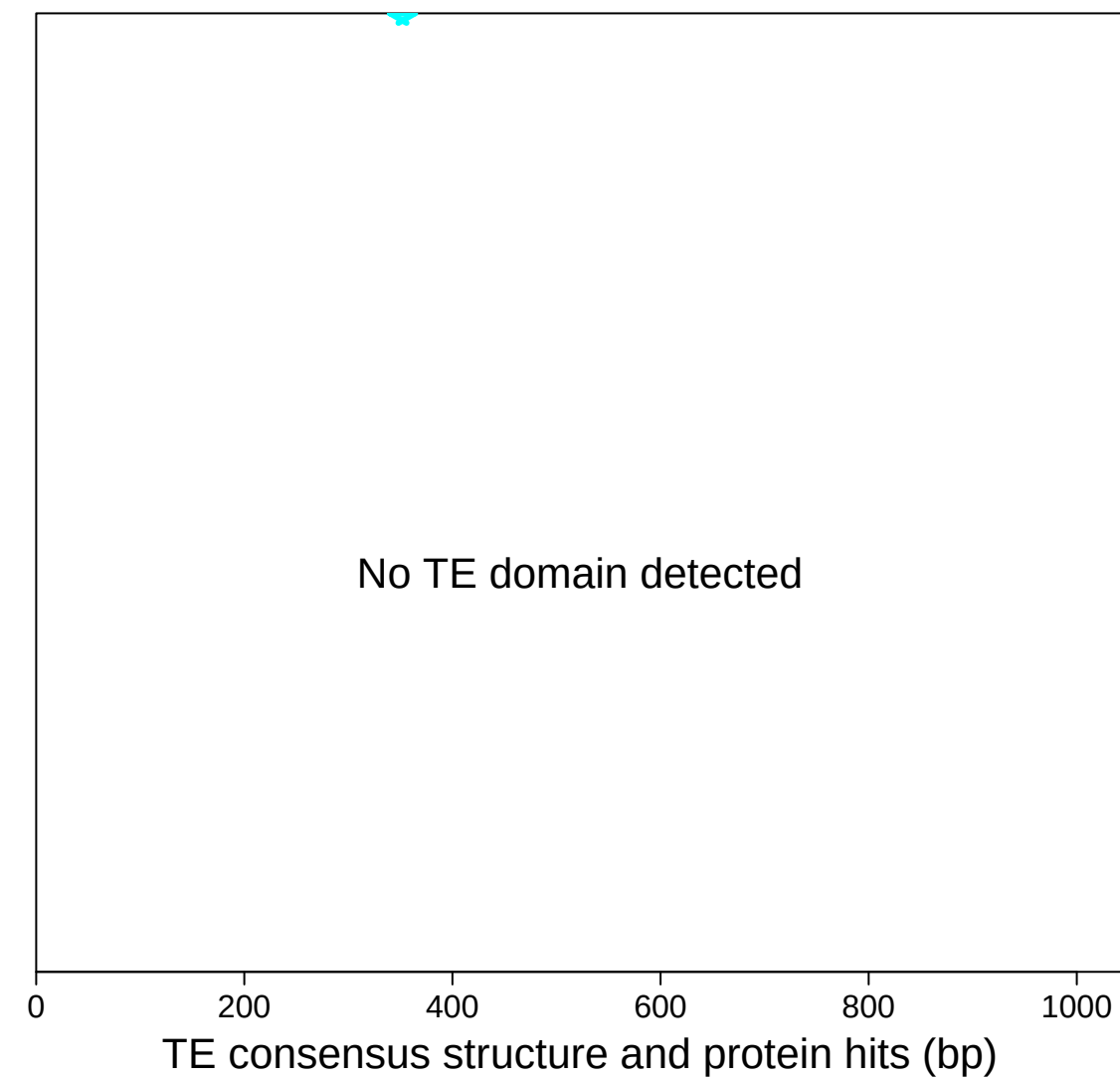
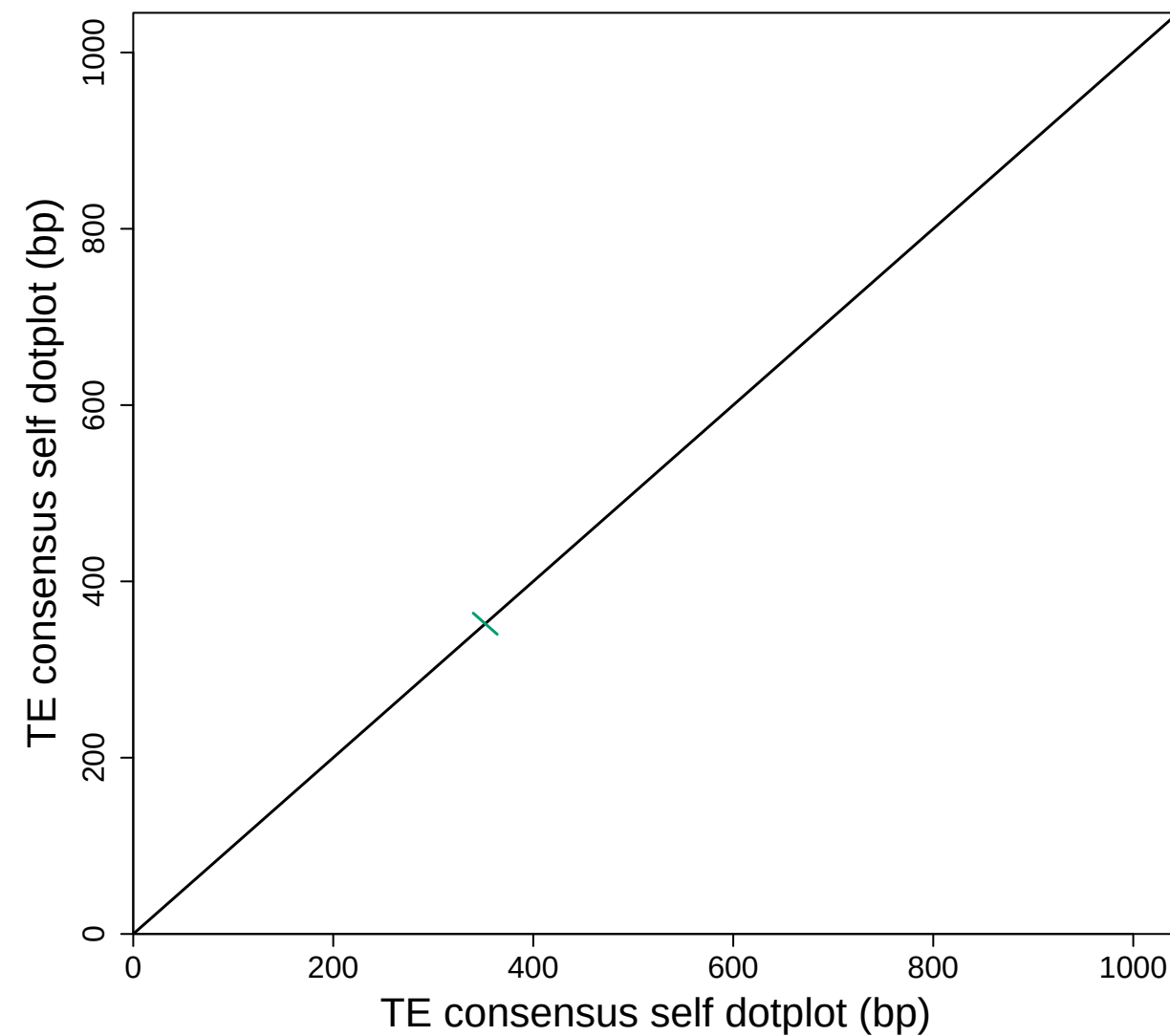
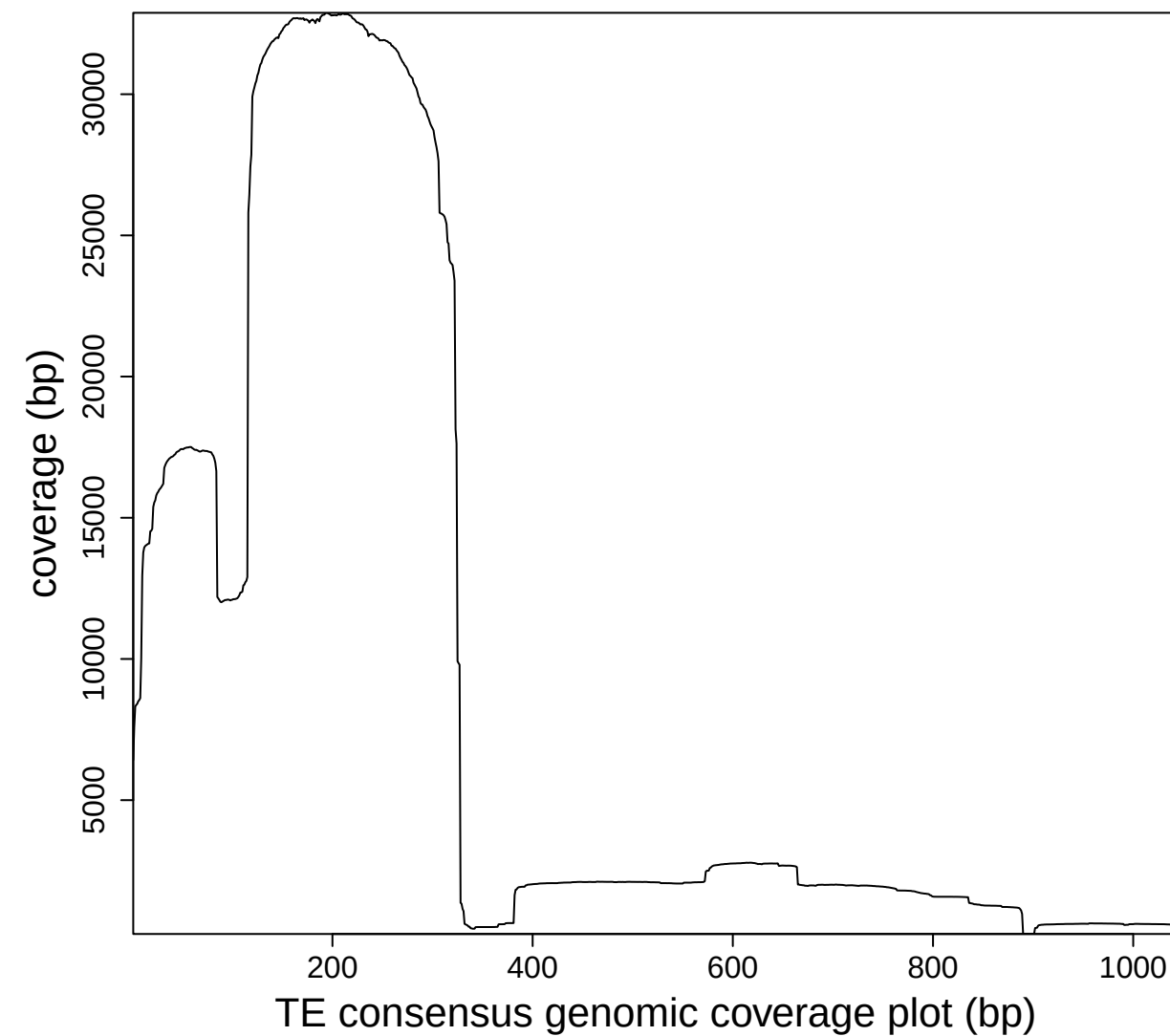
