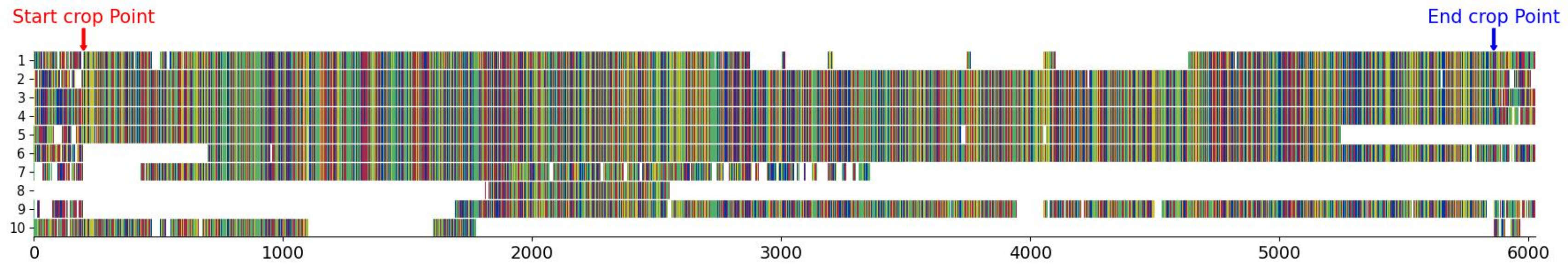


Start Crop Point

End Crop Point

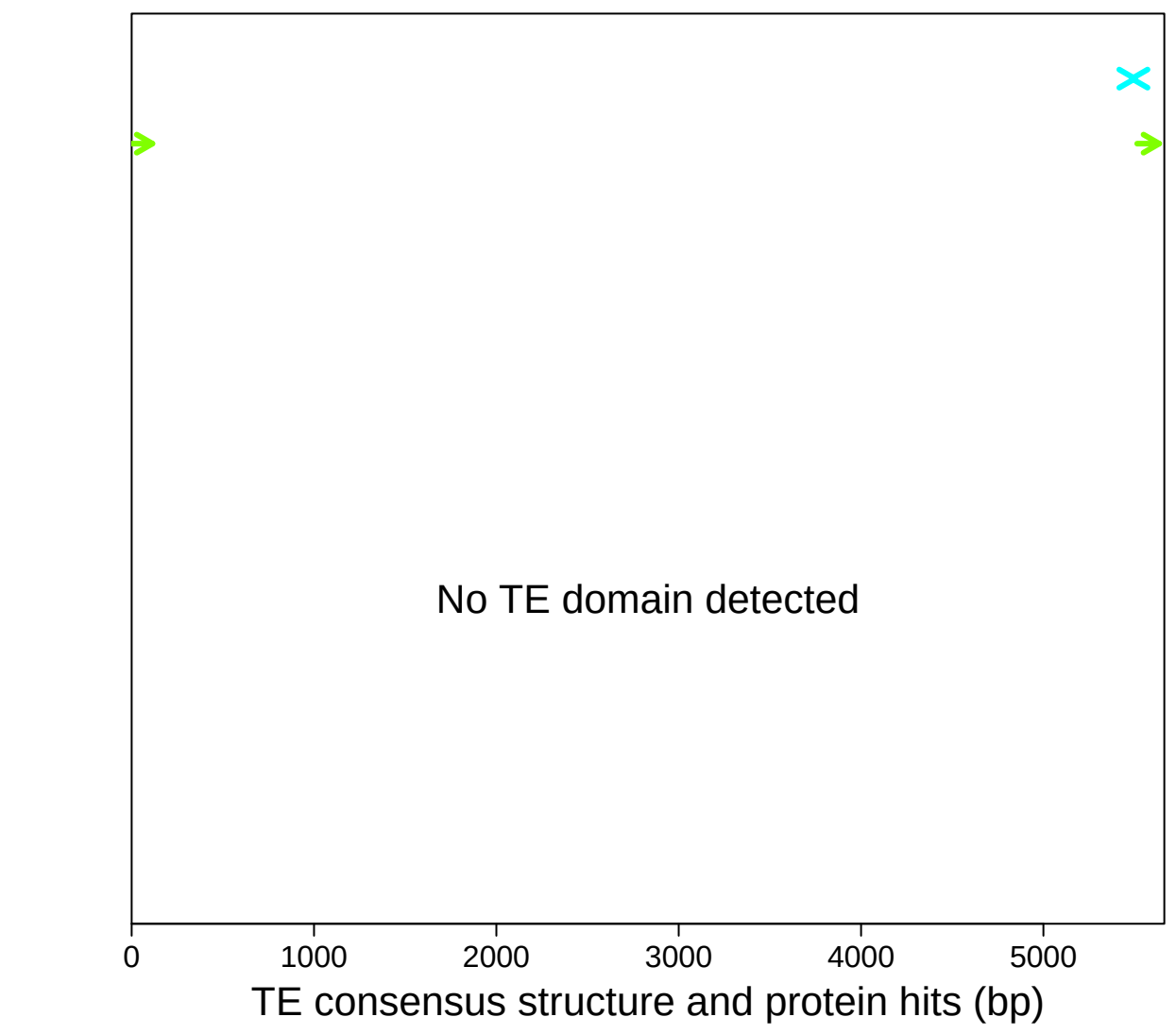
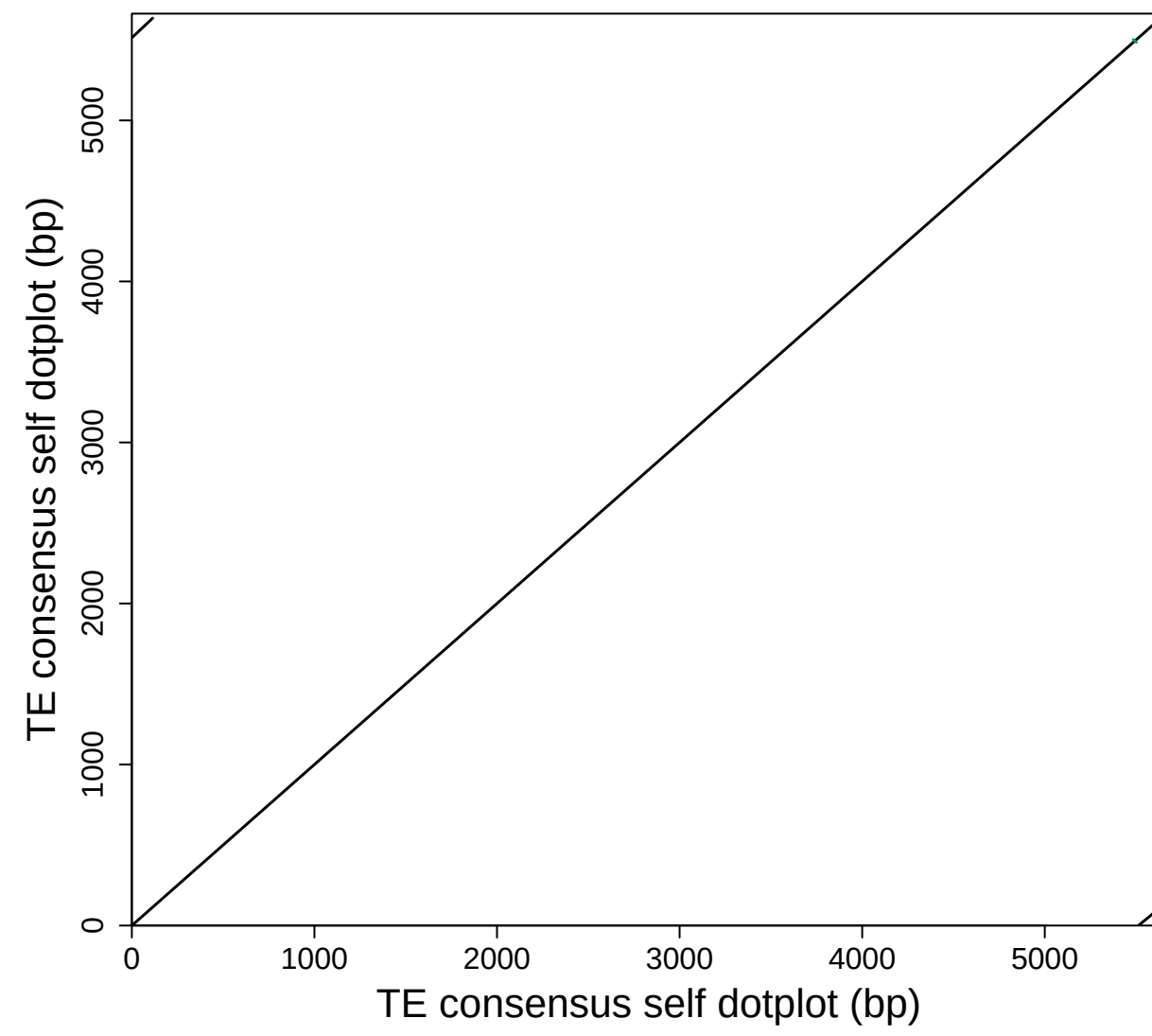
