

Start crop Point

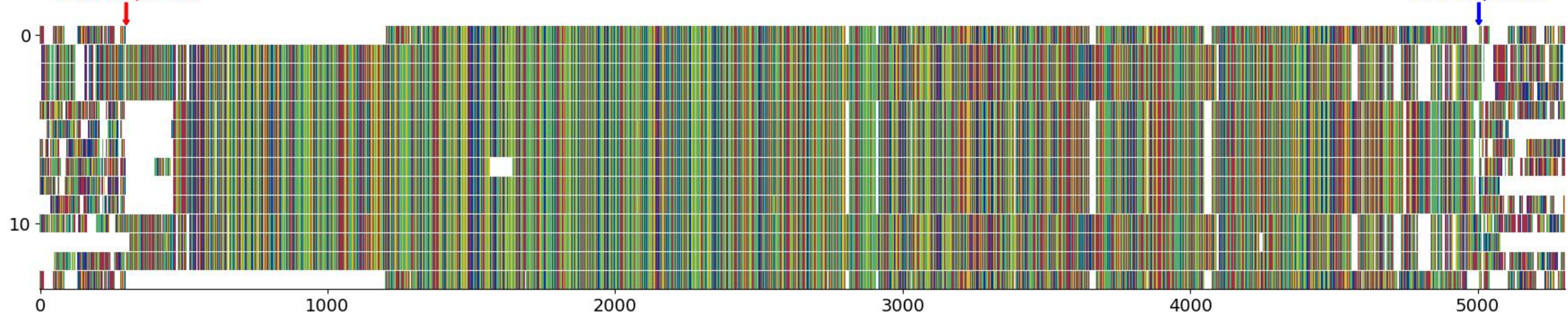
End crop Point

MSA length = 4705



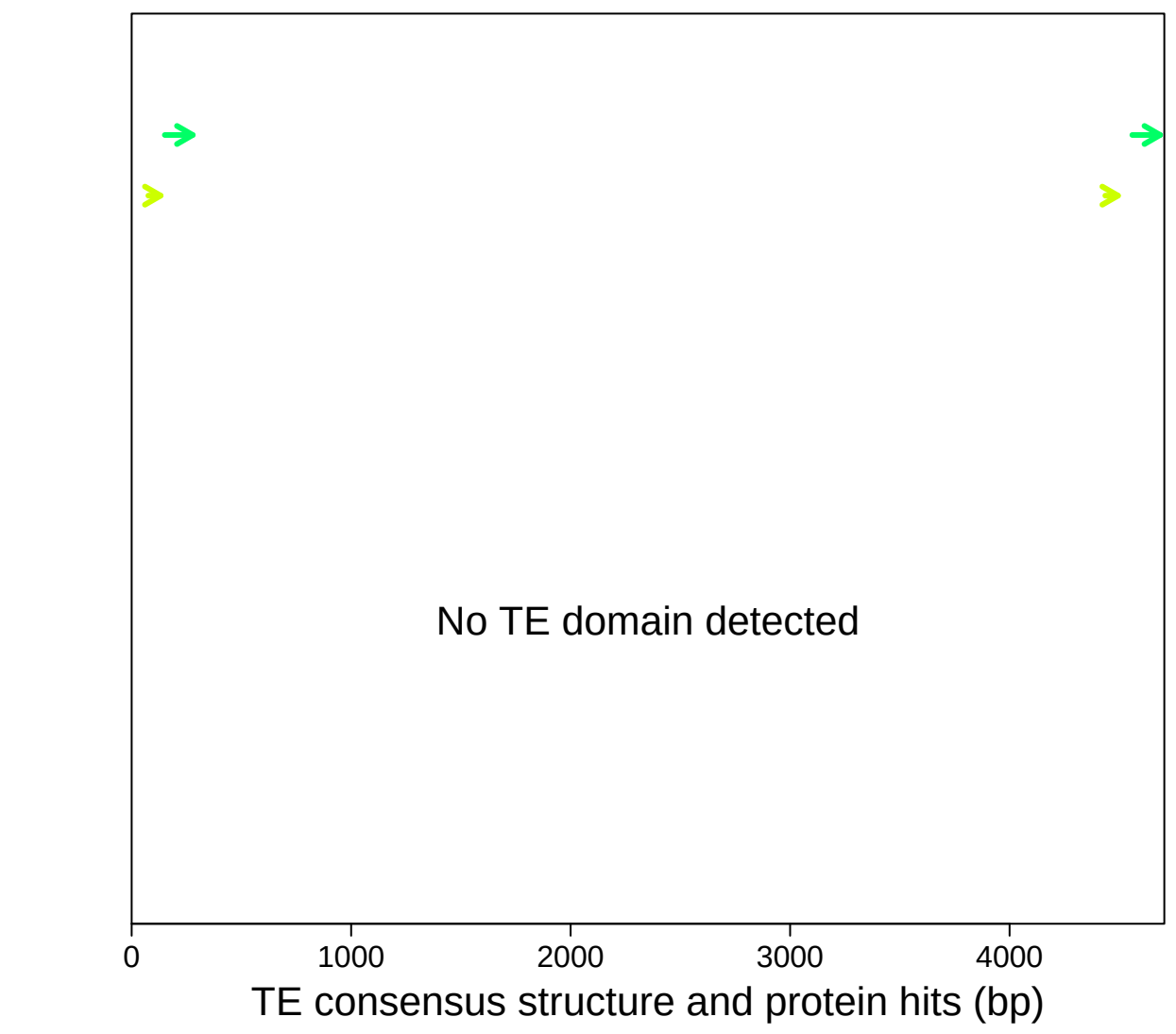