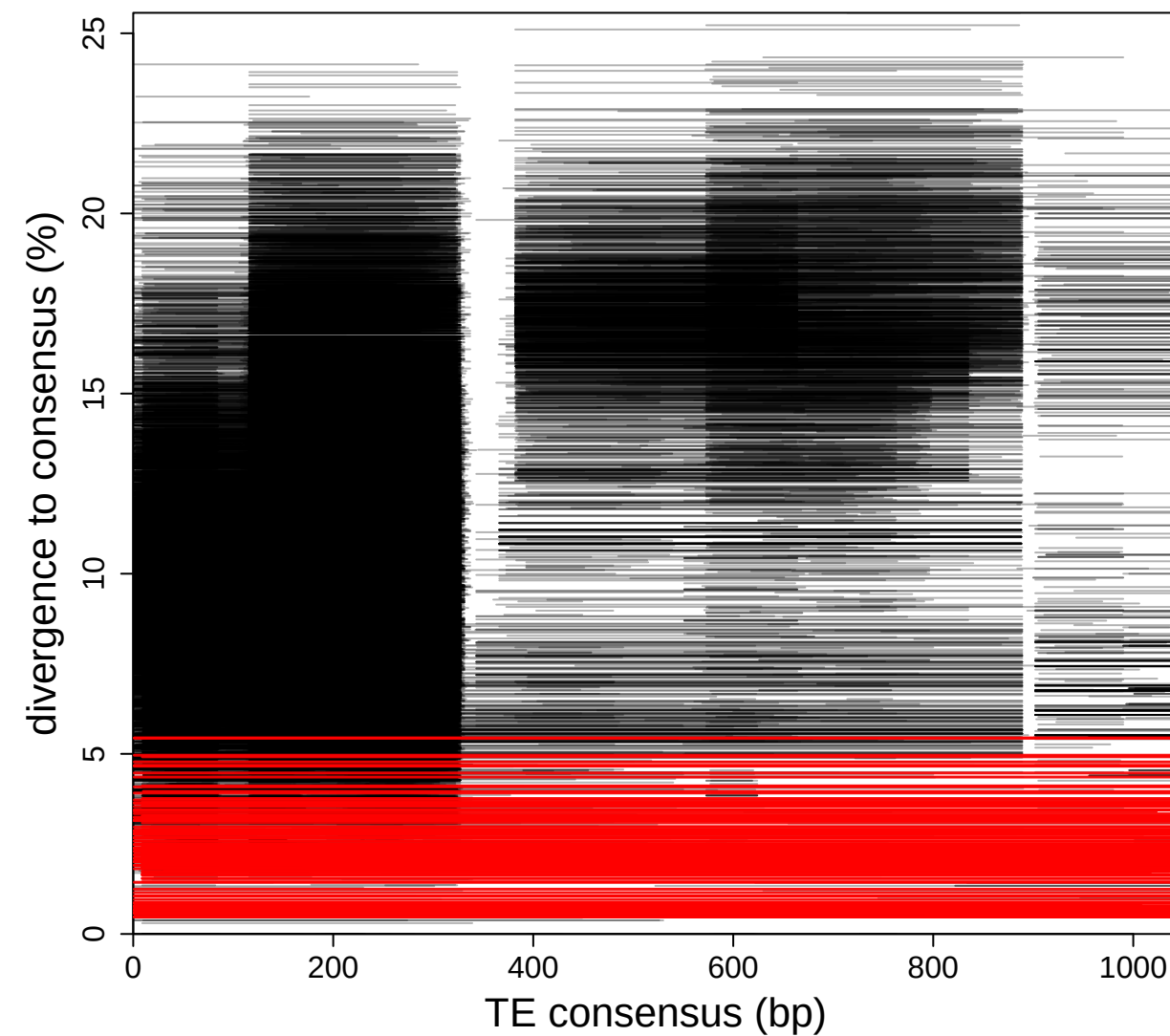
MSA length = 1045





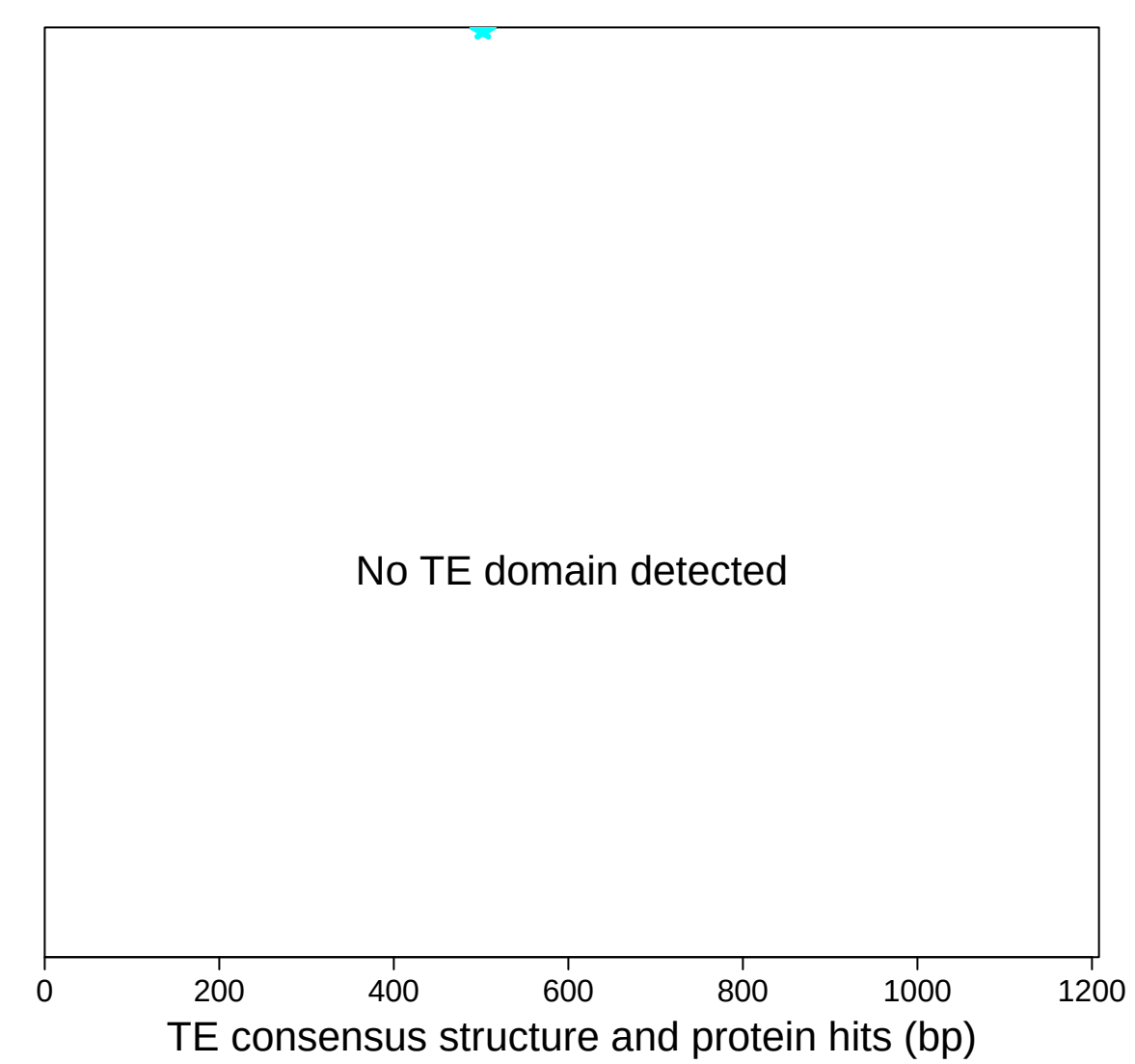
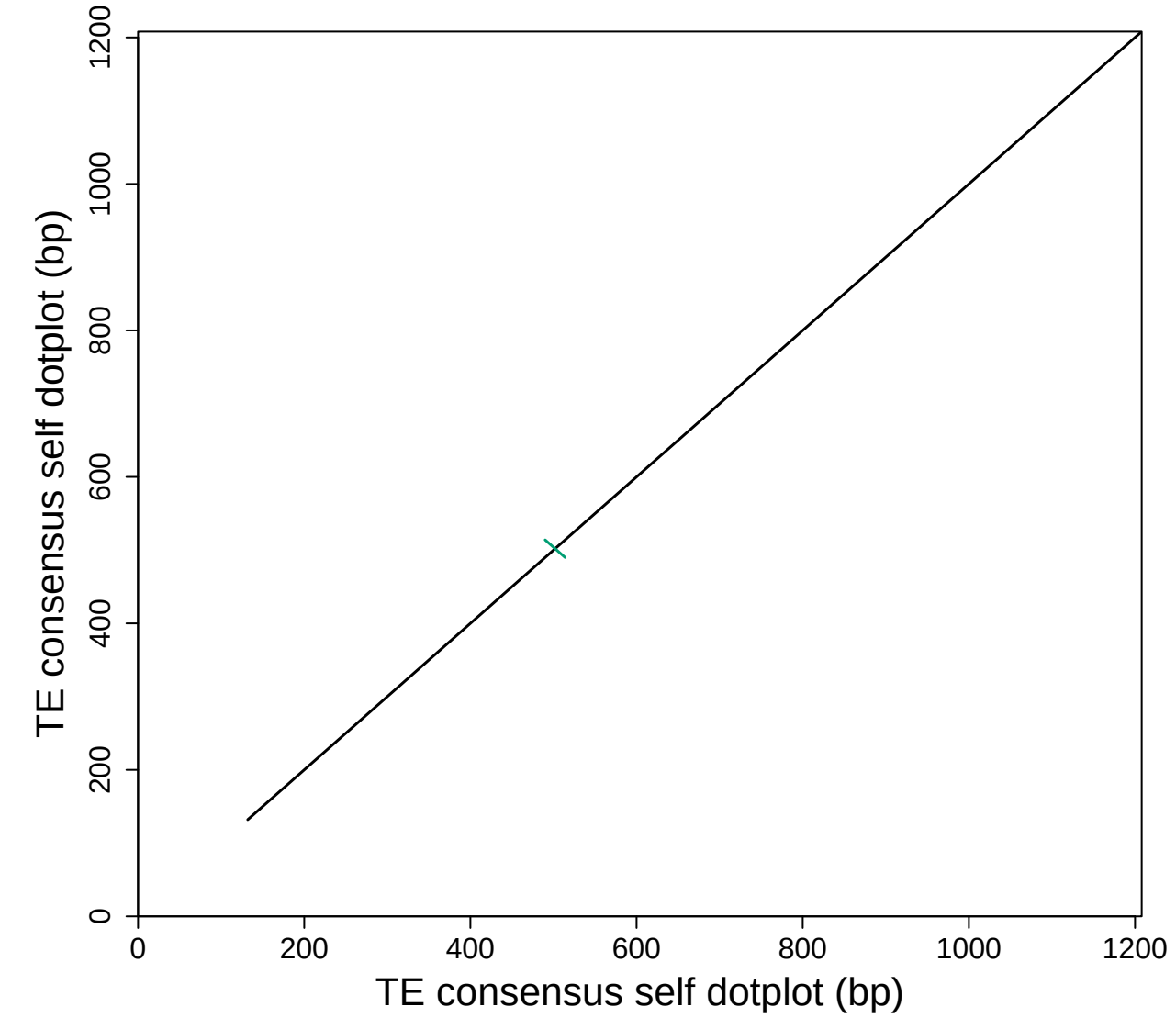