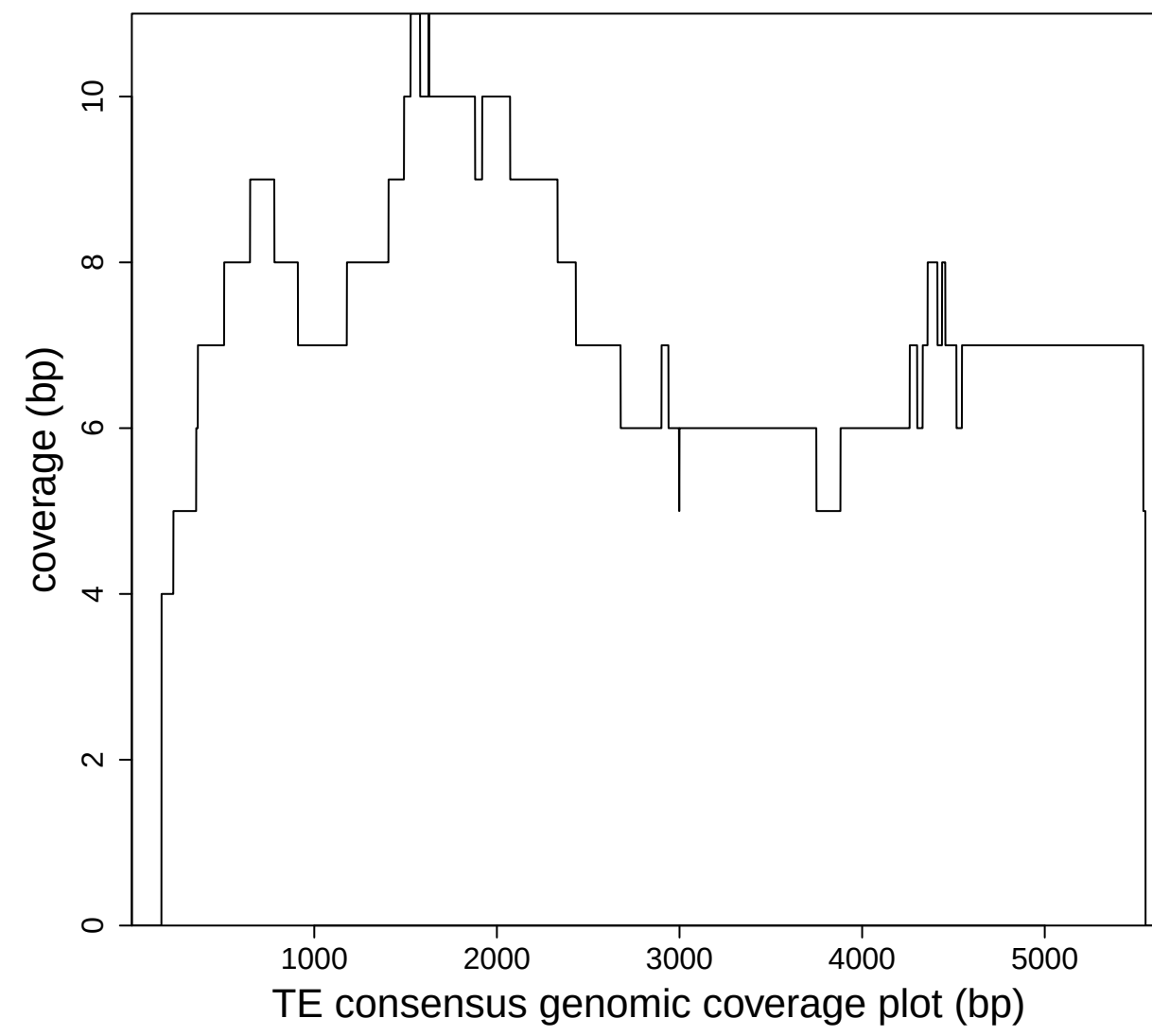
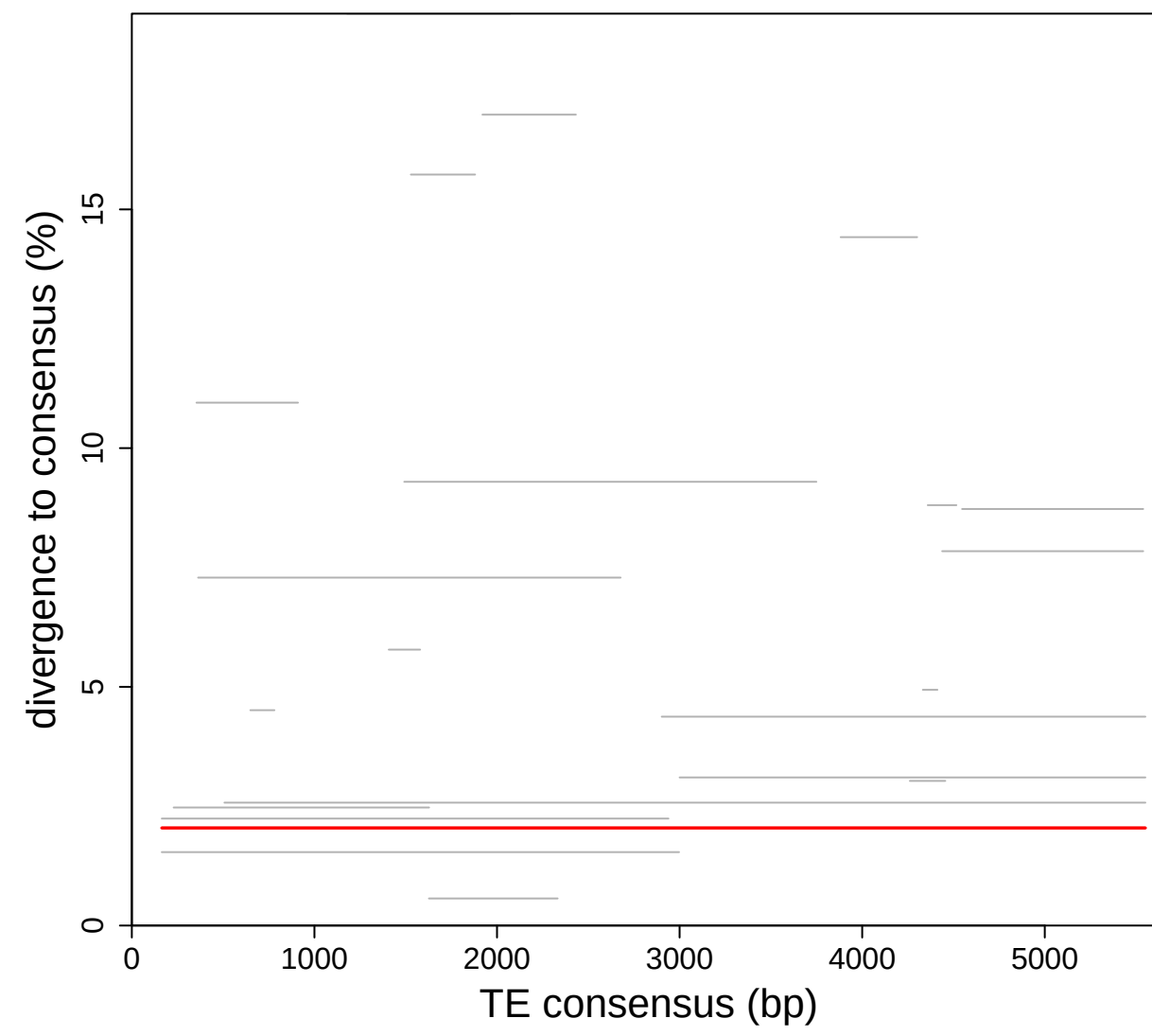
MSA length = 5663





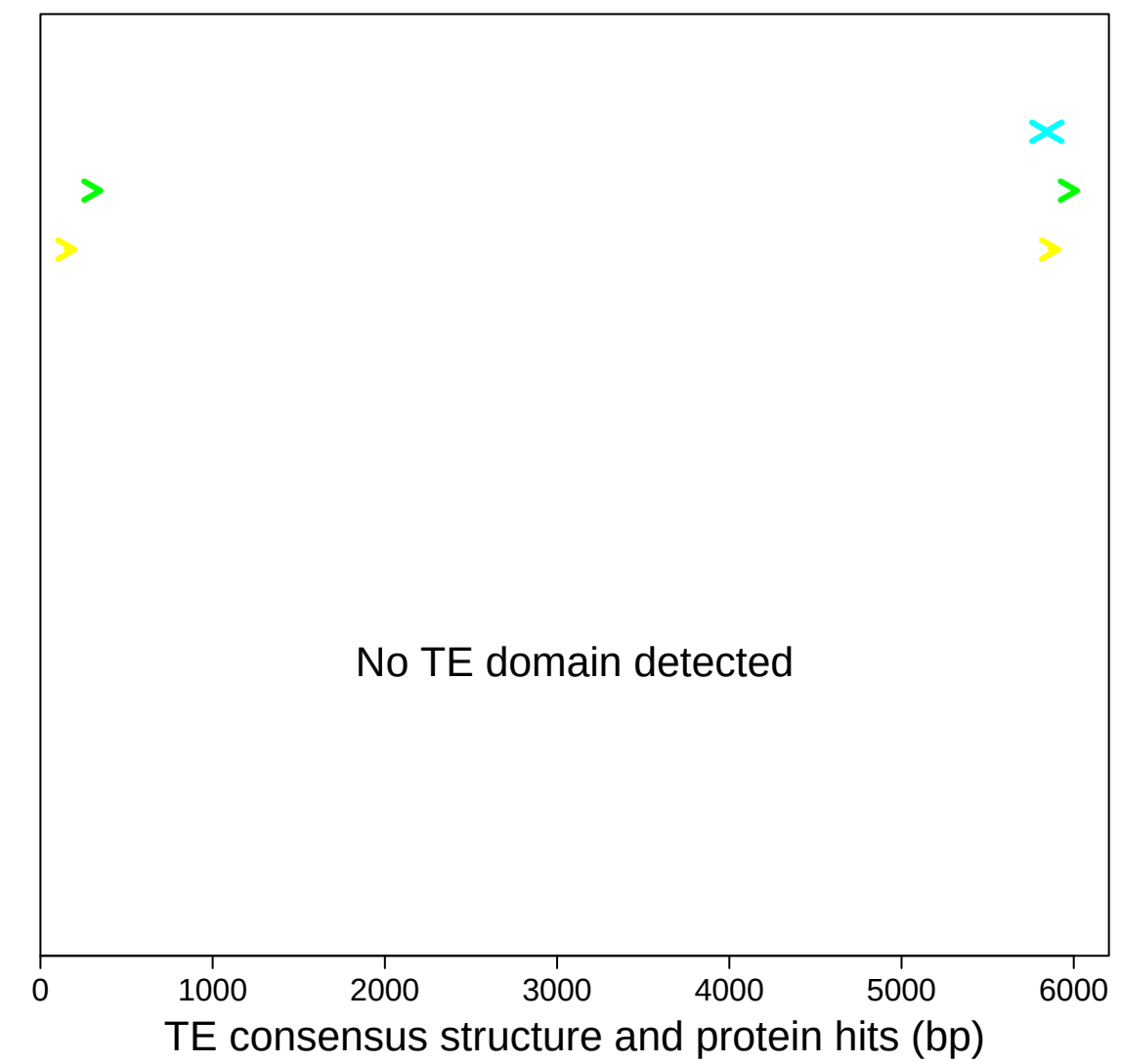