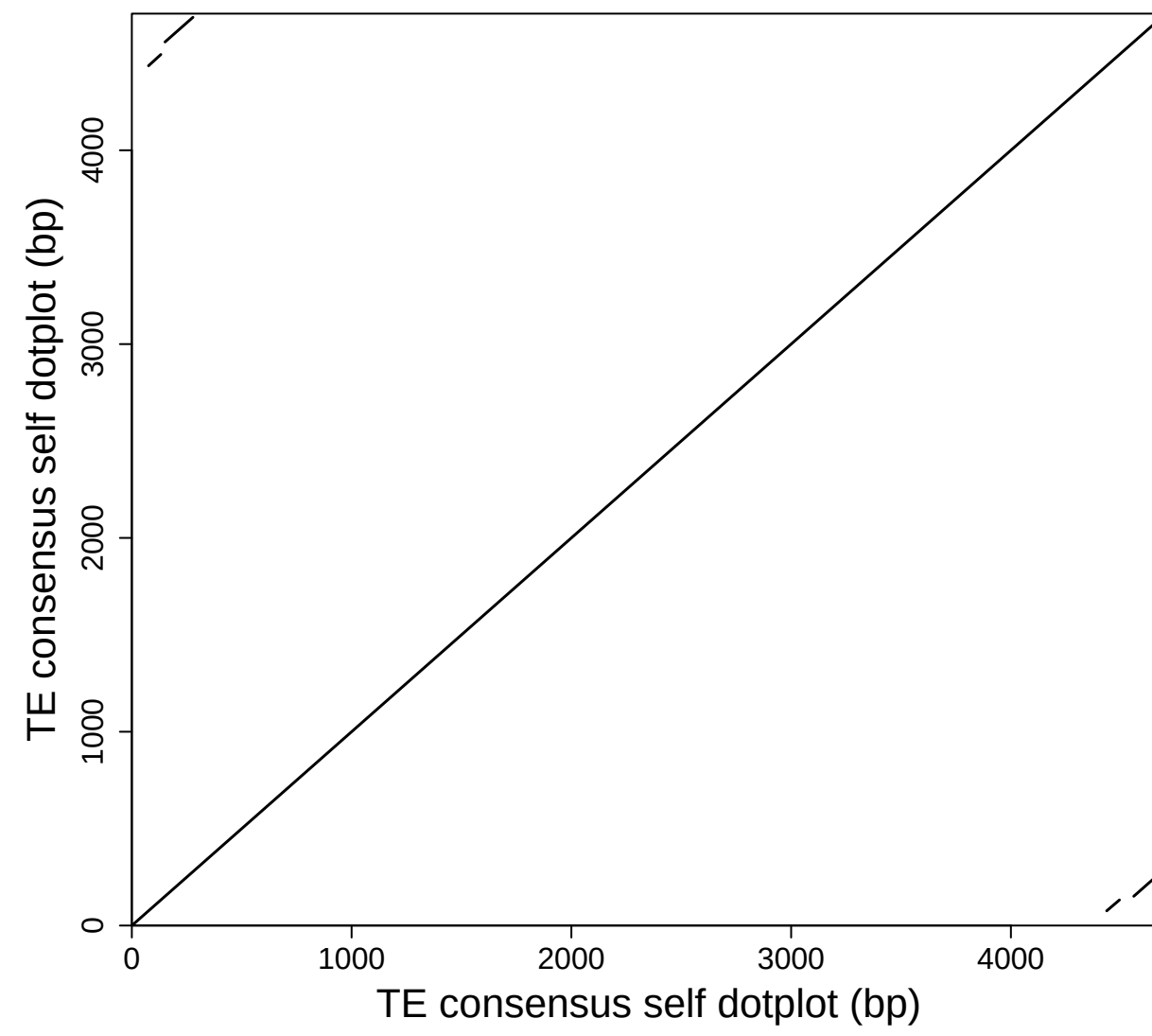
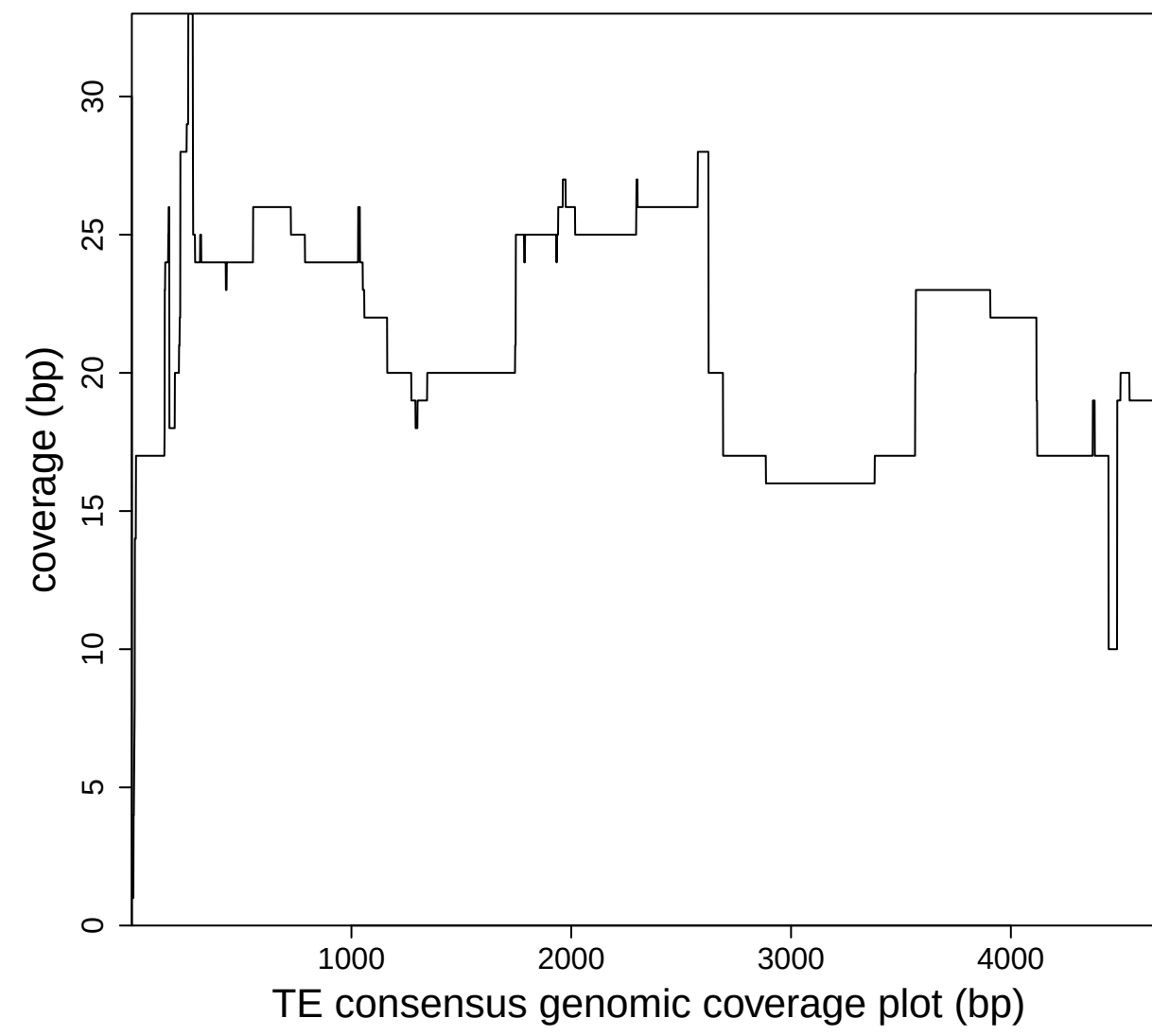
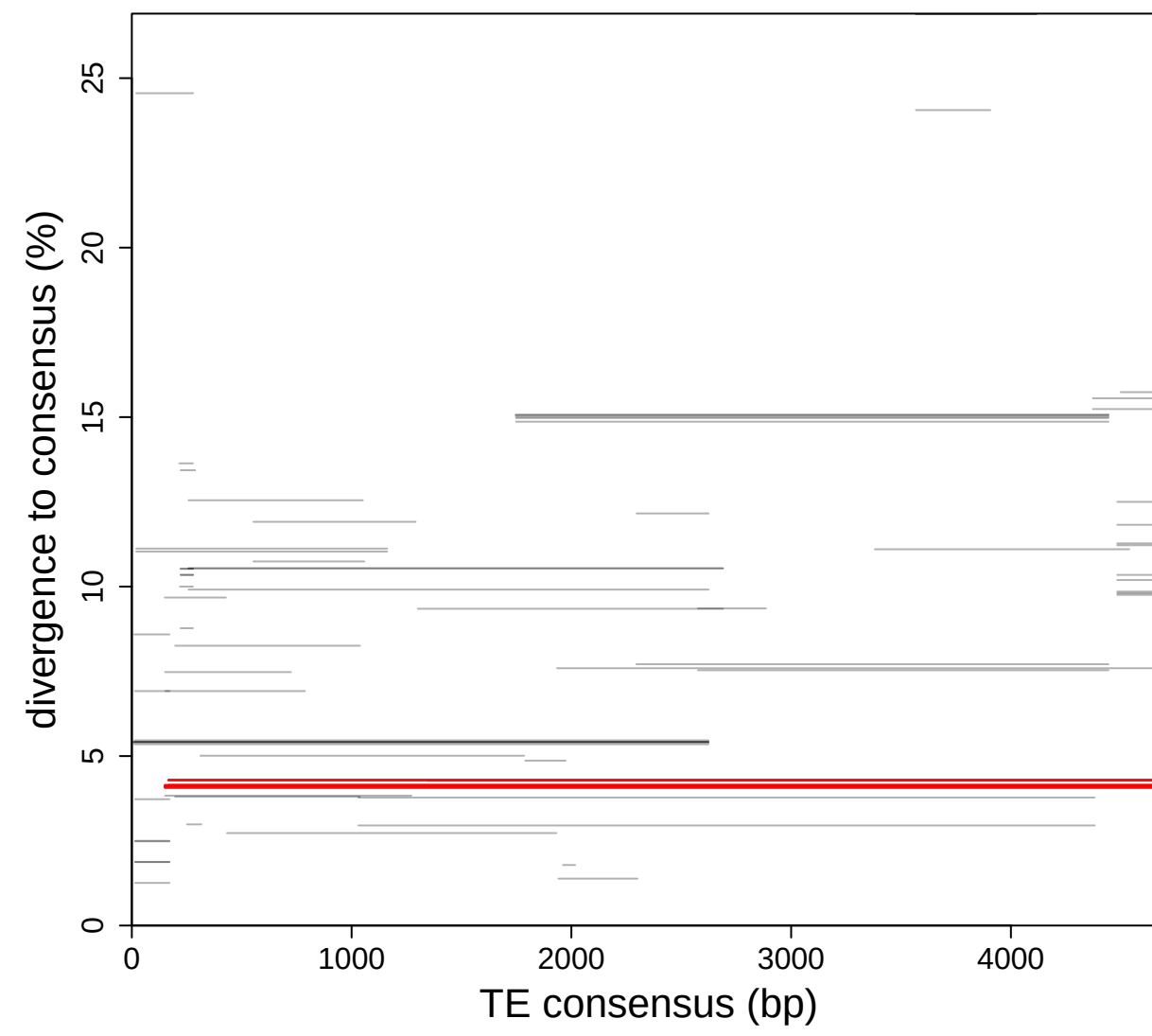