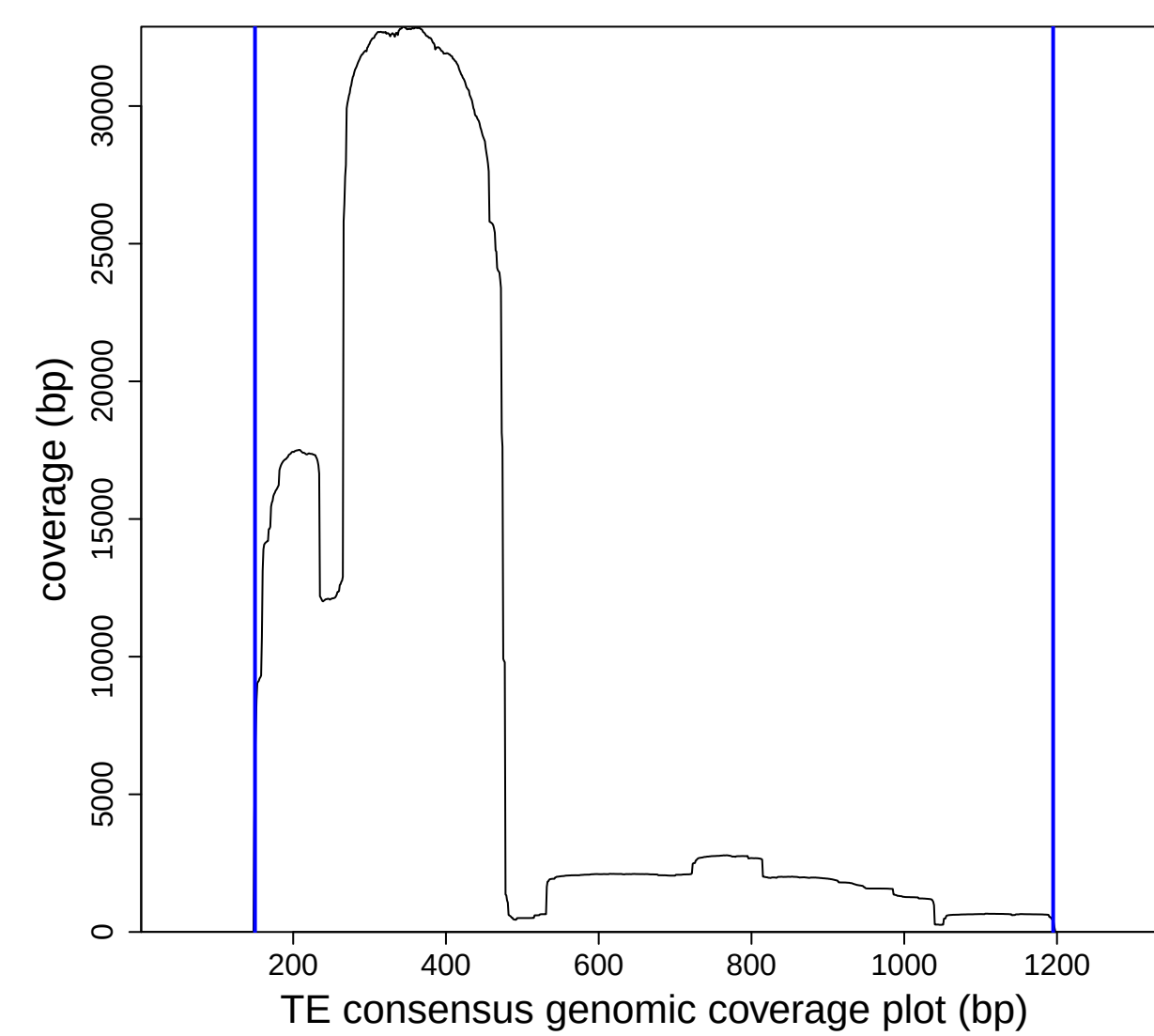
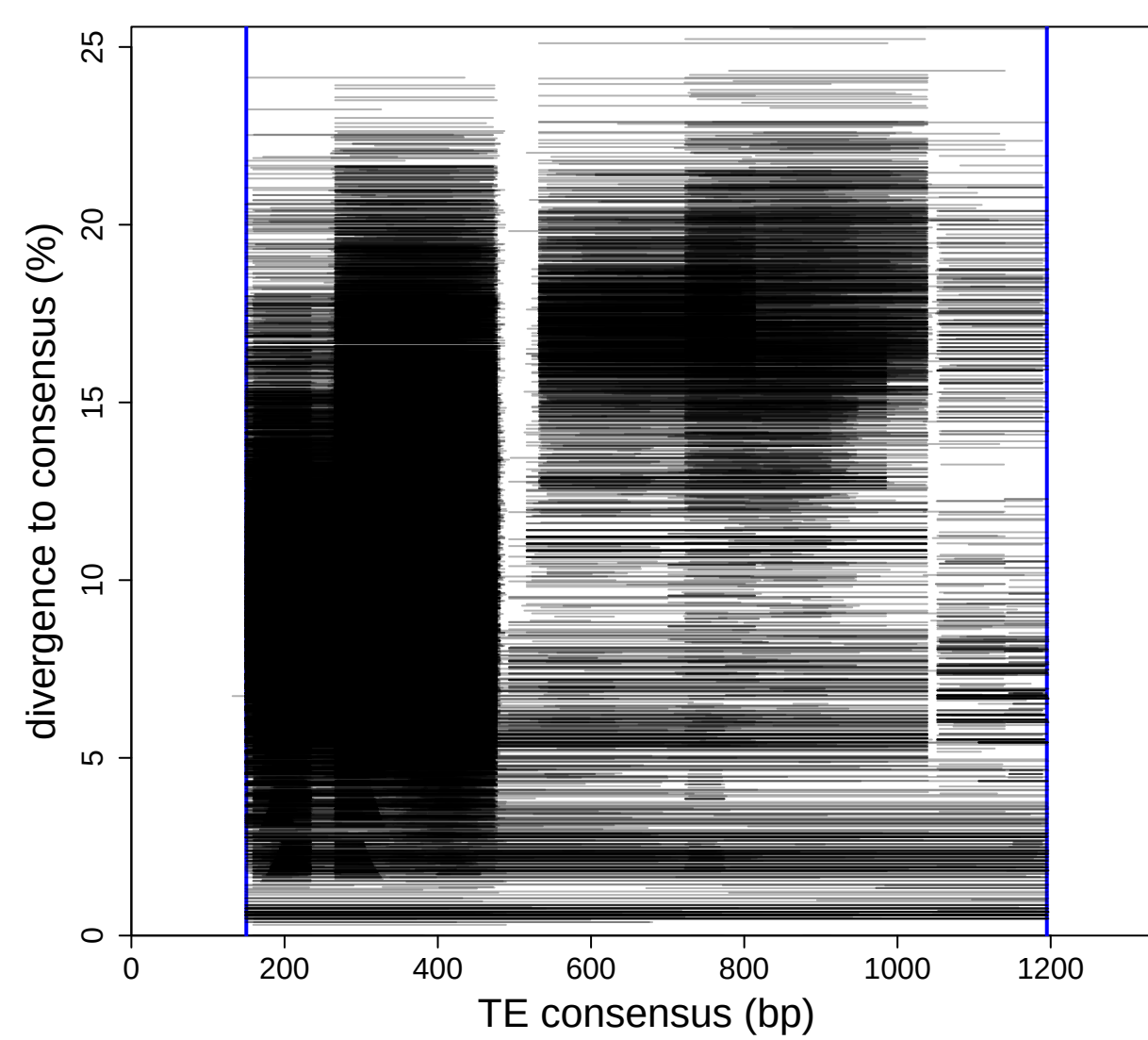
.l.b.bed uf.bed g_3.bed fm_1.bed 0 0 bcln.fa aln.fa cl.fa gs.fa