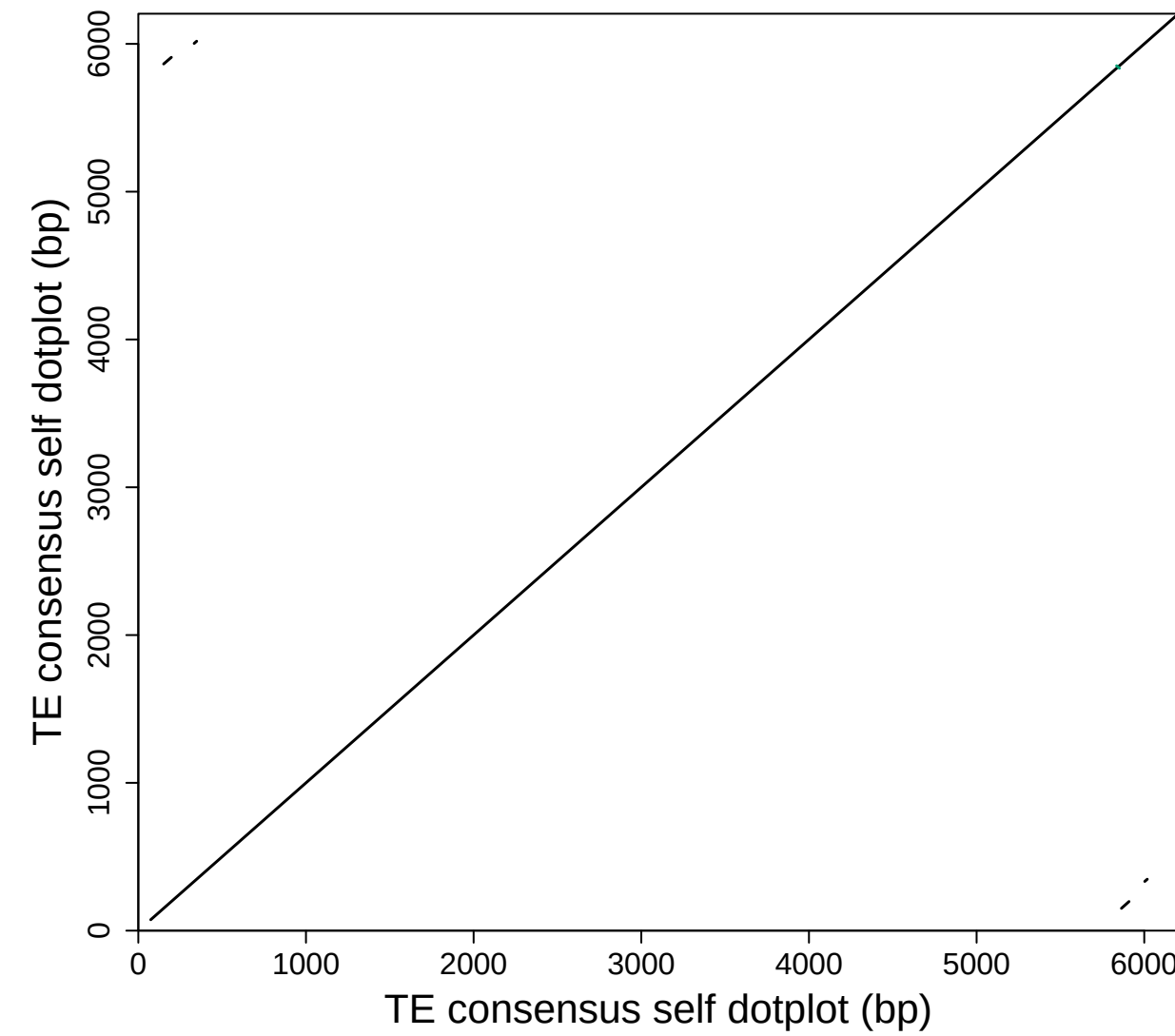
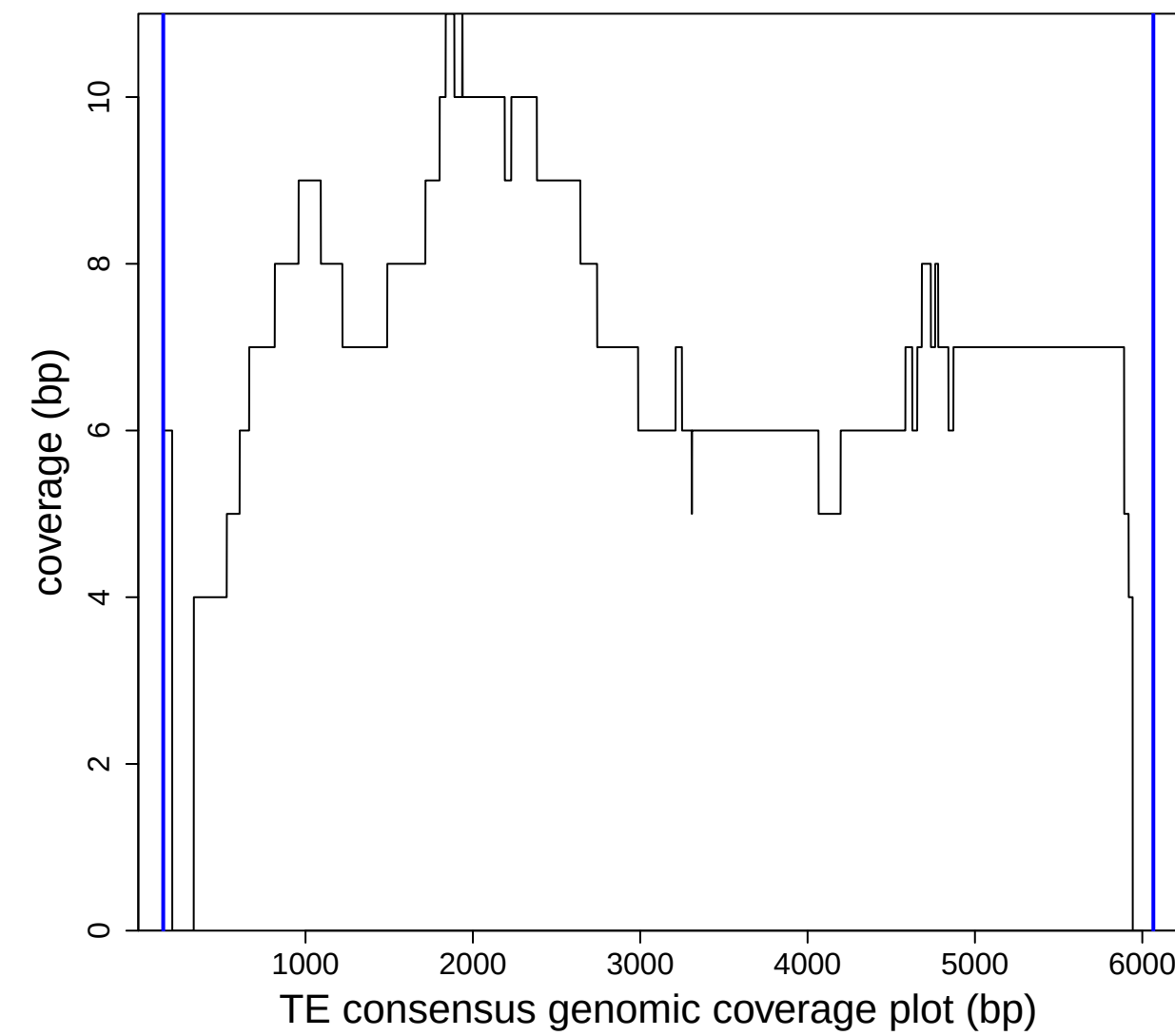
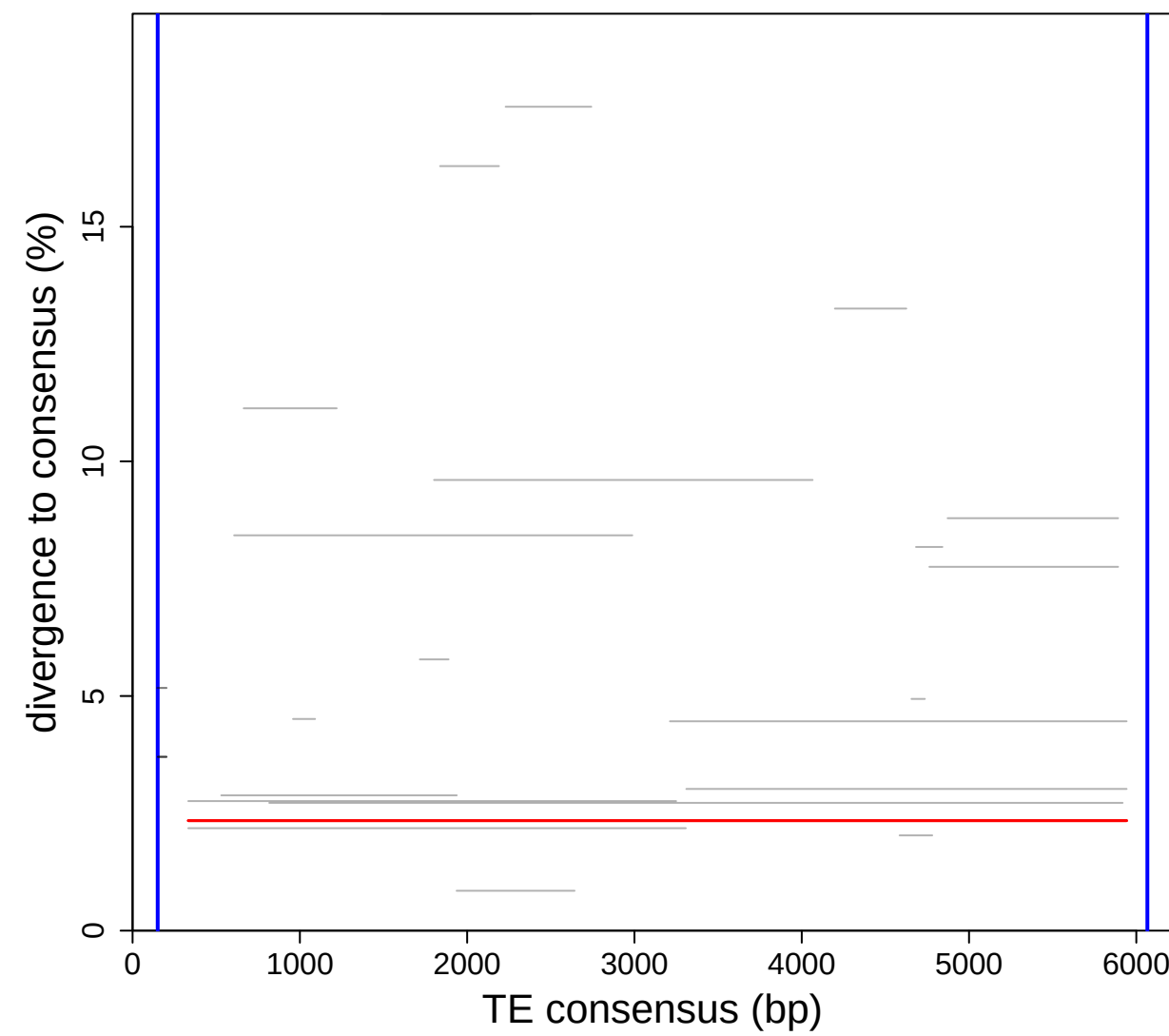
l.b.bed uf.bed g_1.bed