Start crop Point

End crop Point



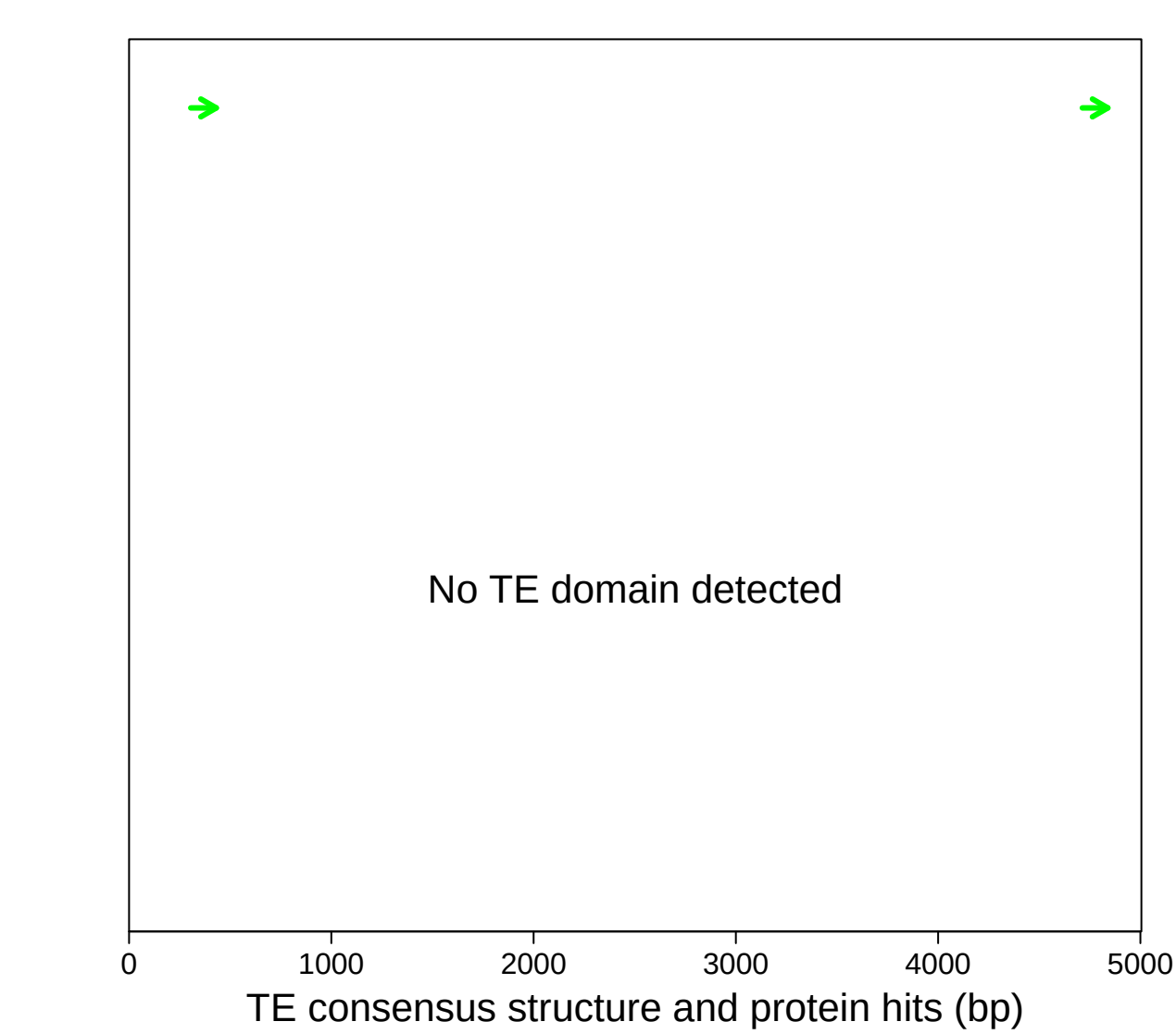
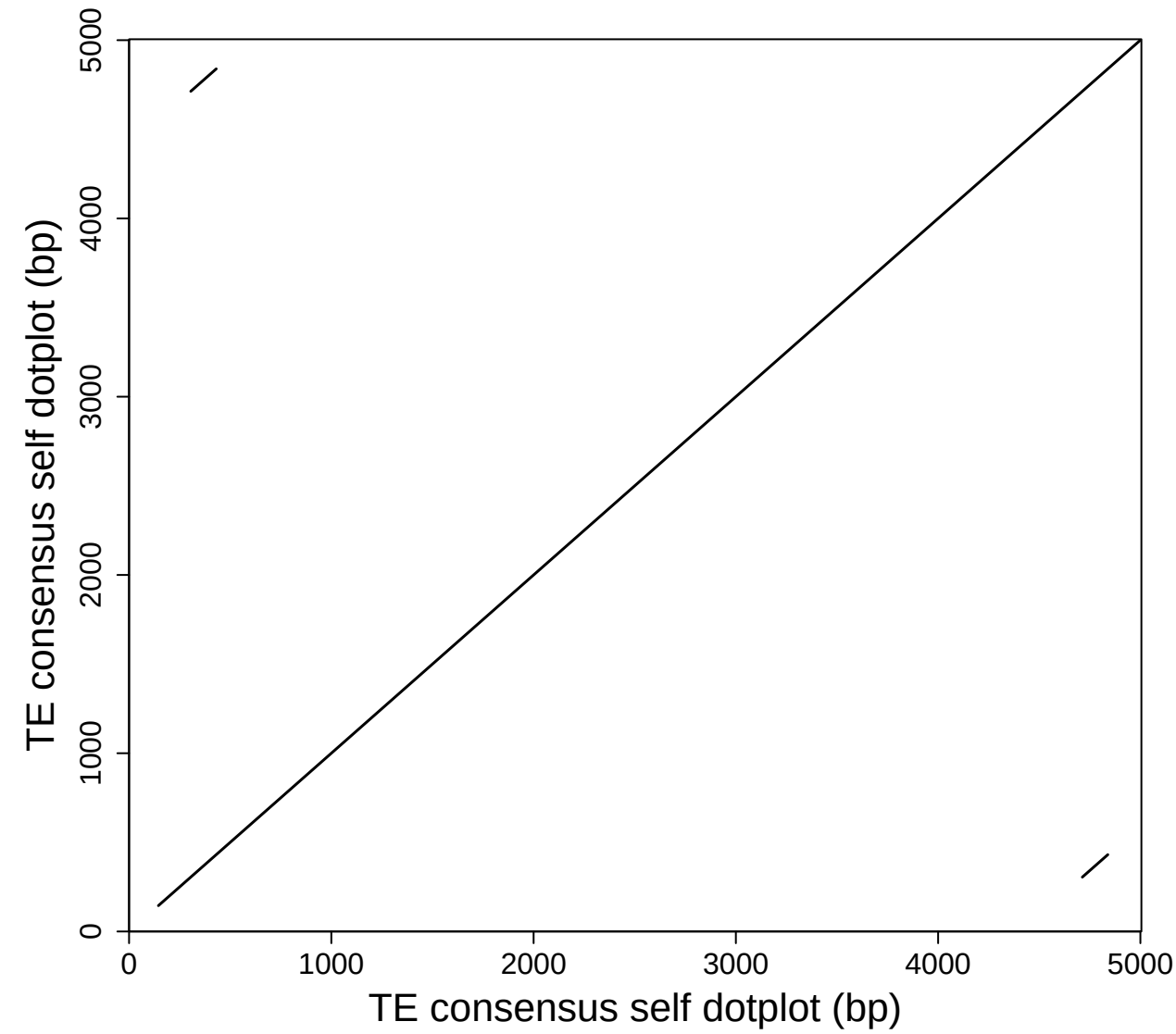
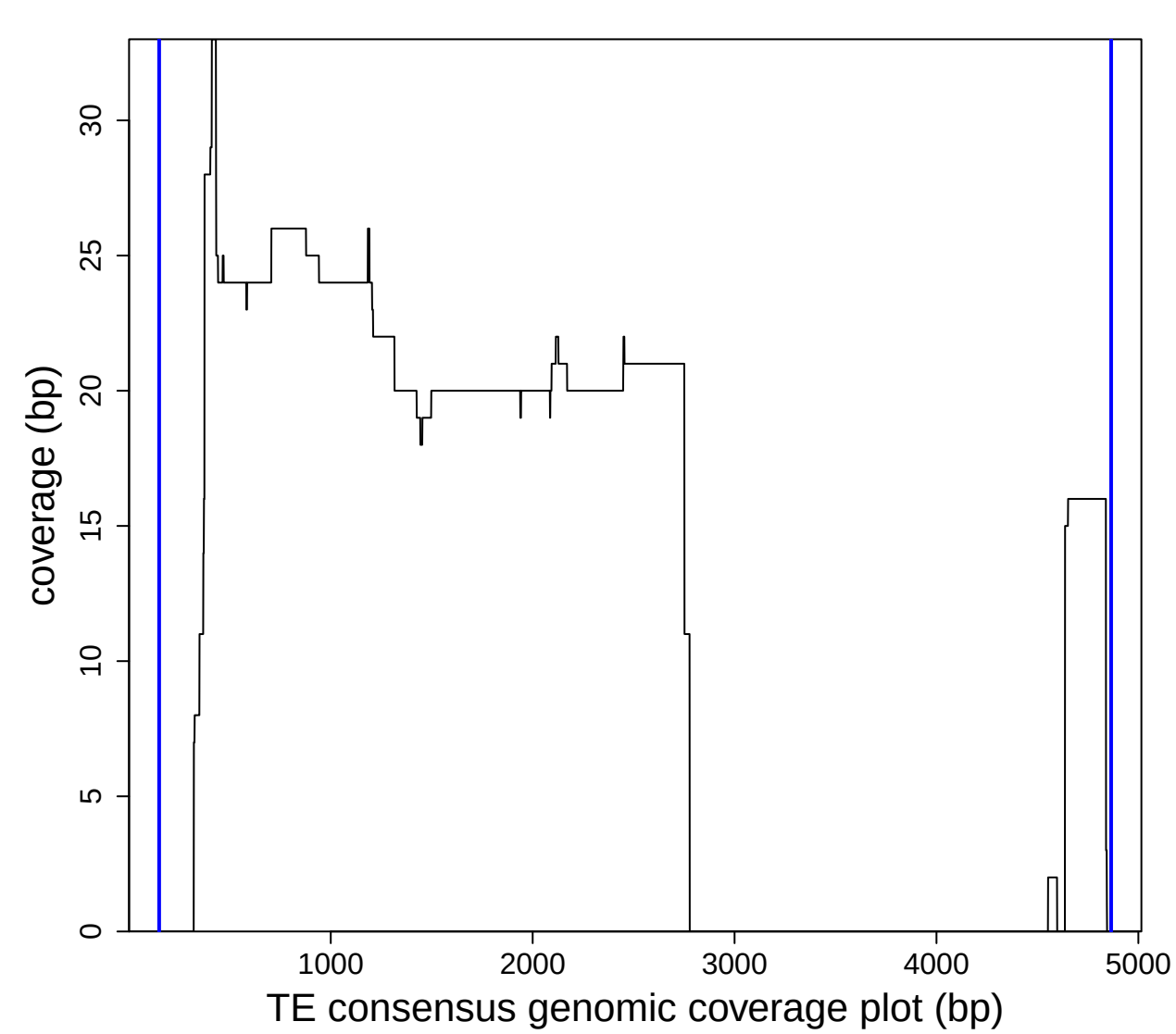
