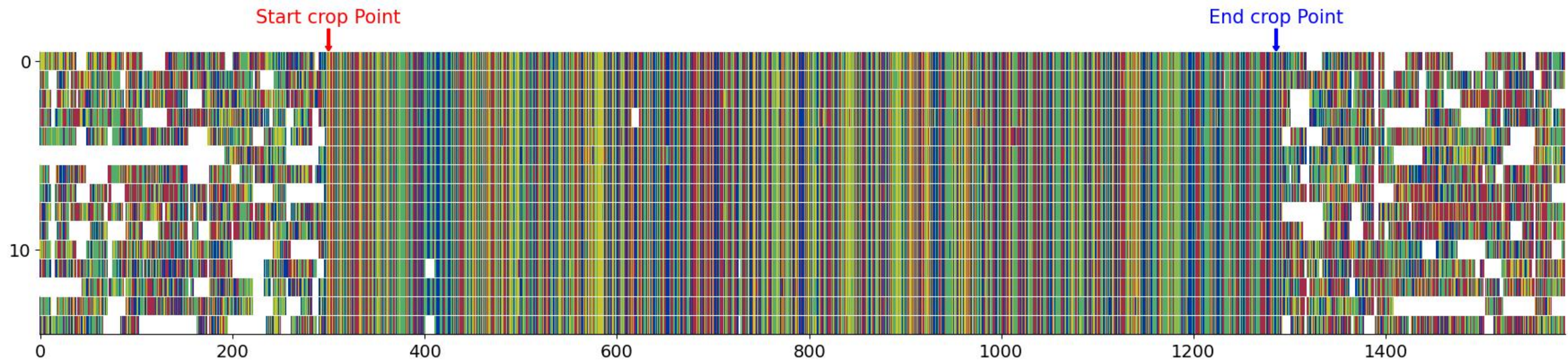
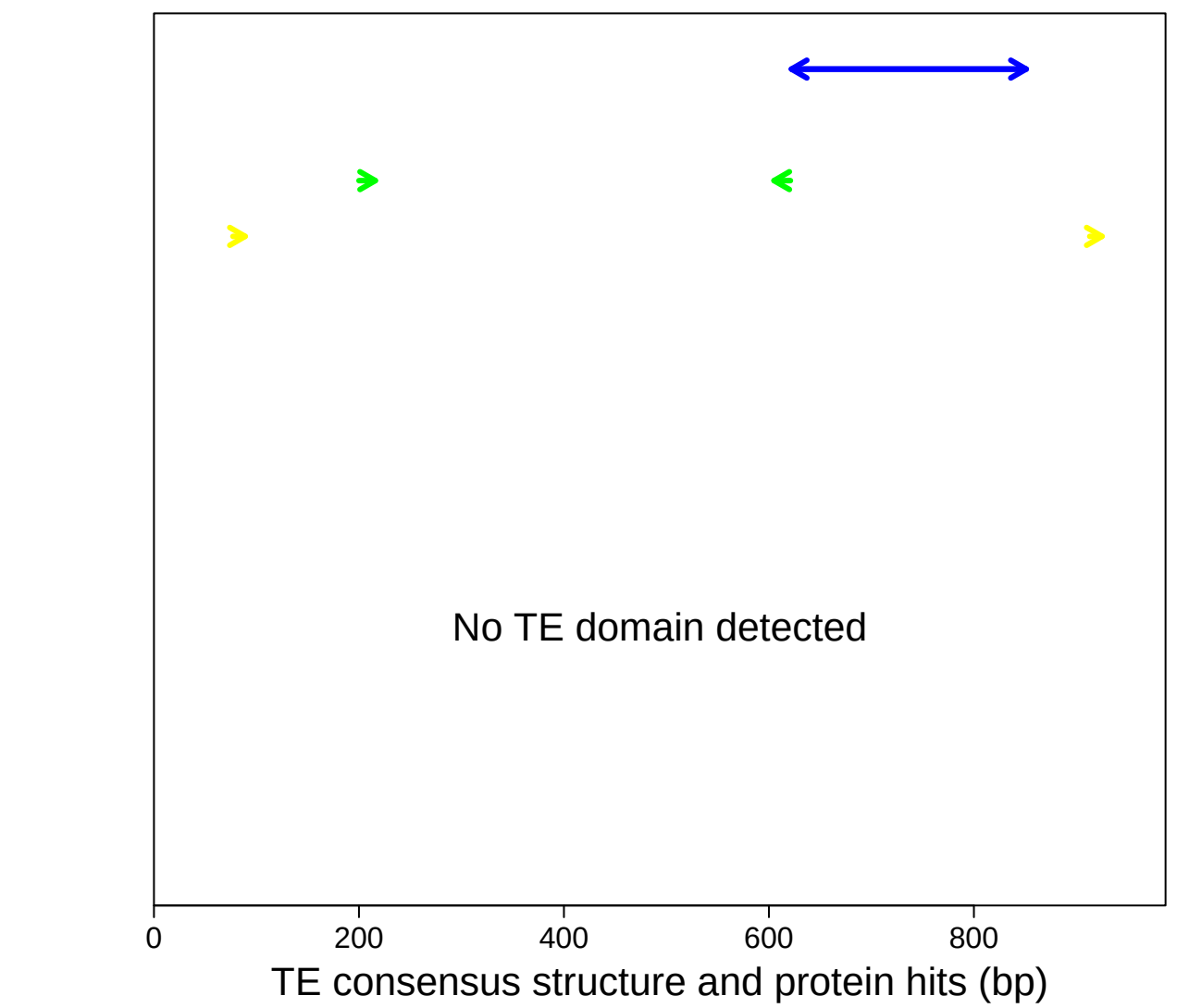
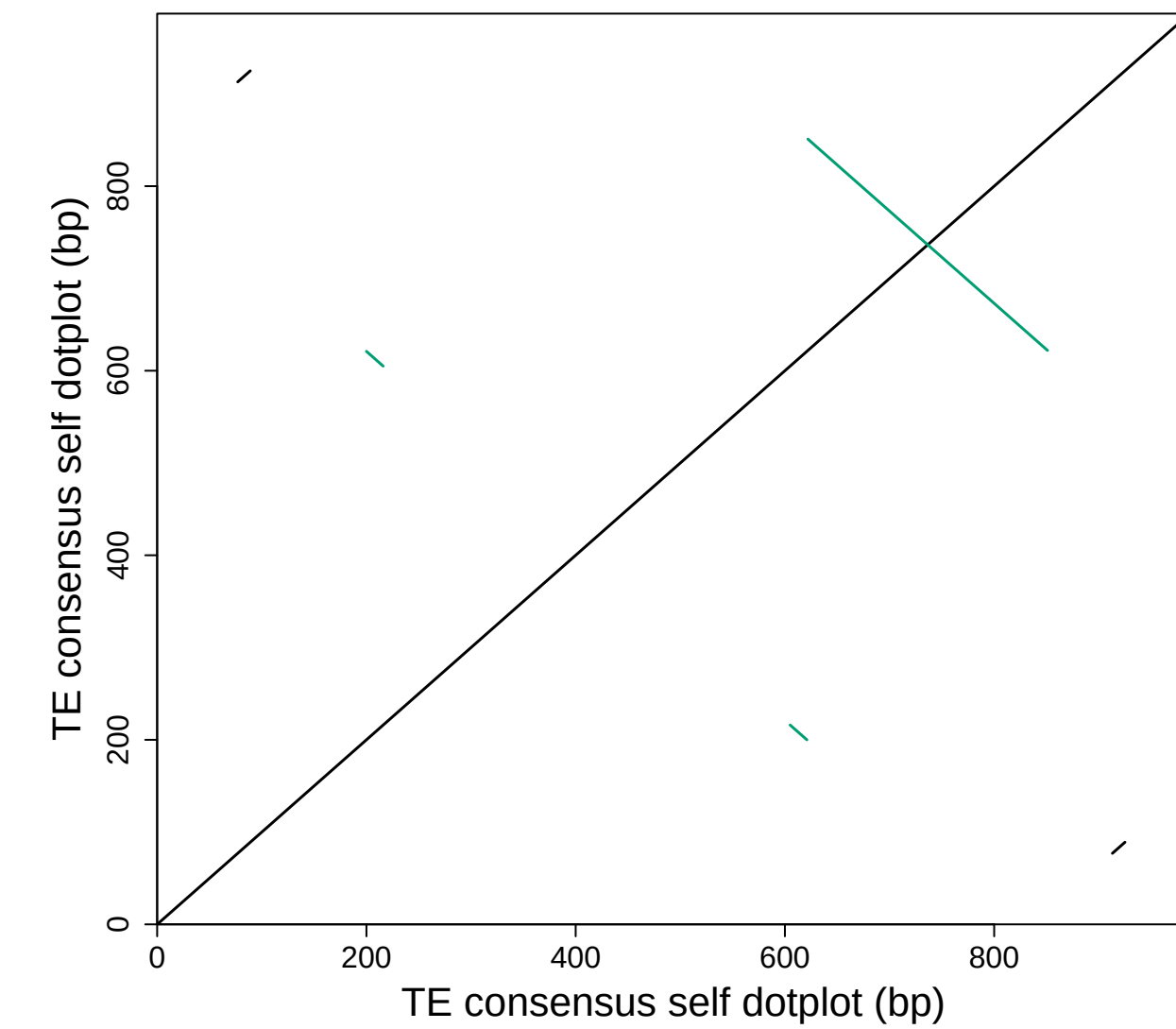
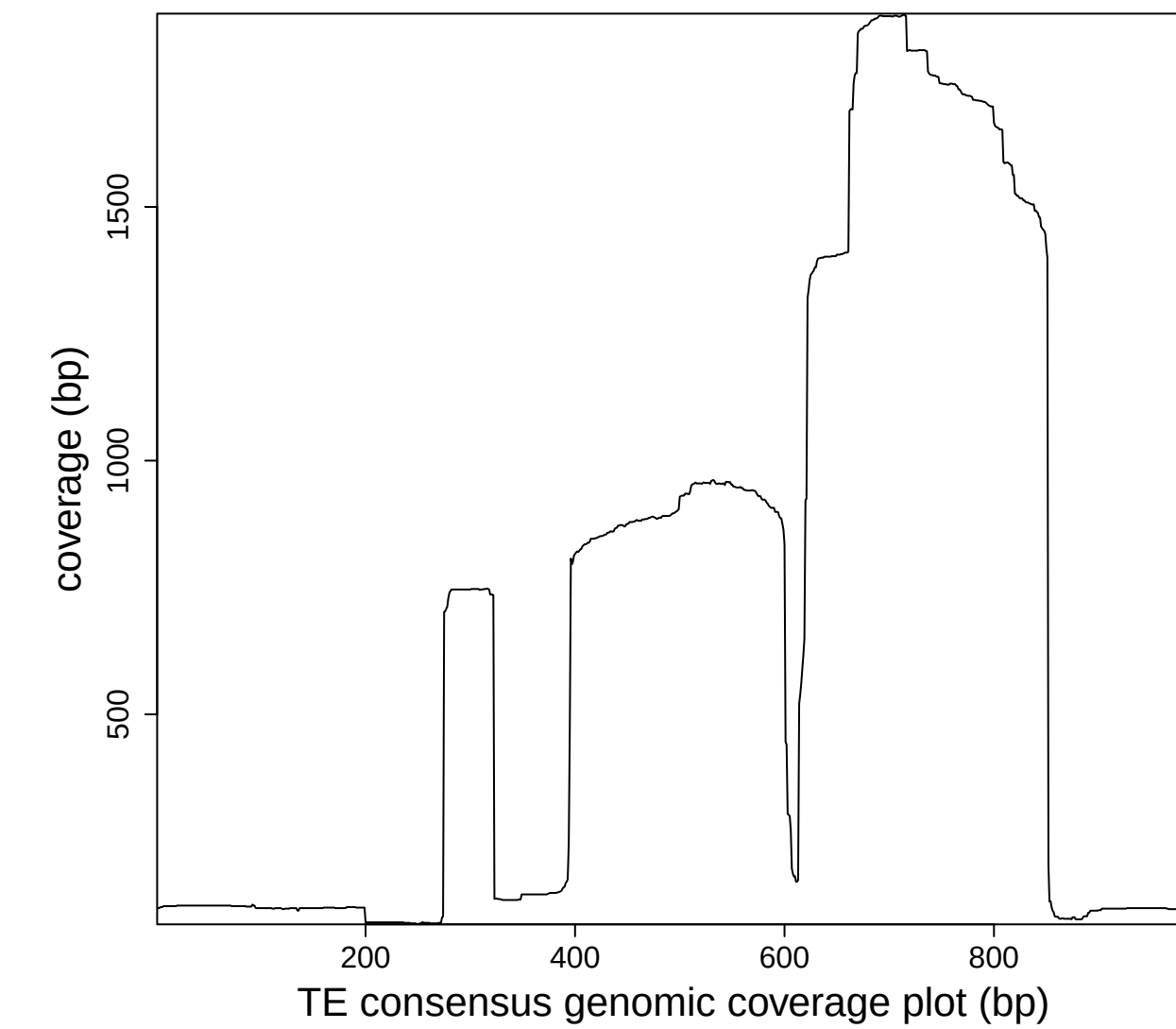