size: 1045bp; fragments: 44973; full length: 211 (>=940.5bp)

After TEtrimmer 1045 bp



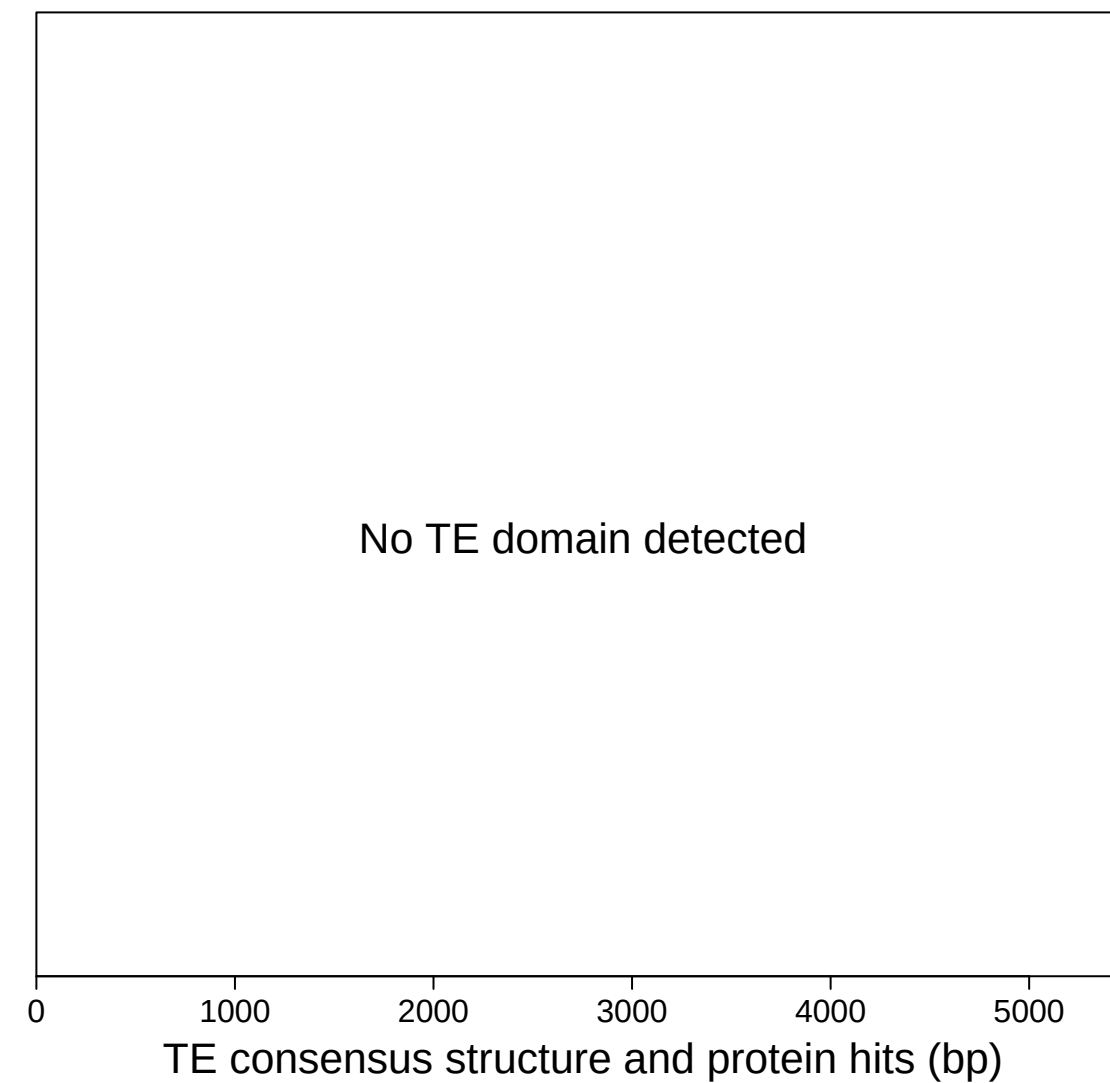
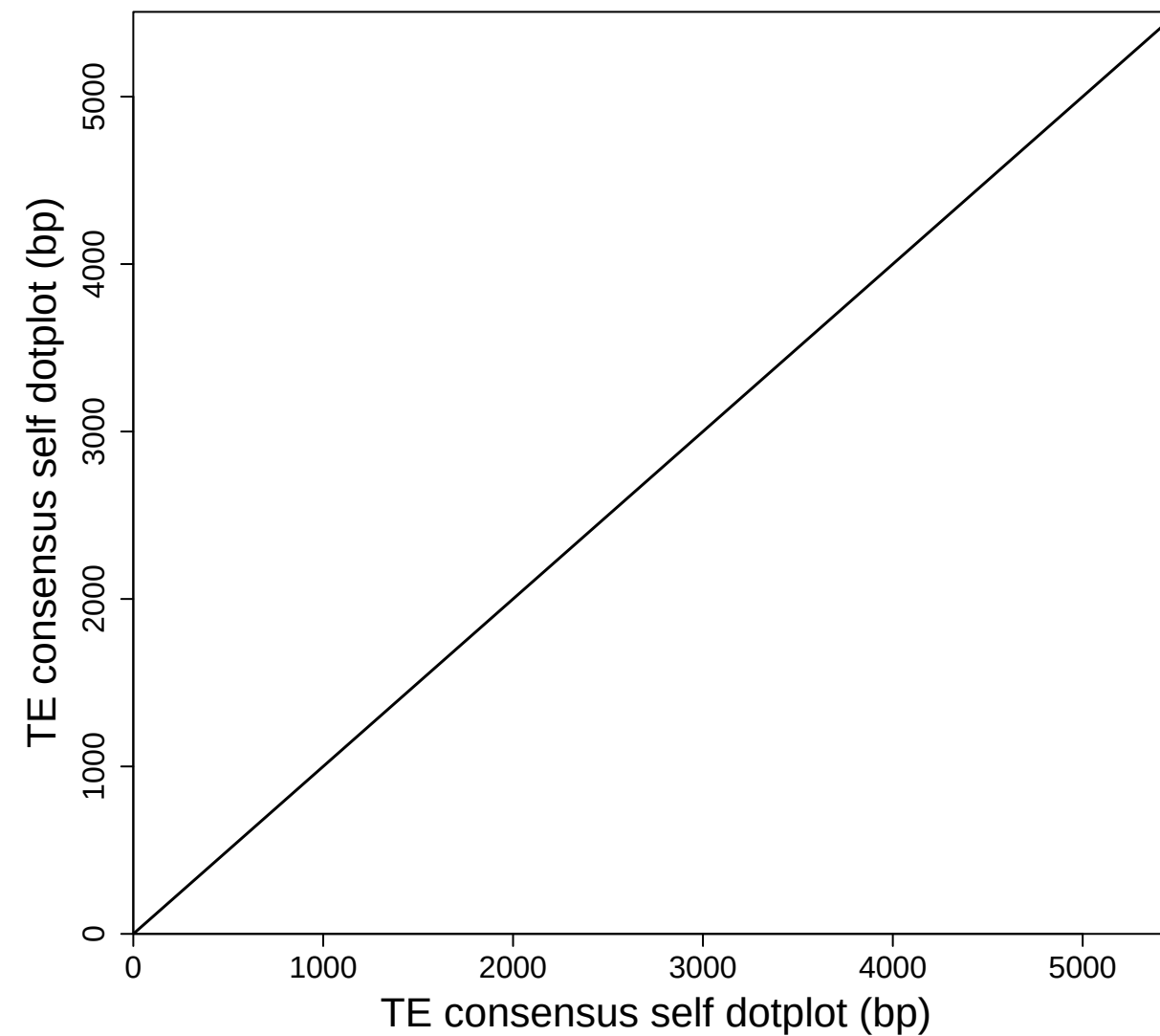