fm_1.bed 0_0_bcln.fa aln.fa cl.fa gs.fa
size: 5663bp; fragments: 23; full length: 2 (>=5096.7bp)

After TEtrimmer 5663 bp

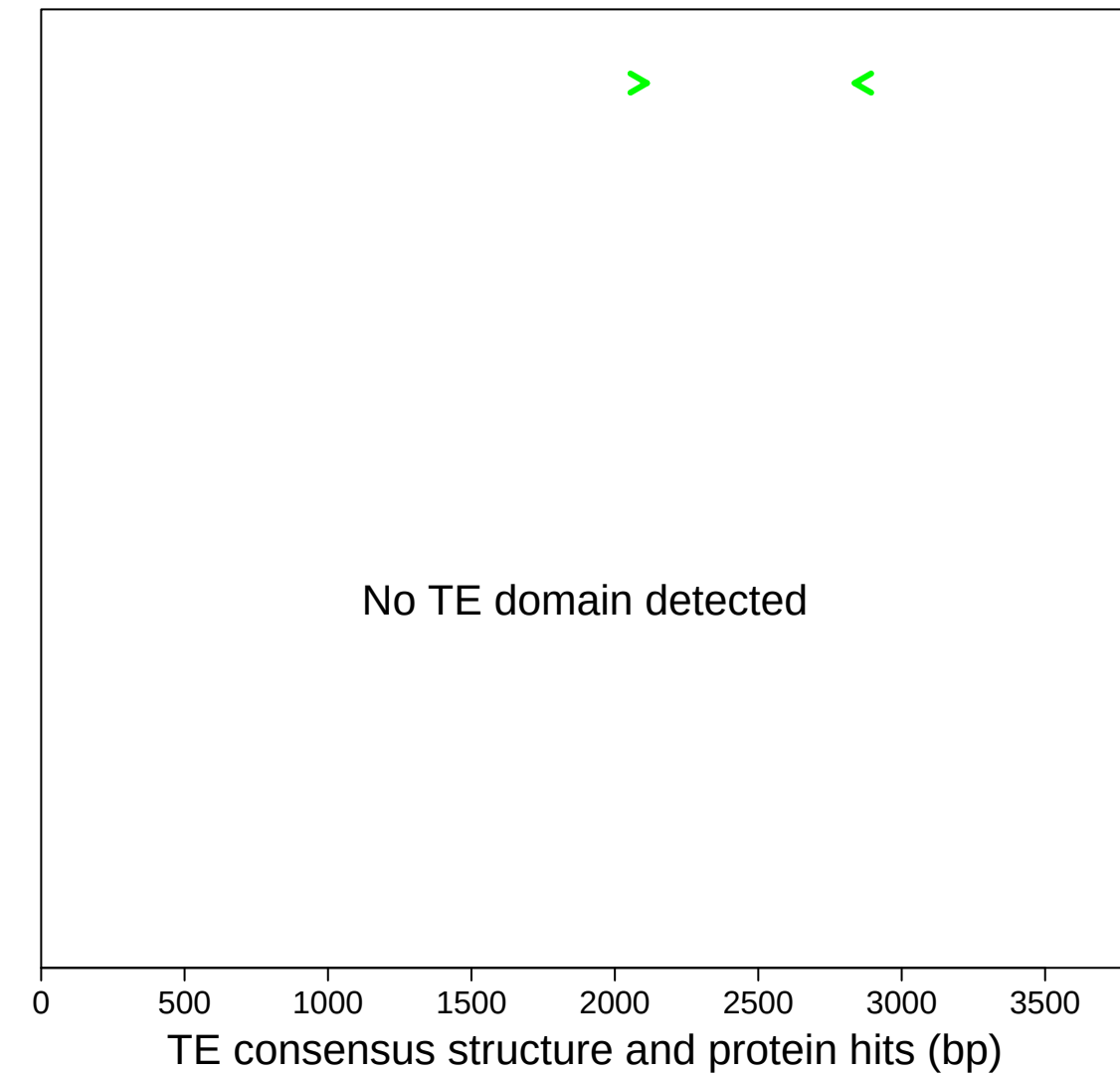