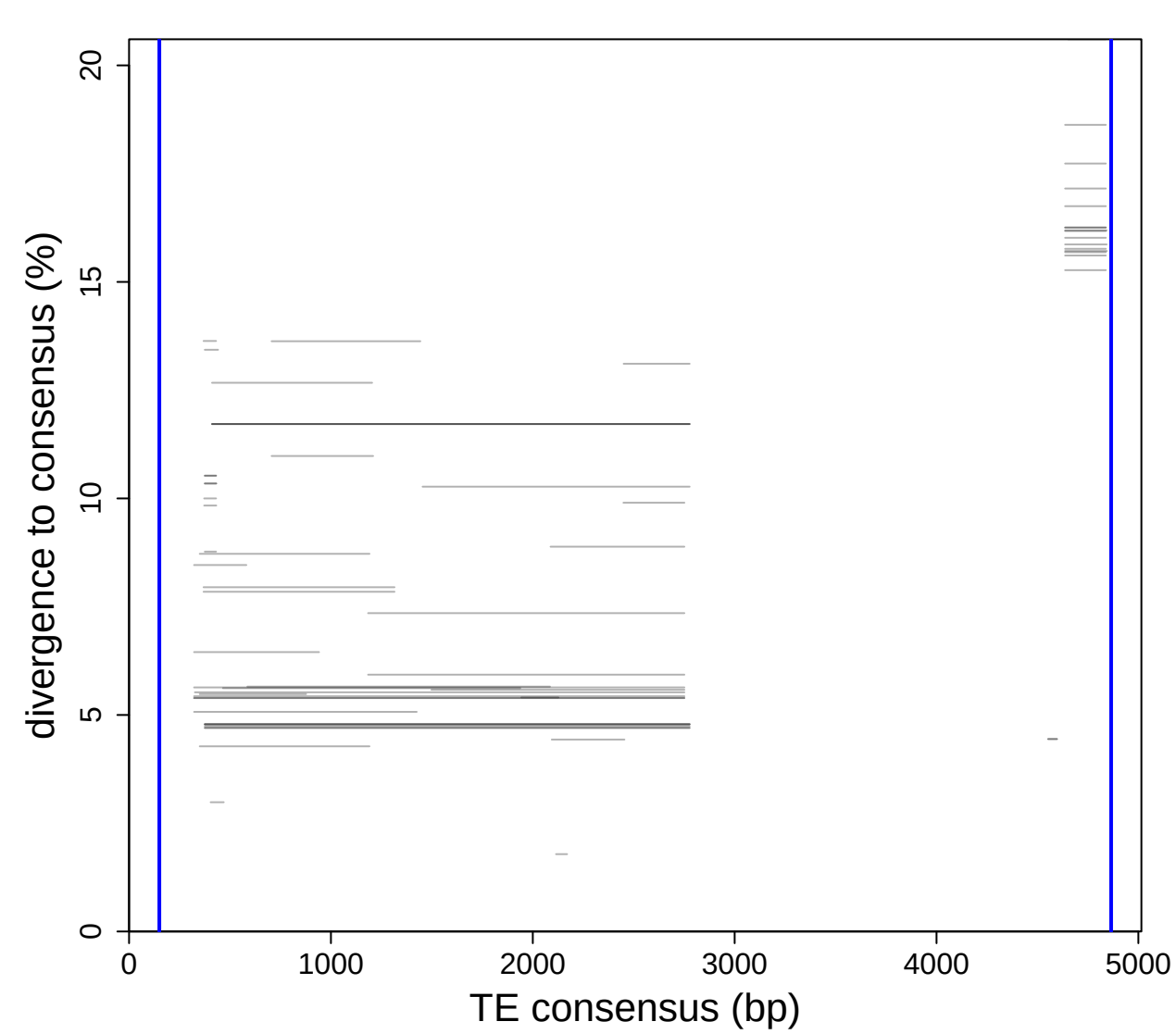
l.b.bed uf.bed g_1.bed fm_1.bed 0_0_bcln.fa aln.fa cl.fa gs.fa