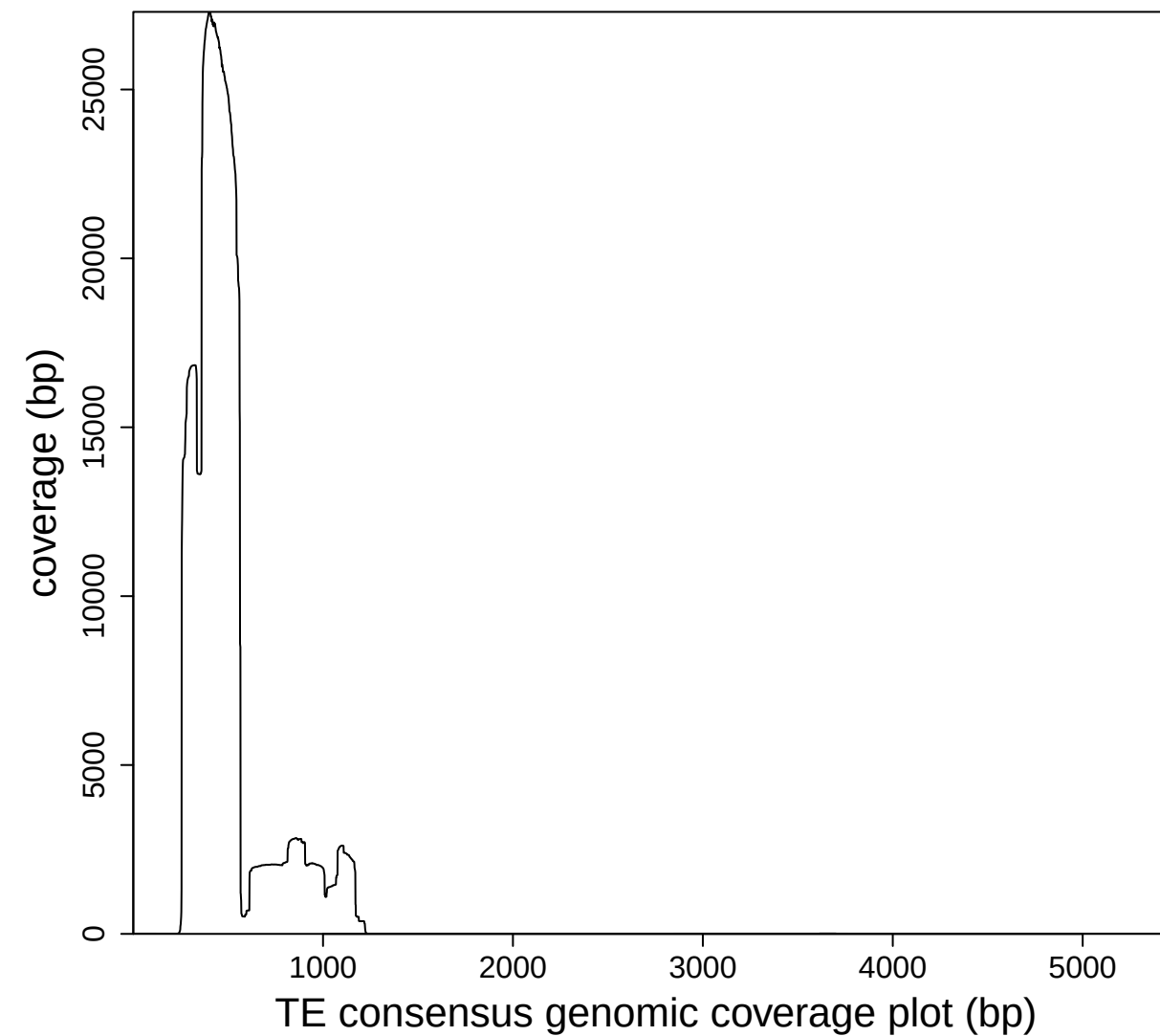
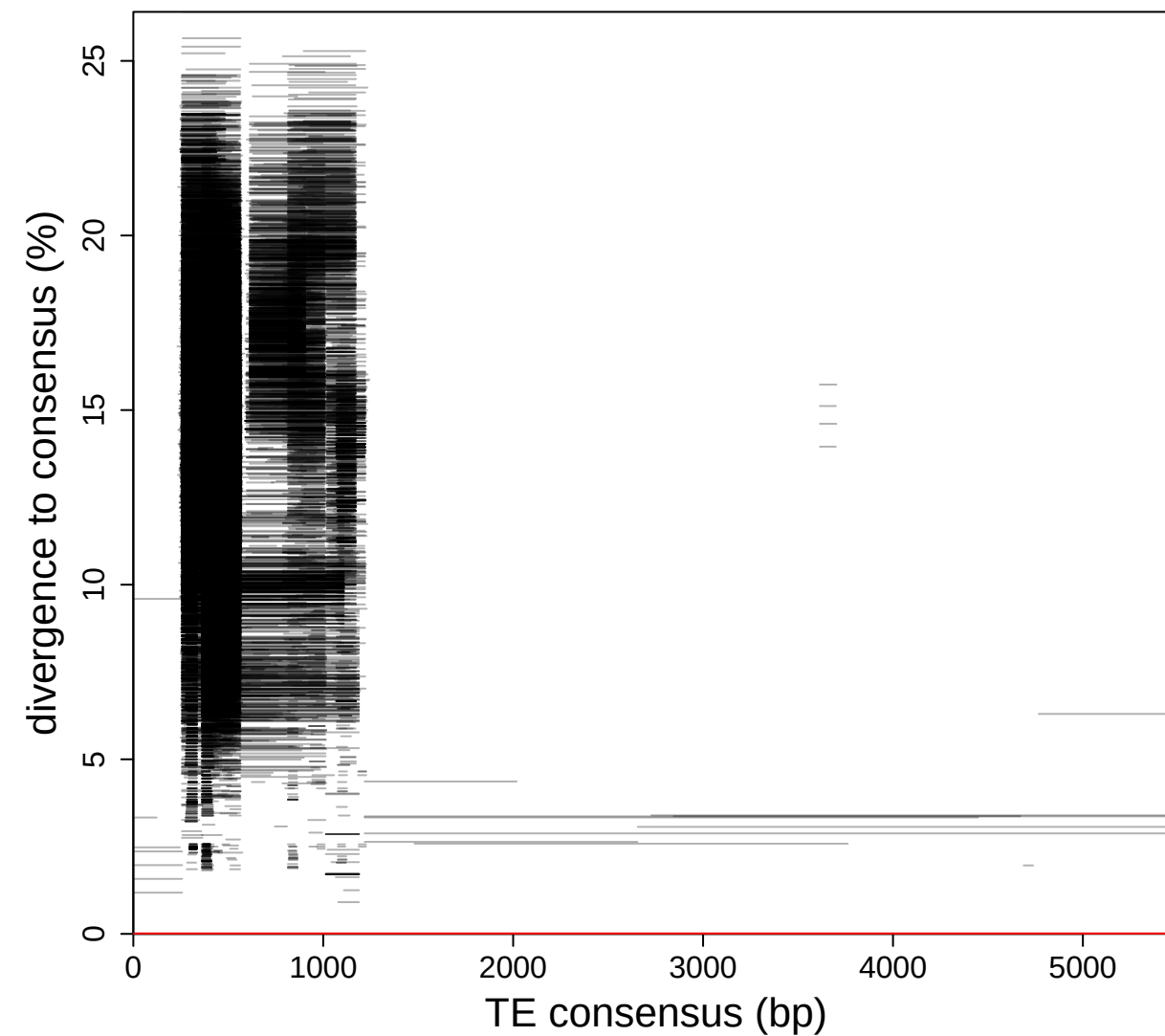
5.fasta.b.bed uf.bed g 3.bed fm 1.bed 0 0 bcln.fa aln.fa cl.fa