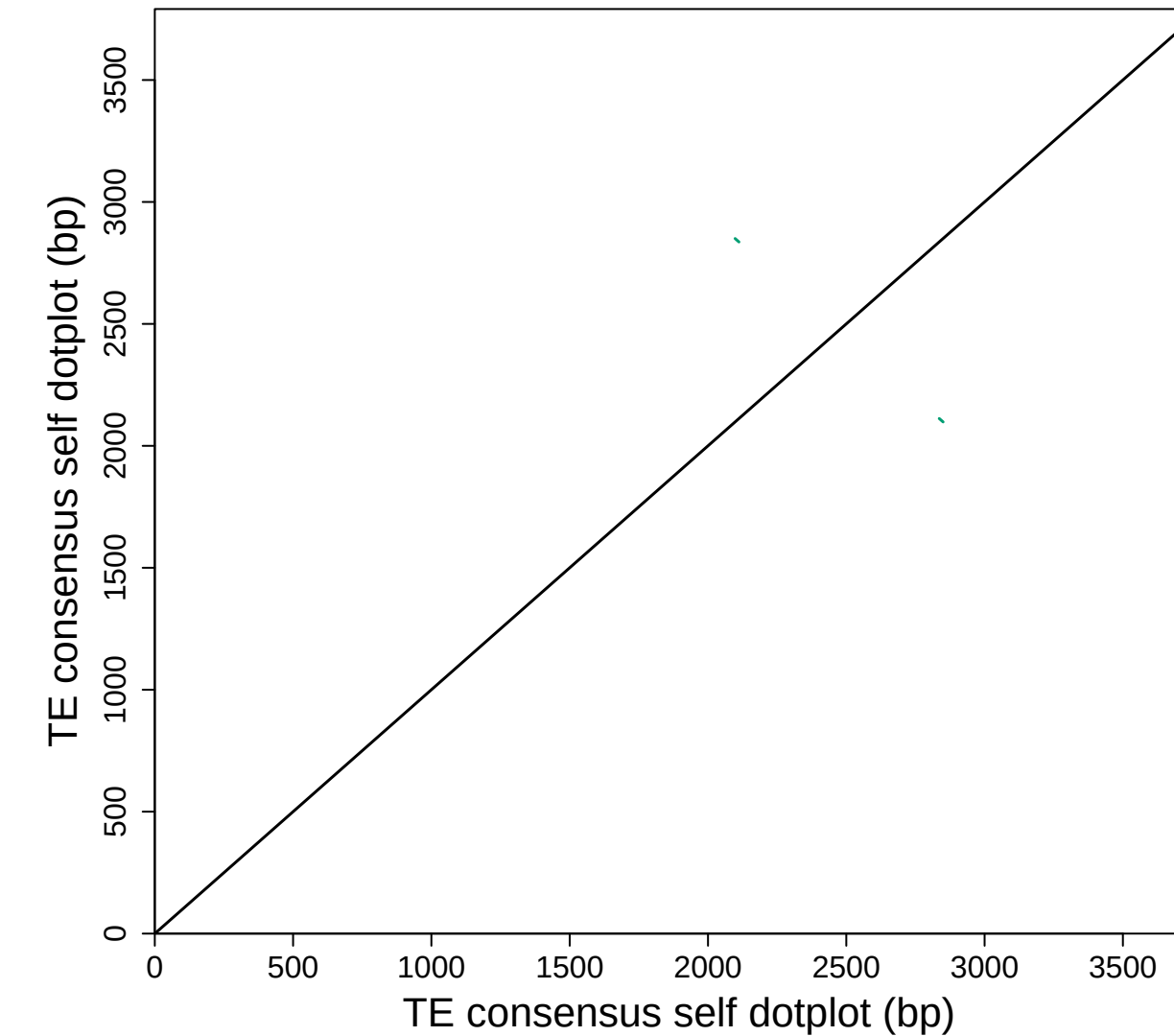
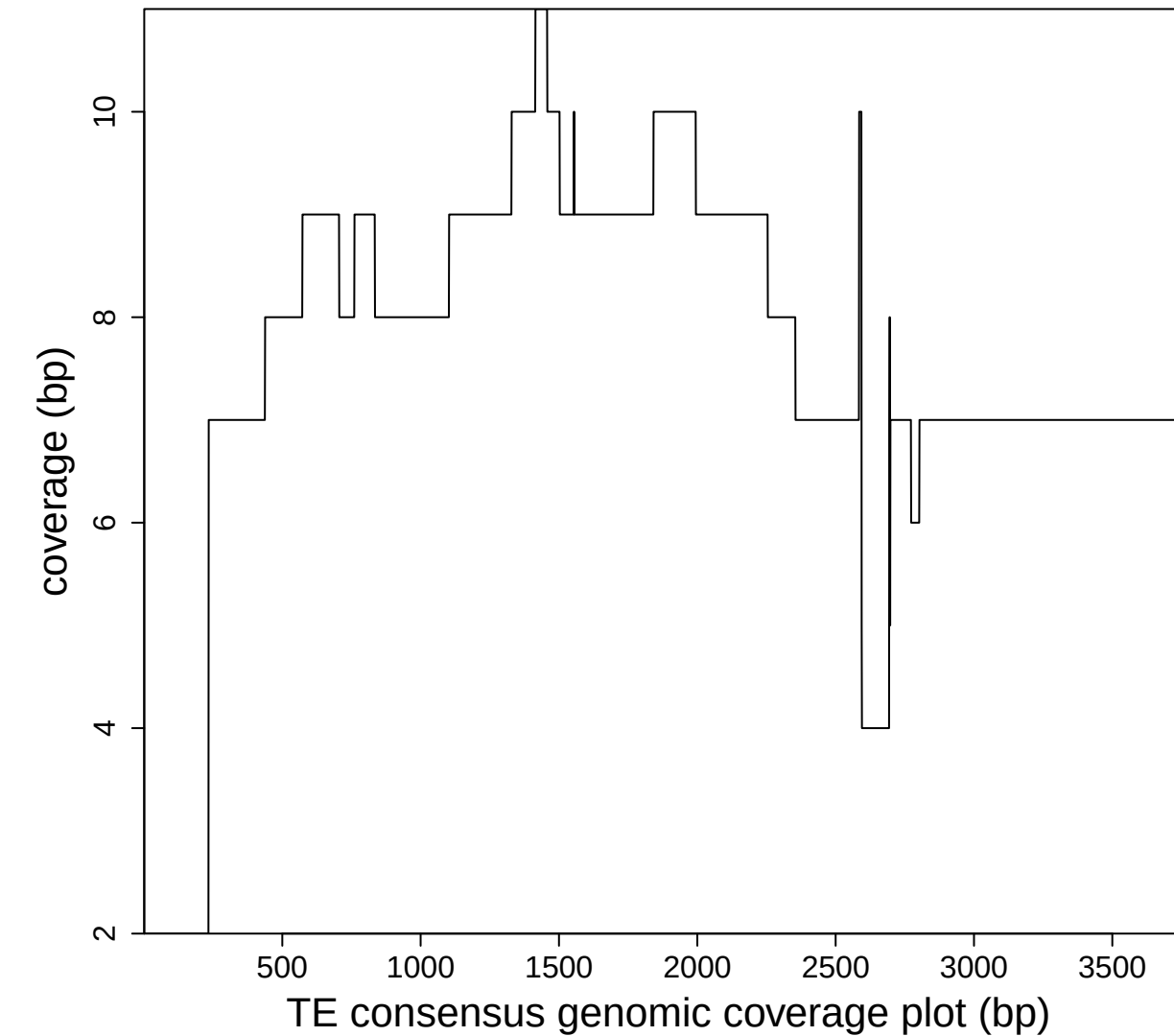
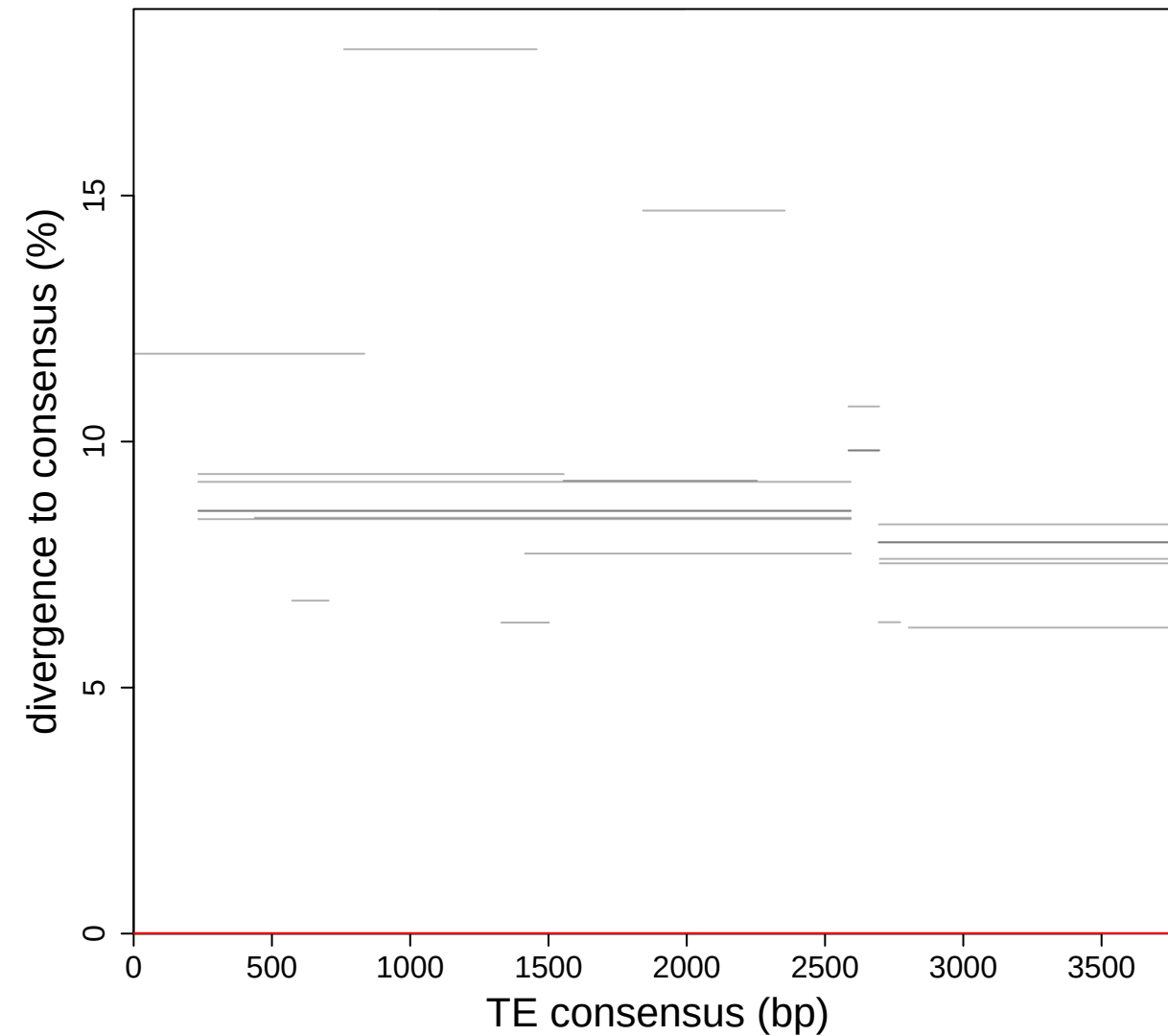