size: 4705bp; fragments: 81; full length: 5 (>=4234.5bp)

After TEtrimmer 4705 bp

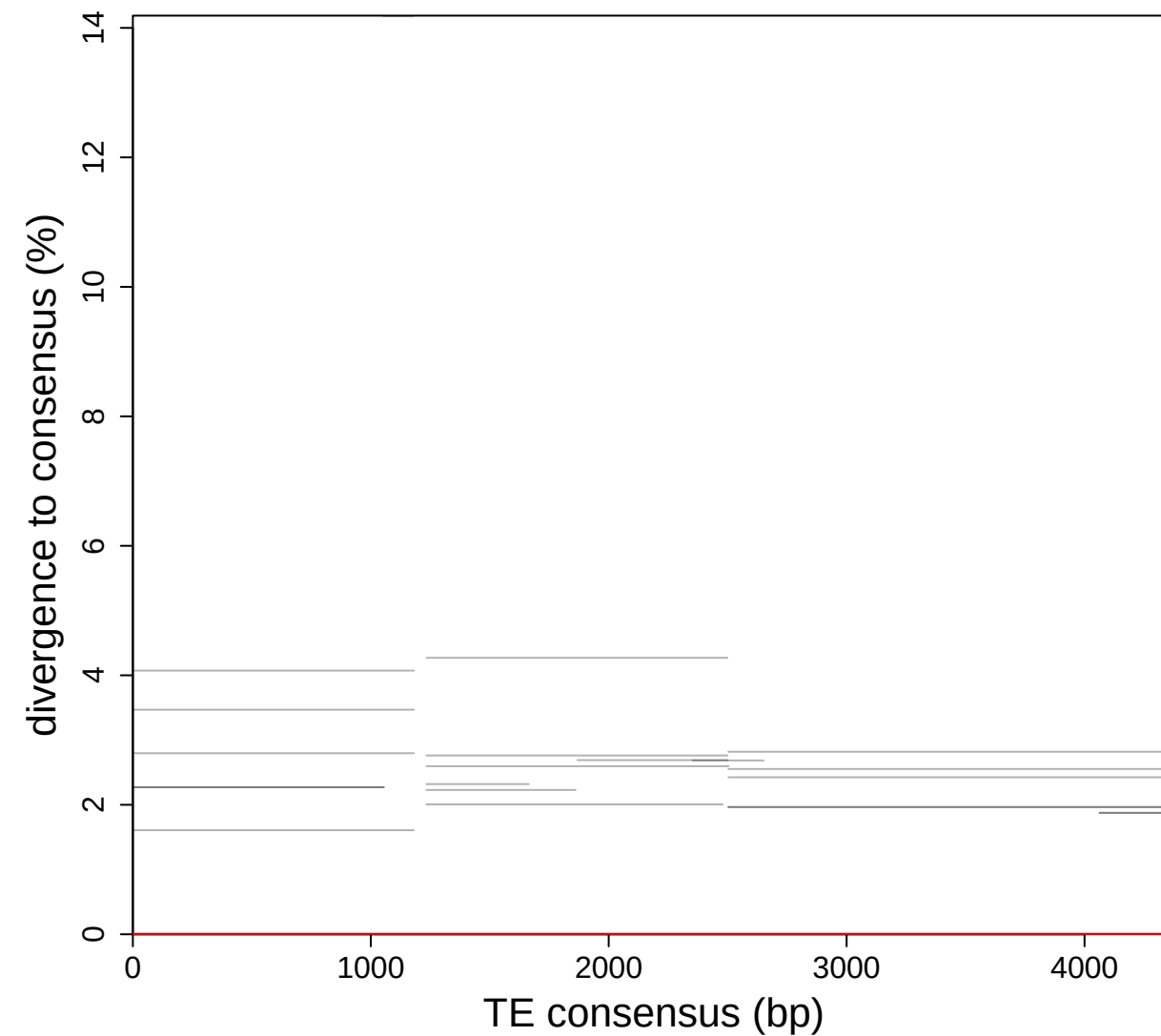


2.fasta.b.bed uf.bed