size: 1345bp; fragments: 45007; full length: 0 (>=1210.5bp)

After TEtrimmer Extended plot Blue lines are boundaries

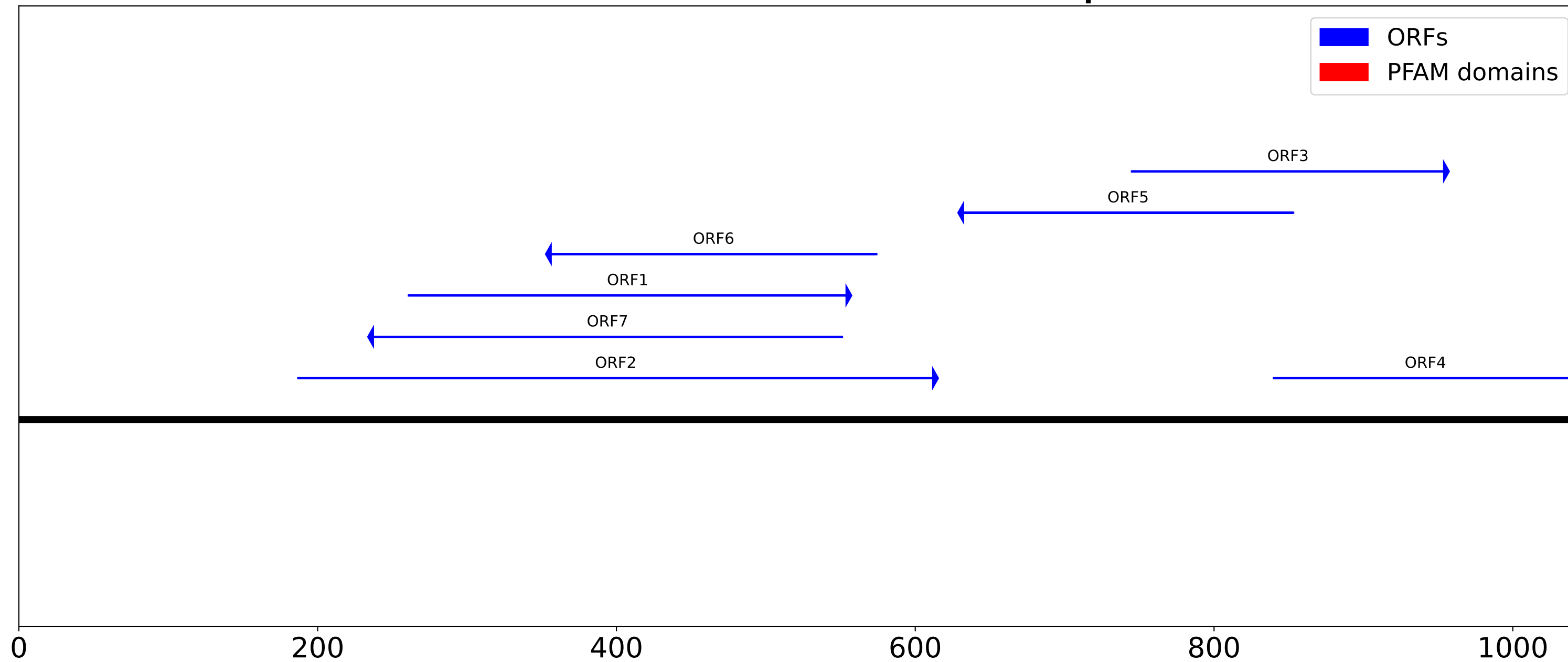


TE: ltr_1_family_1065
size: 5506bp; fragments: 36439; full length: 1 (≥ 4955.4 bp)

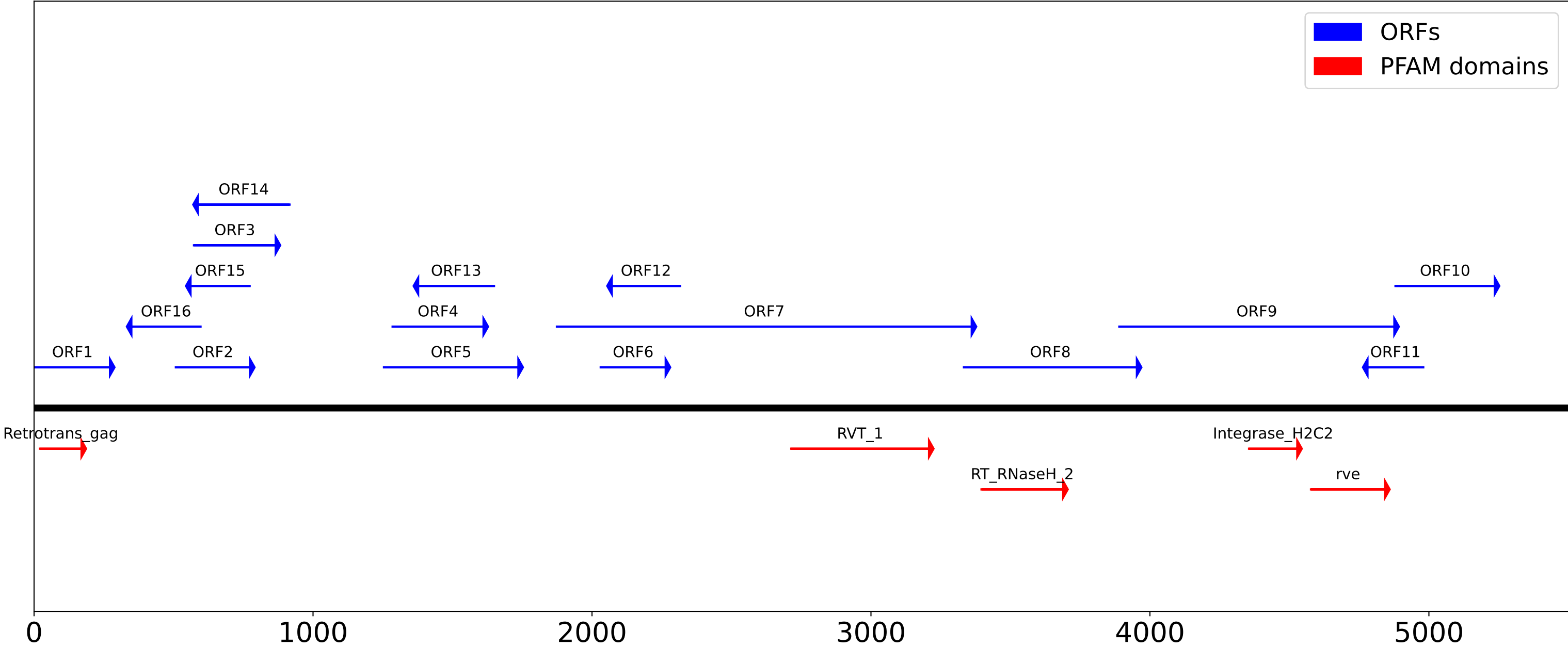
Before TEtrimmer 5506 bp



After TEtrimmer ORF and PFAM domain plot



Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