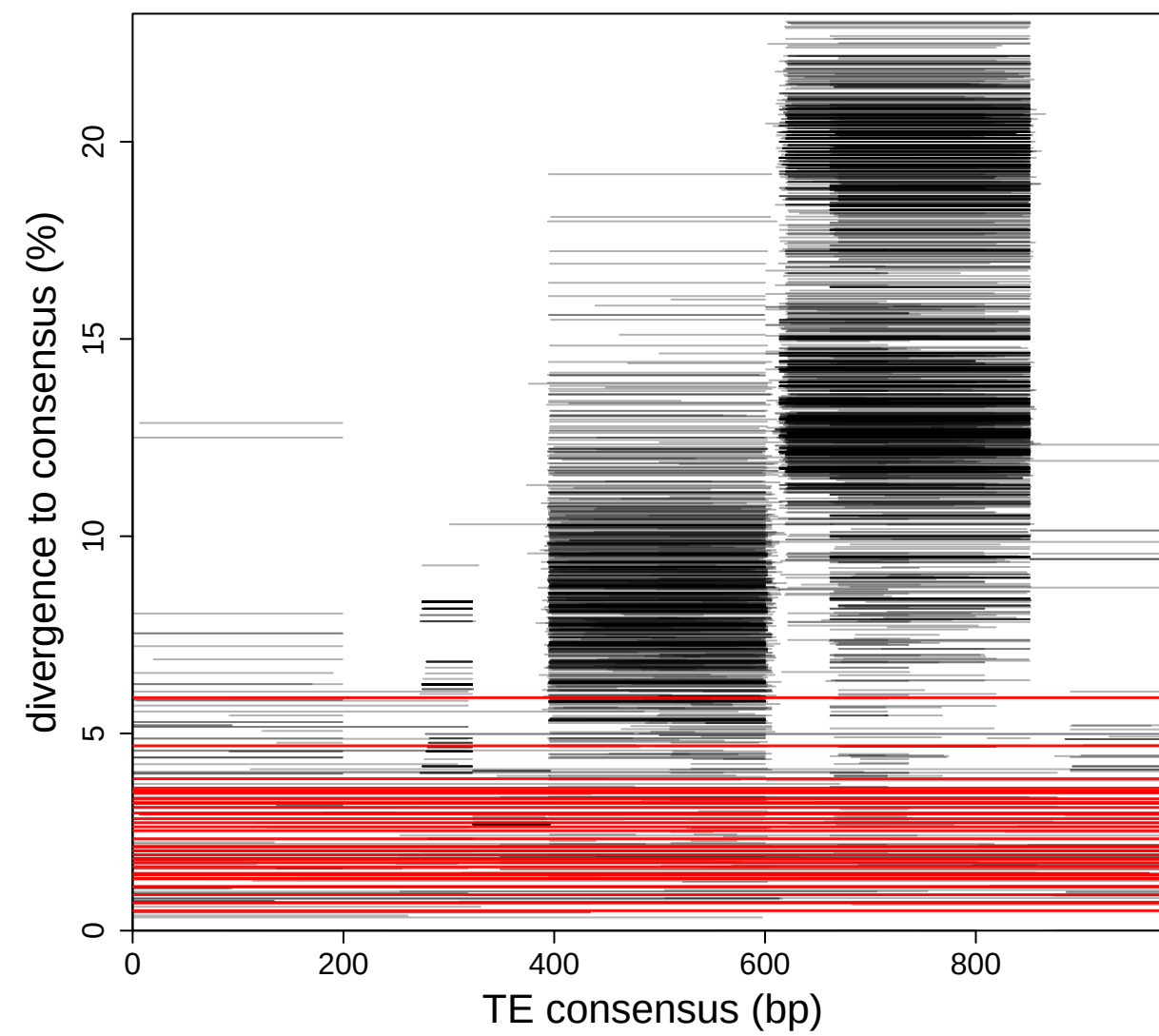




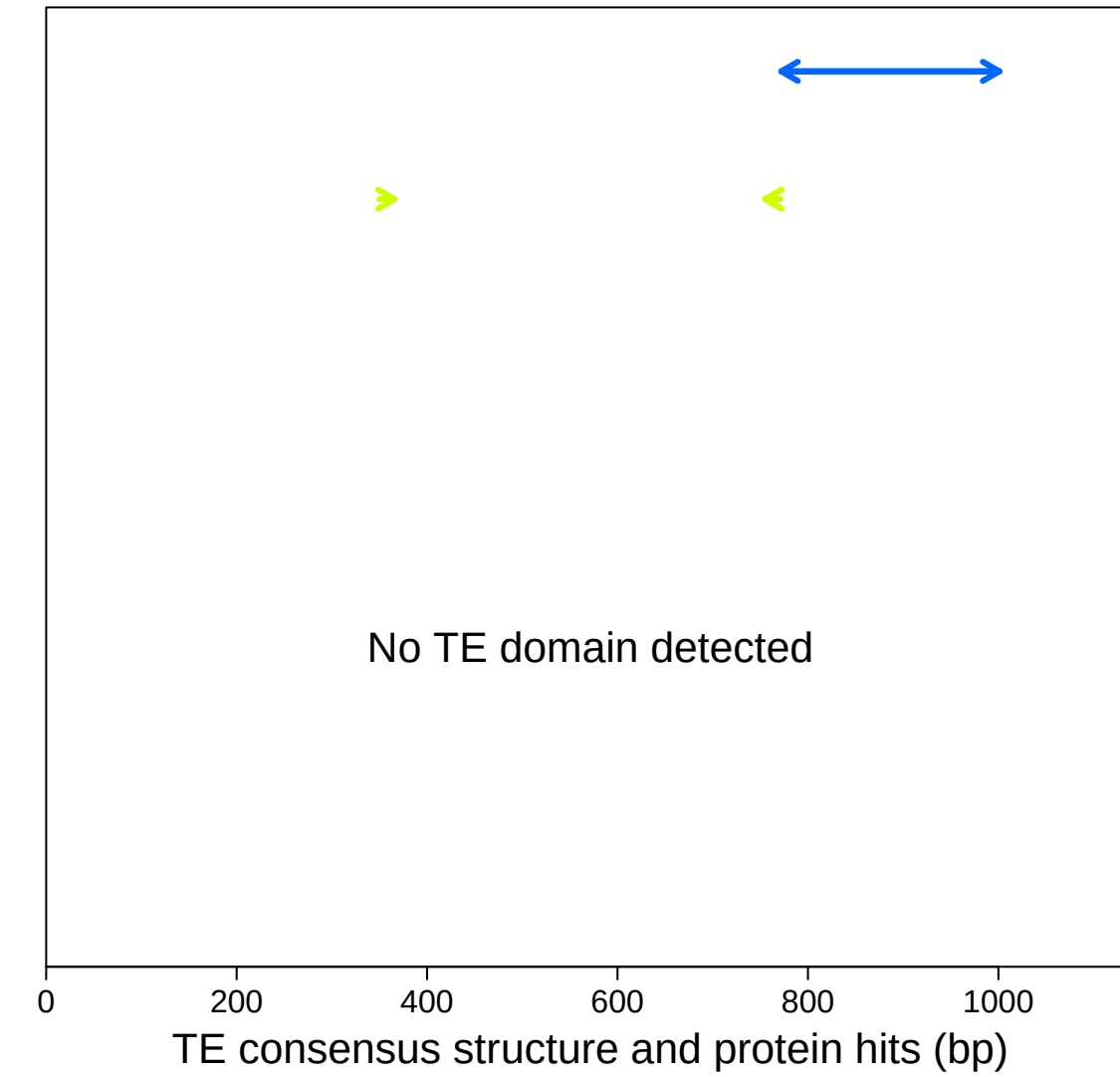
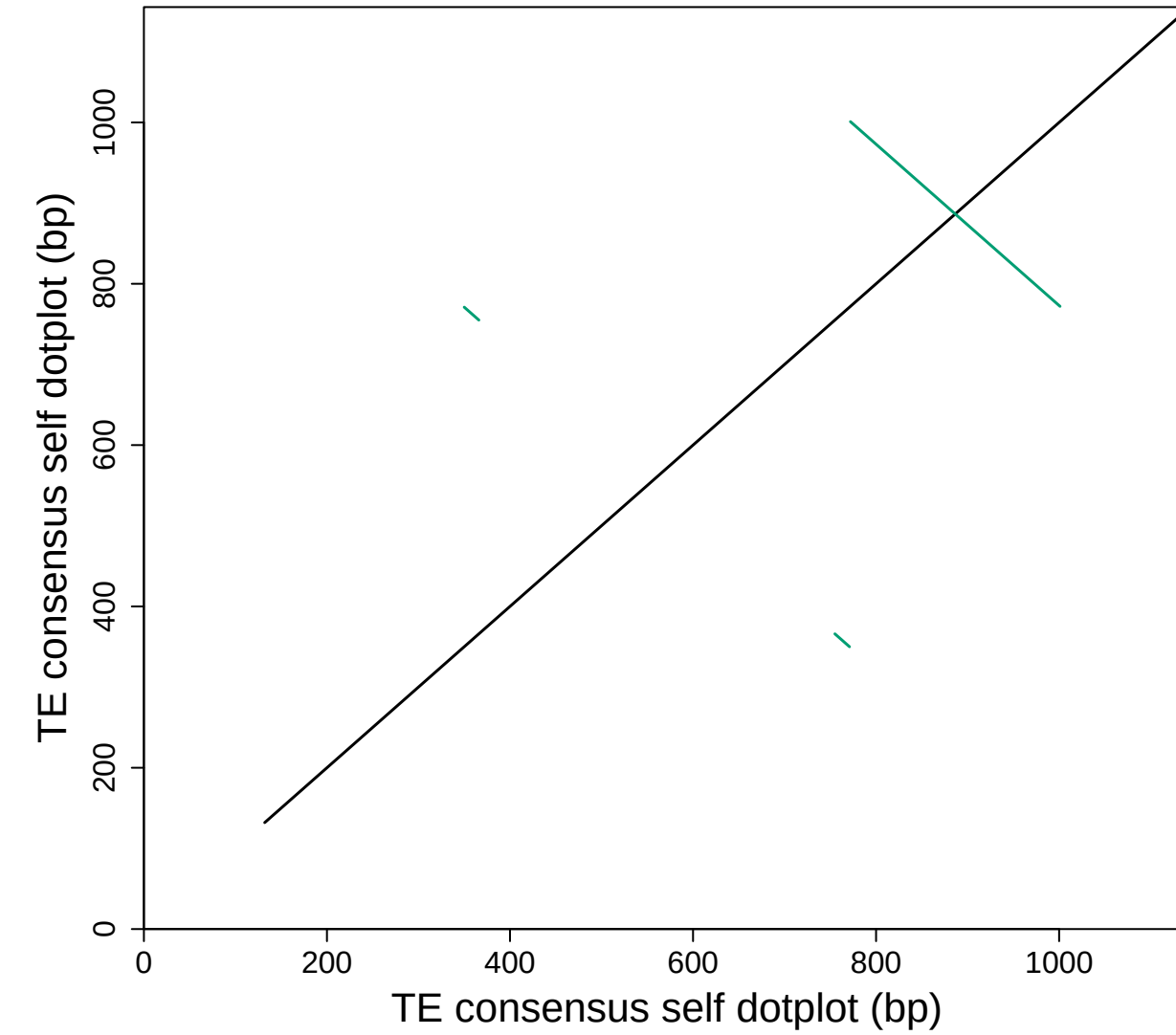