5.fasta.b.bed uf.bed g 1.bed fm 1.bed 0 0 bcln.fa aln.fa cl.fa
size: 6216bp; fragments: 29; full length: 2 (>=5594.4bp)

After TEtrimmer Extended plot Blue lines are boundaries

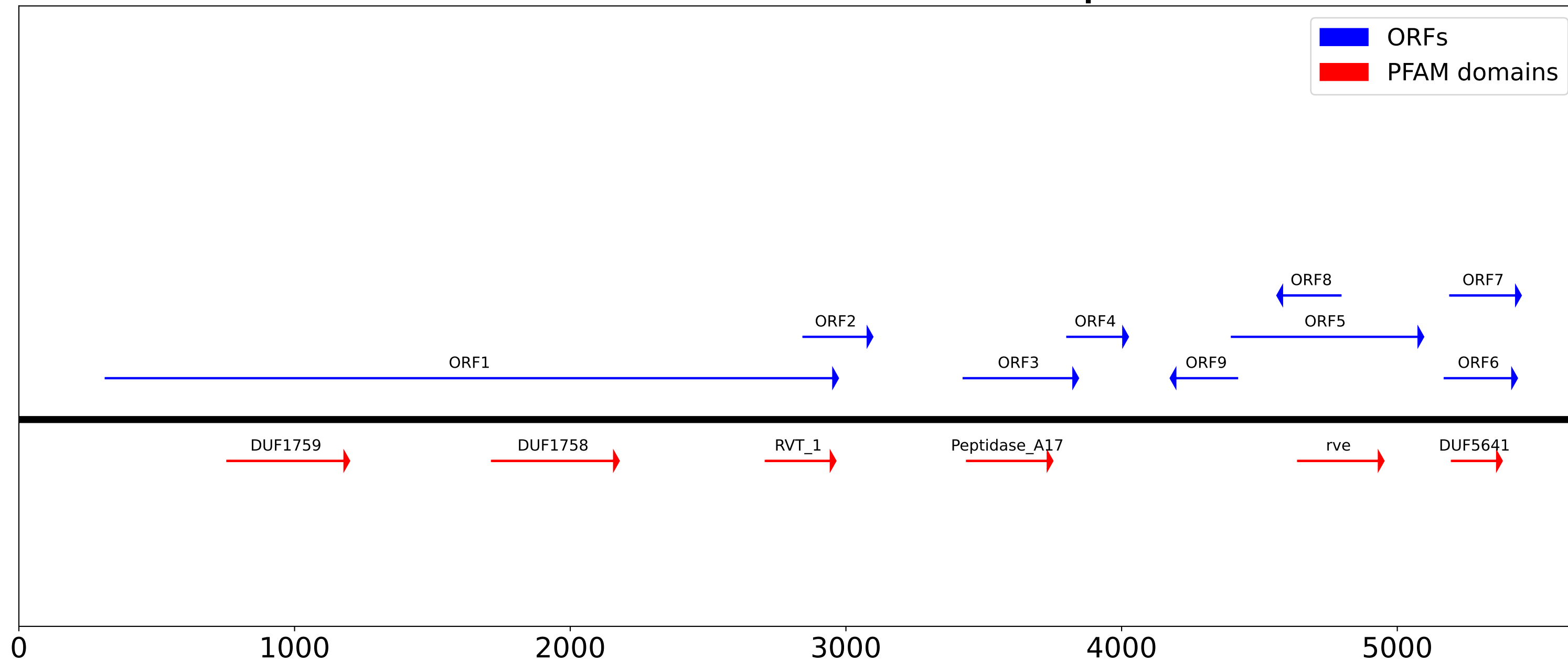


TE: ltr_1_family_1045
size: 3791bp; fragments: 25; full length: 1 (≥ 3411.9 bp)