g 1.bed fm 1.bed 0 0 bcln.fa aln.fa cl.fa
size: 5015bp; fragments: 65; full length: 0 (>=4513.5bp)

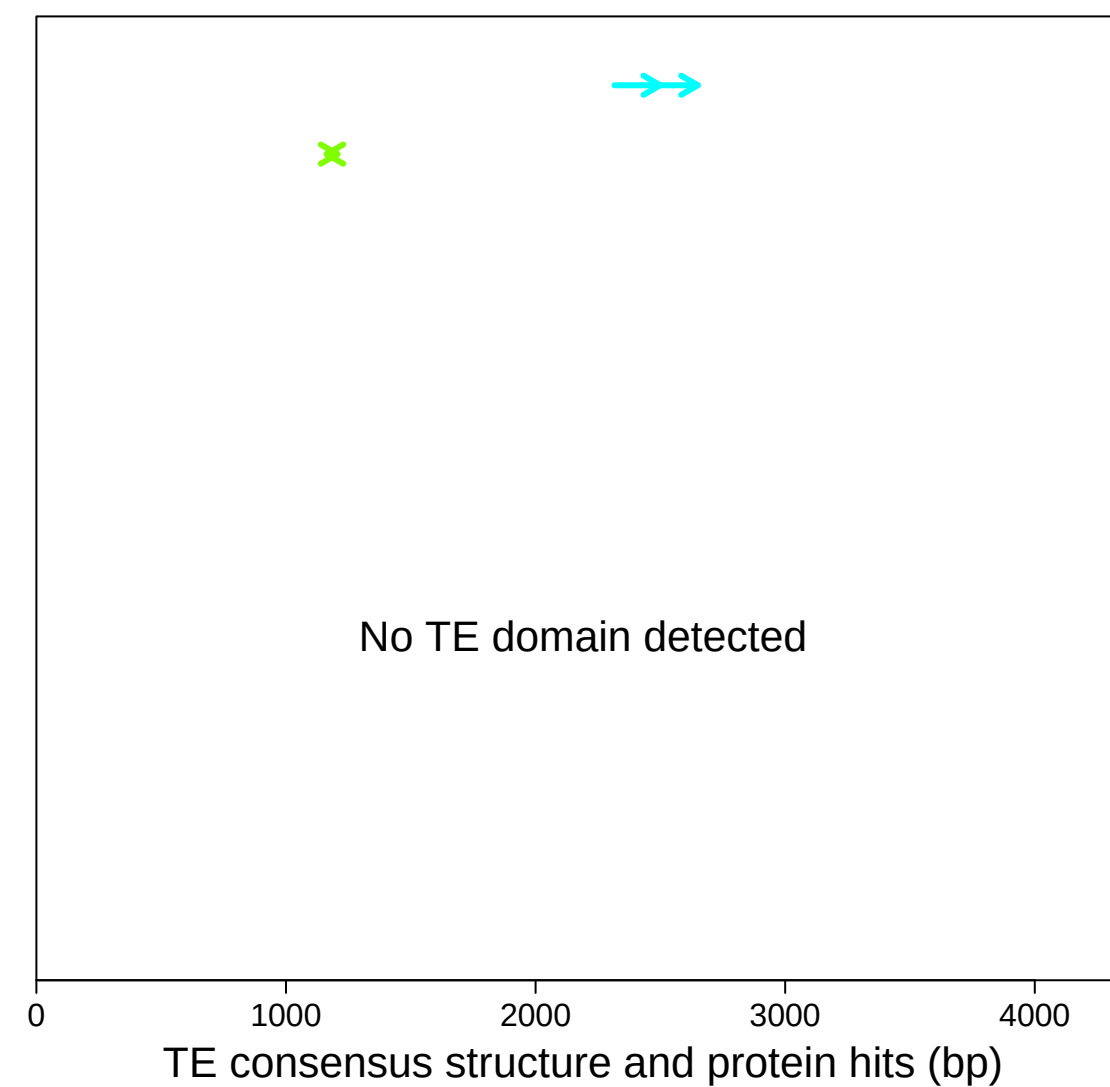