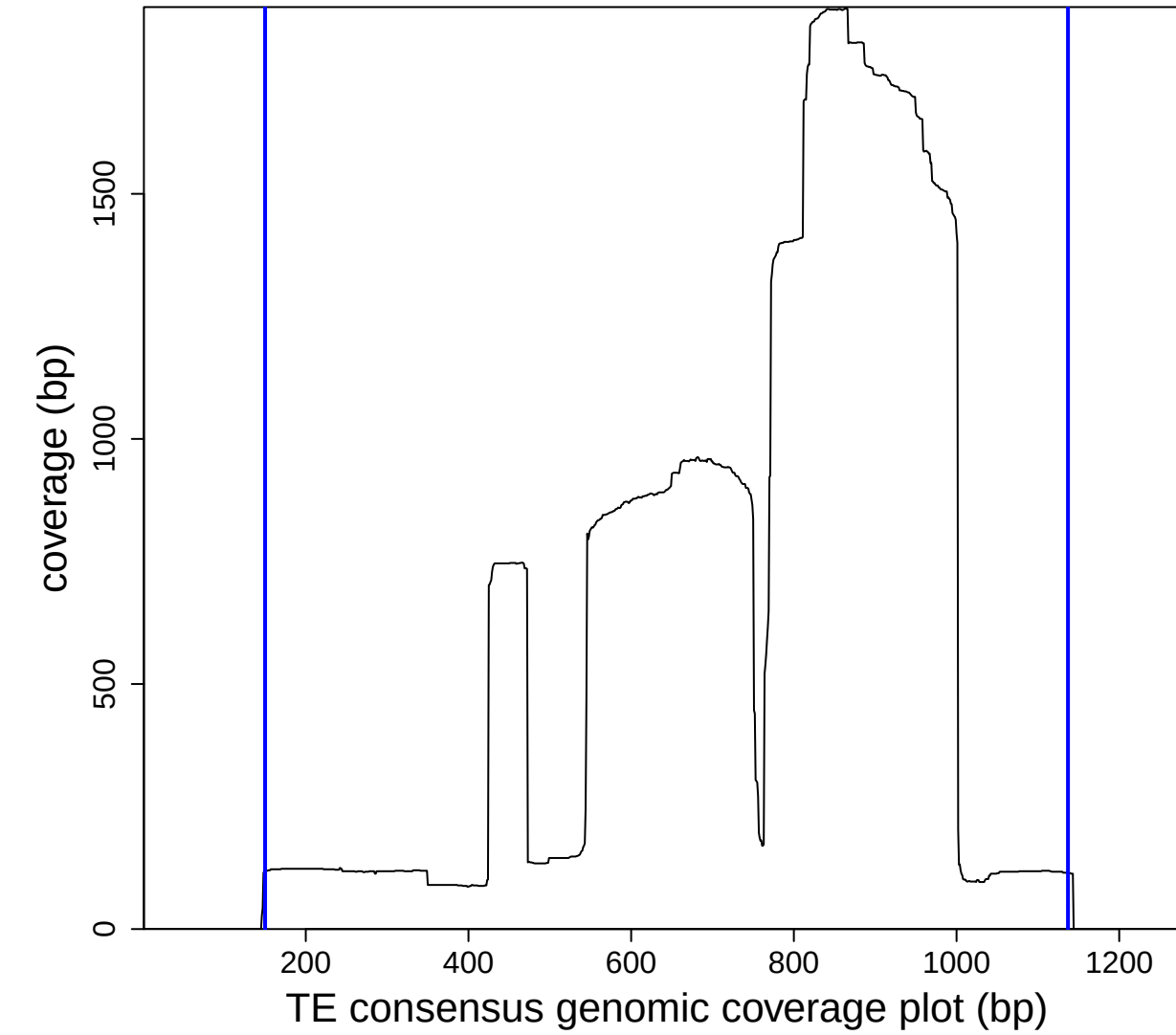
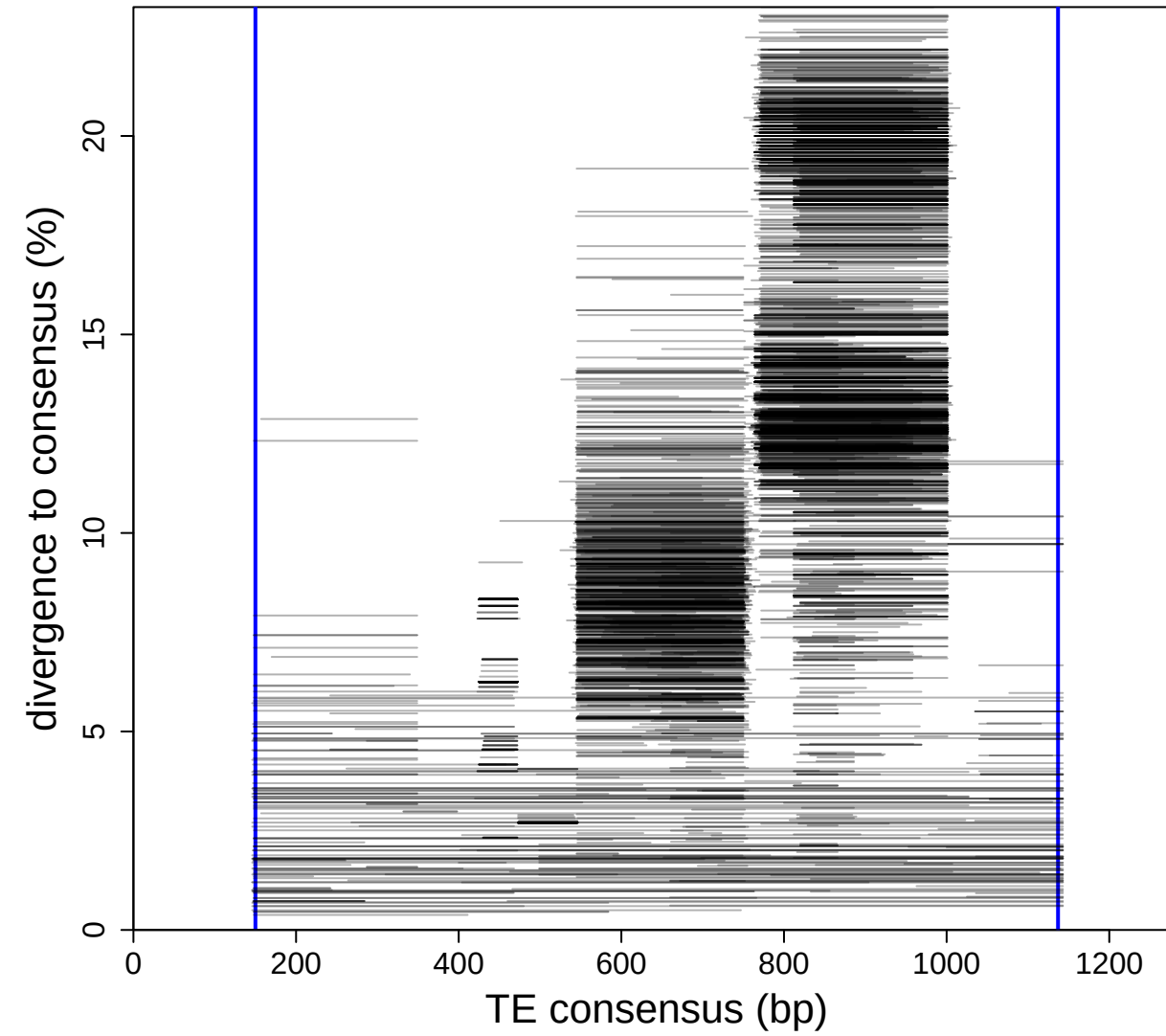
b.bed uf.bed g 3.bed fm 1.bed 0 0 bcln.fa aln.fa cl.fa gs.fa