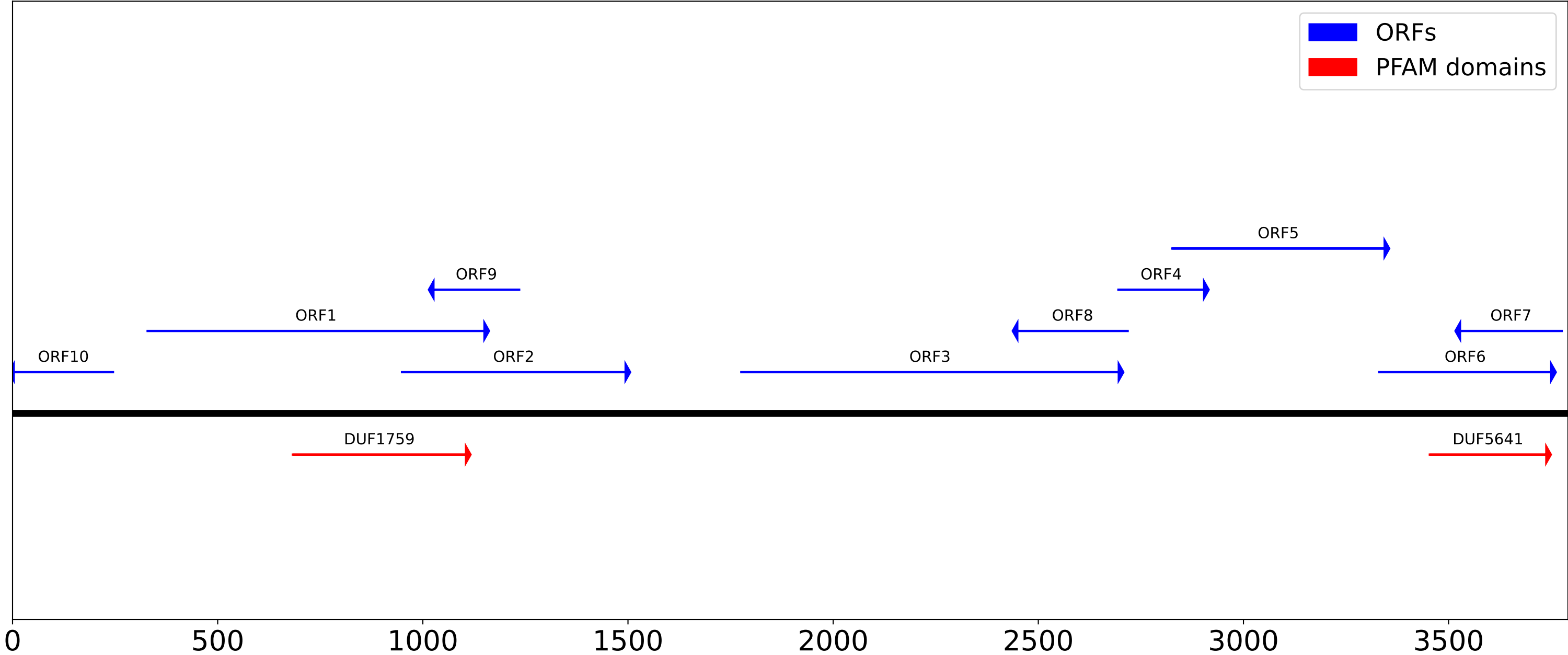


Before TEtrimmer 3791 bp

After TEtrimmer ORF and PFAM domain plot



Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