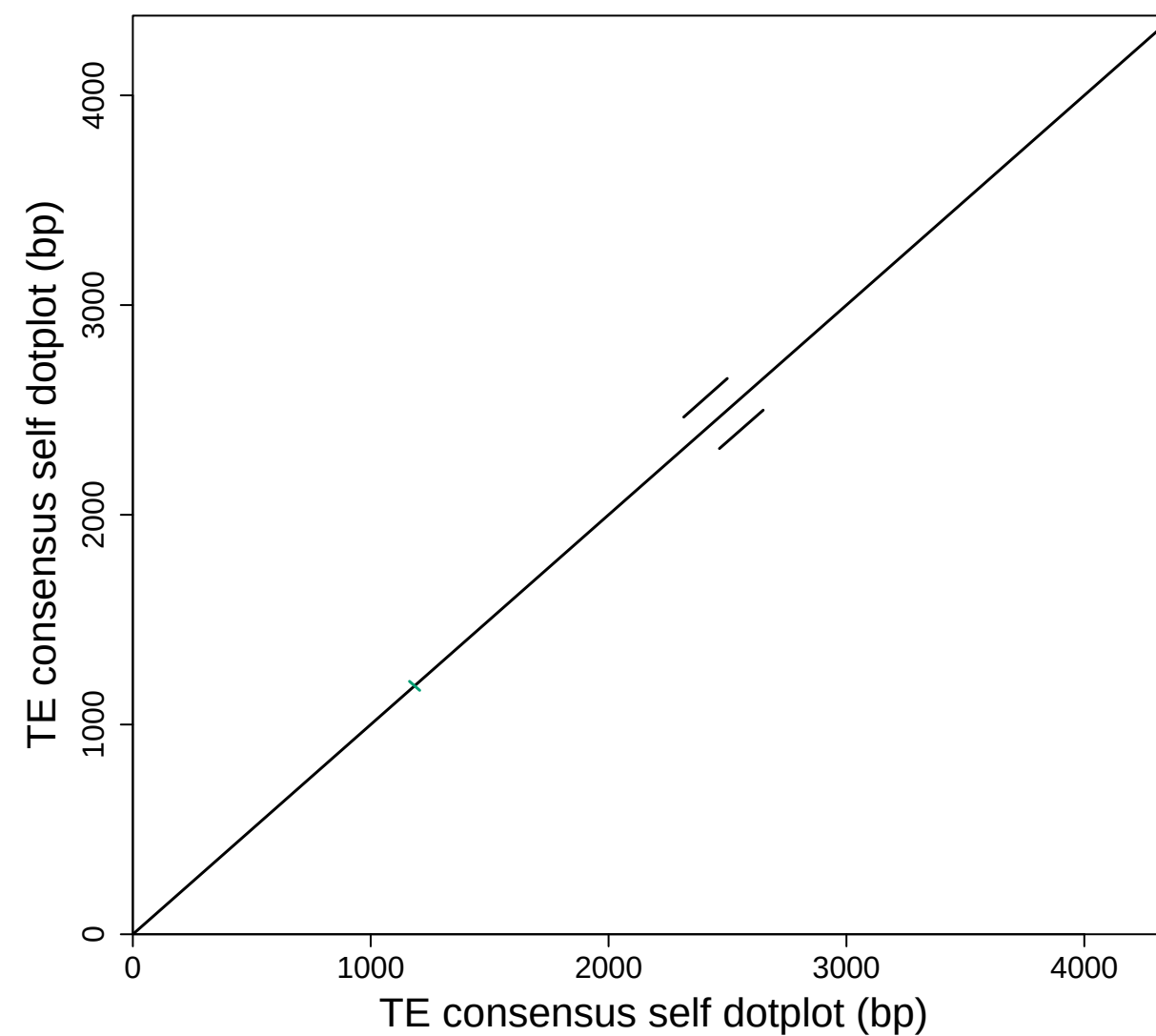
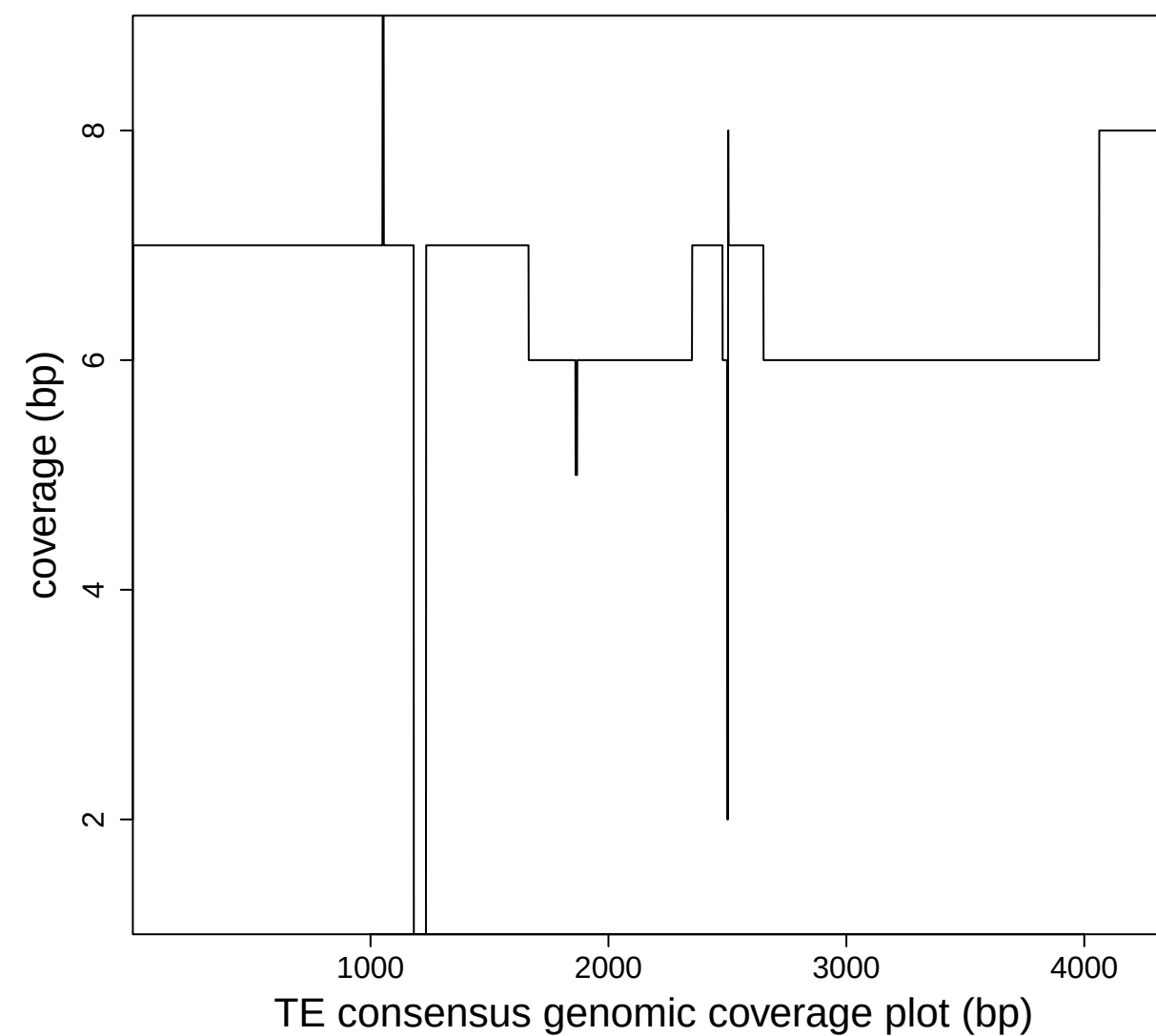
After TEtrimmer Extended plot Blue lines are boundaries