size: 987bp; fragments: 3671; full length: 43 (>=888.3bp)

After TEtrimmer 987 bp



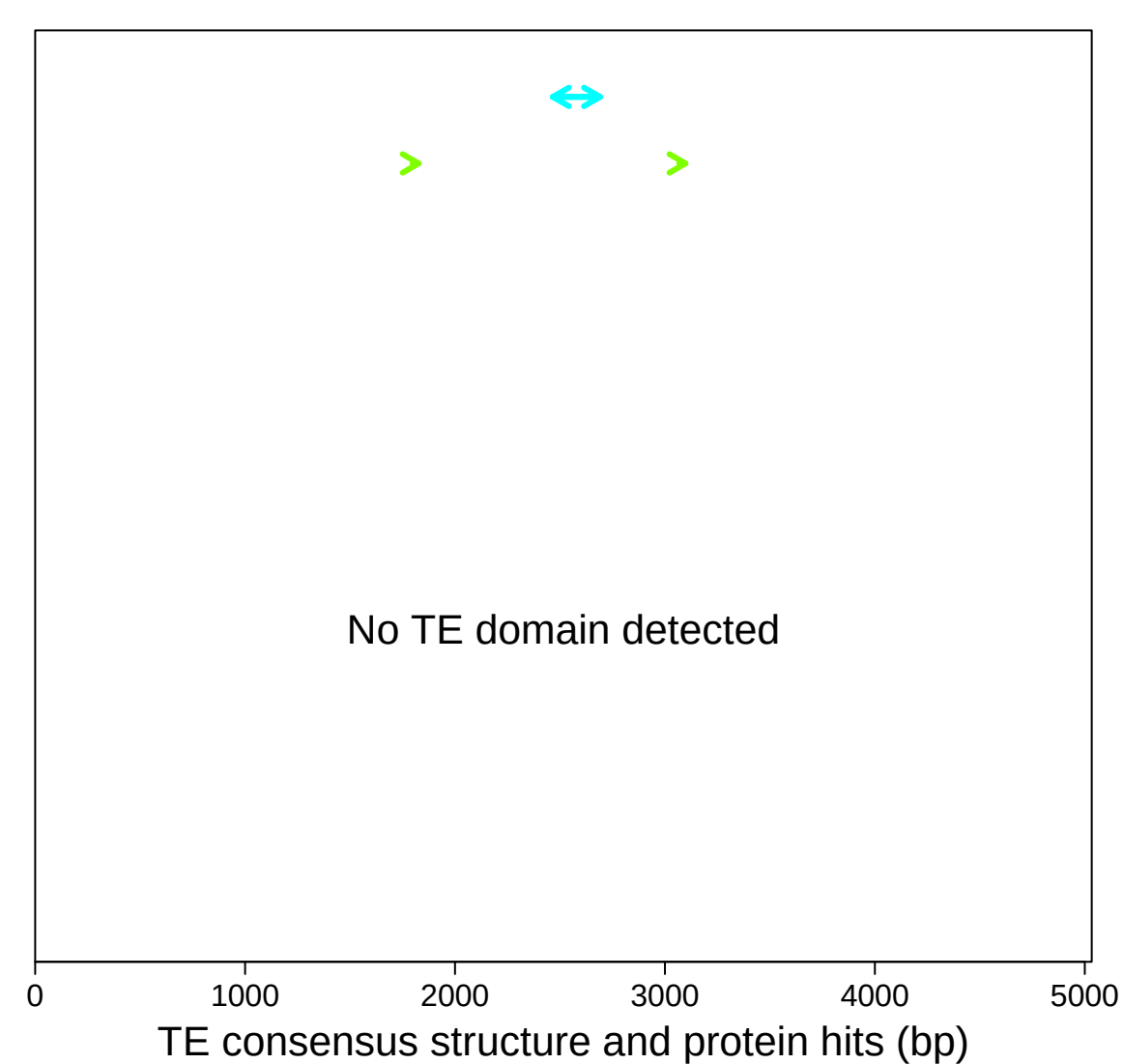
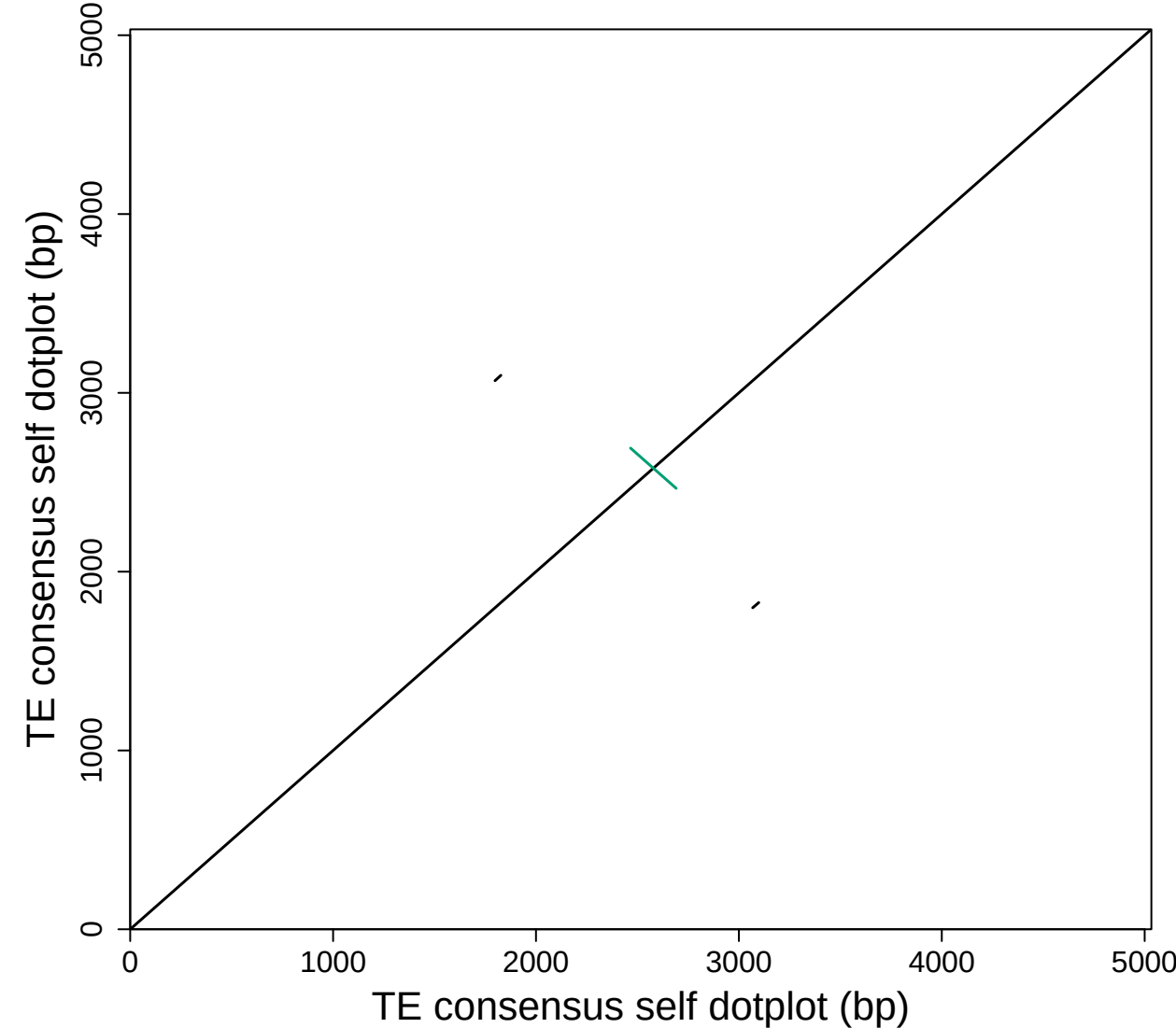
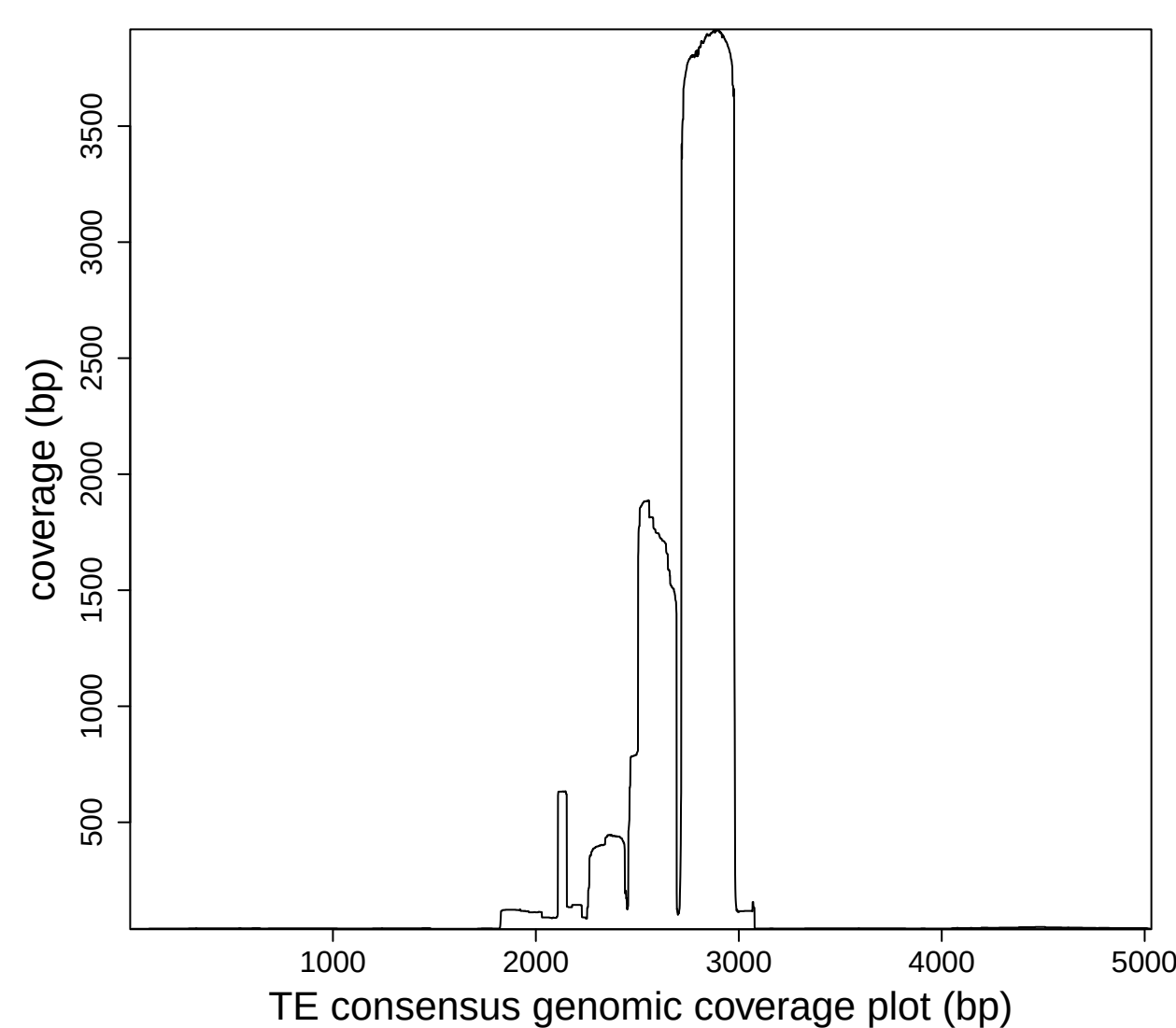
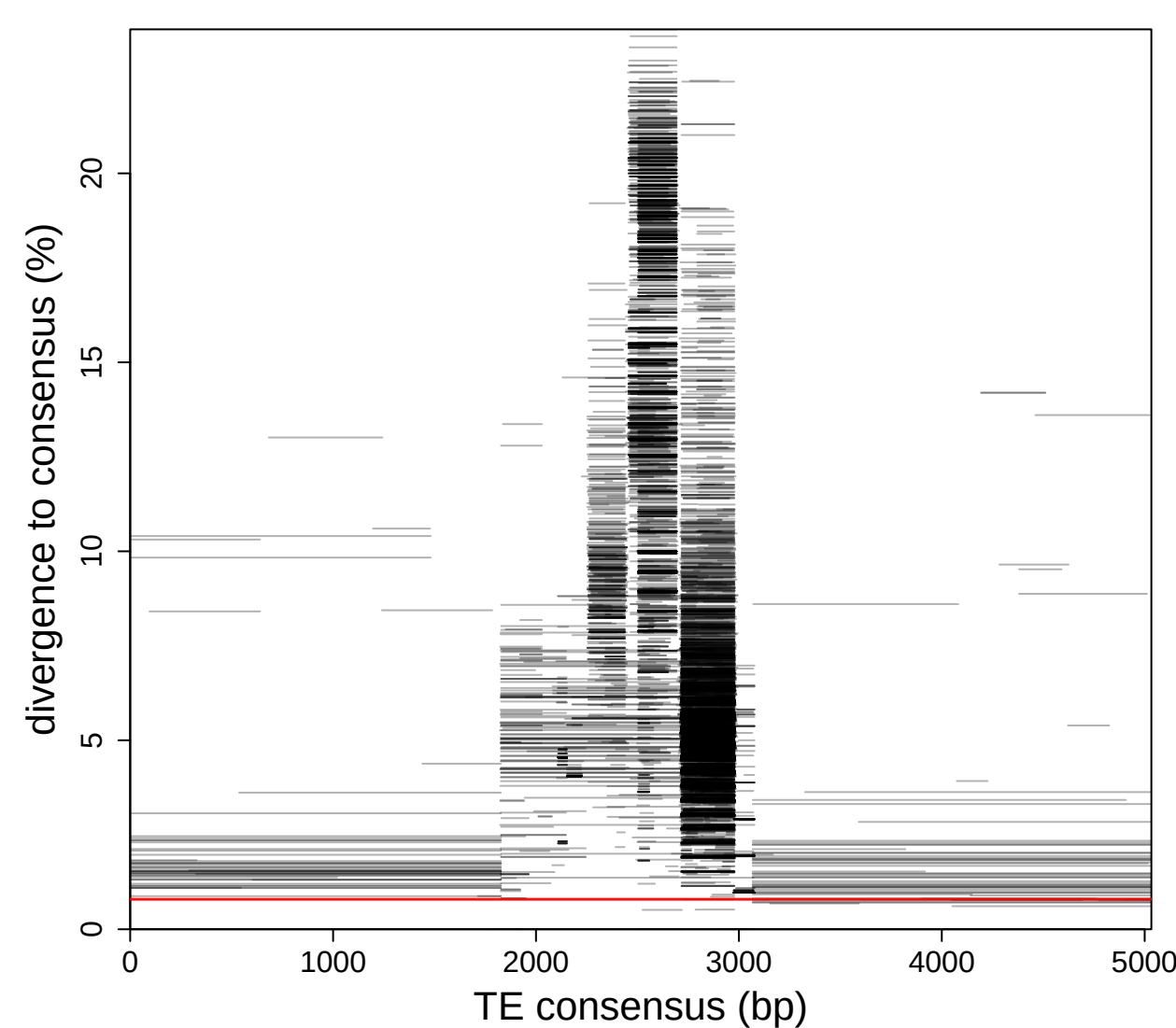
.fasta.b.bed_uf.bed_g_3.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa
size: 1287bp; fragments: 3672; full length: 0 (>=1158.3bp)

After TEtrimmer Extended plot Blue lines are boundaries

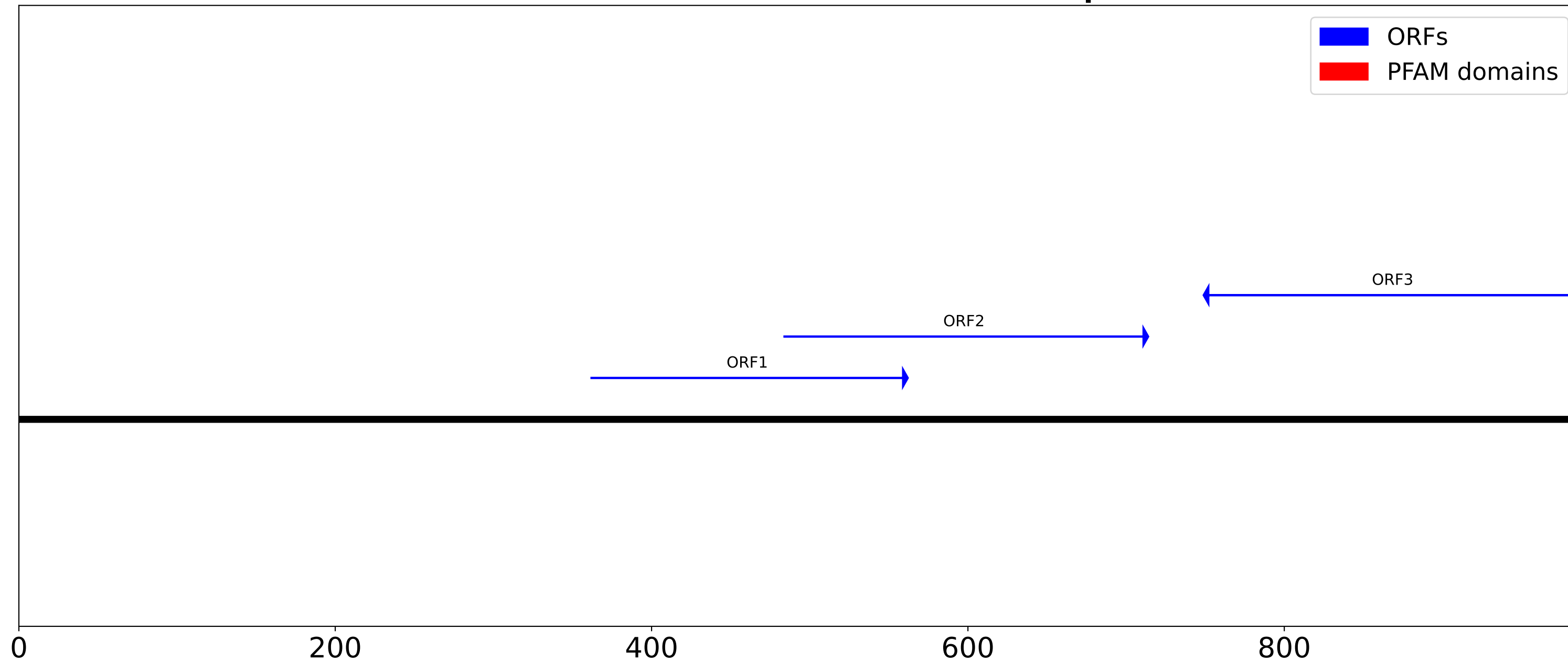


TE: ltr_1_family_957
size: 5033bp; fragments: 7444; full length: 1 (≥ 4529.7 bp)

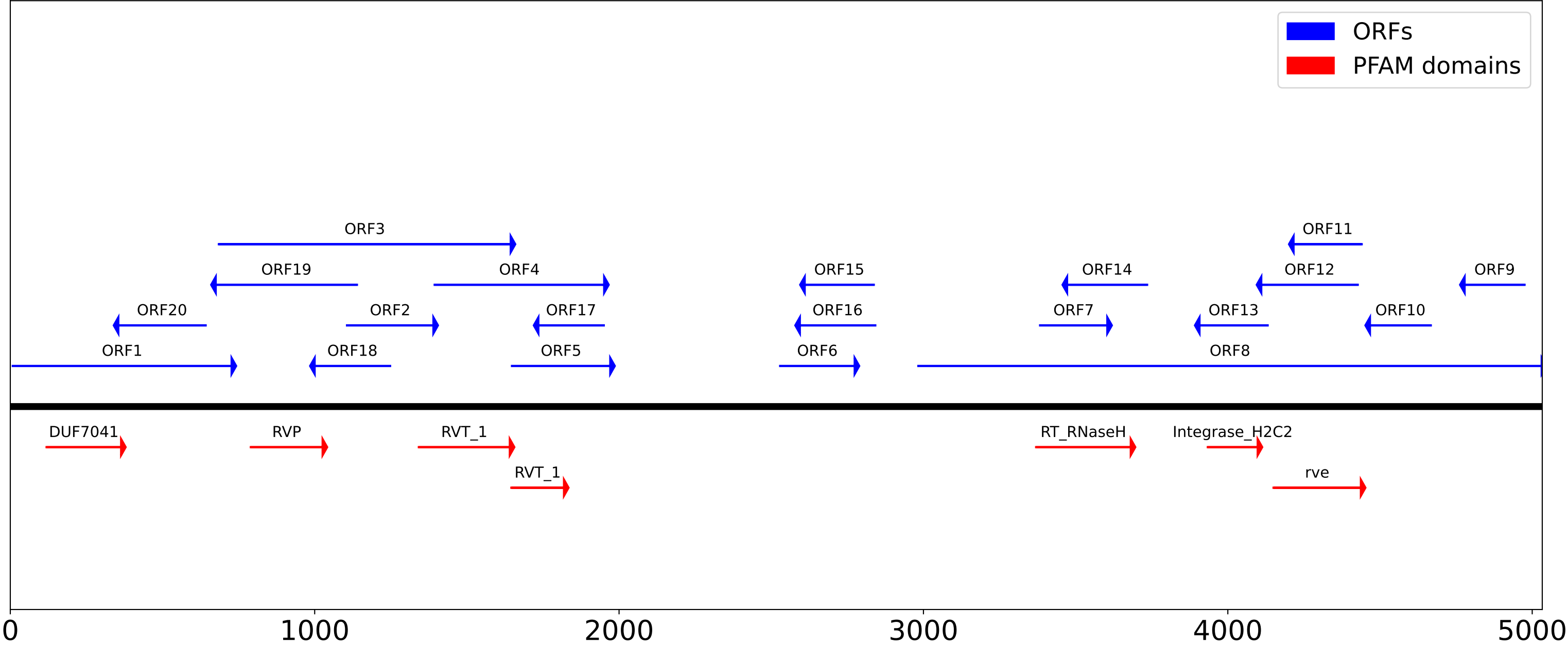
Before TEtrimmer 5033 bp



After TEtrimmer ORF and PFAM domain plot



Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