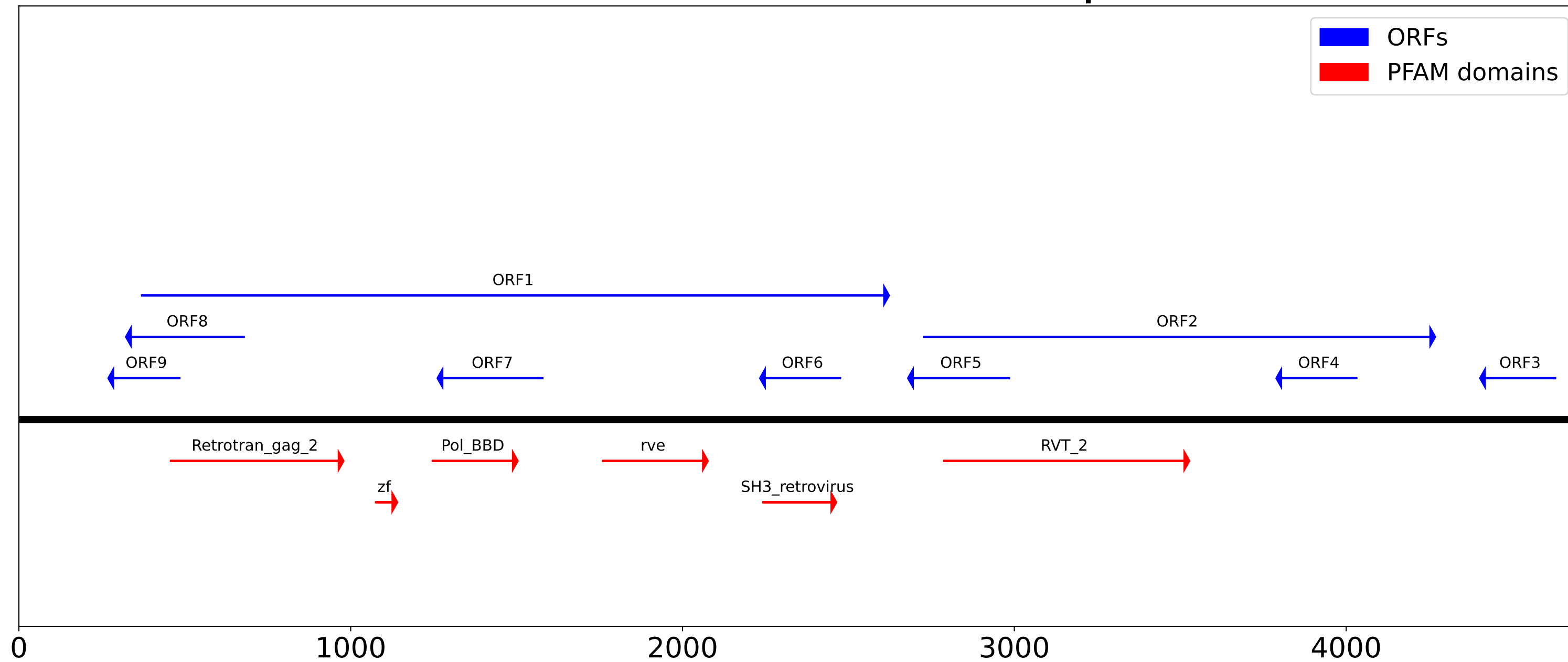
TE: ltr_1_family_1062
size: 4380bp; fragments: 25; full length: 1 (≥ 3942 bp)



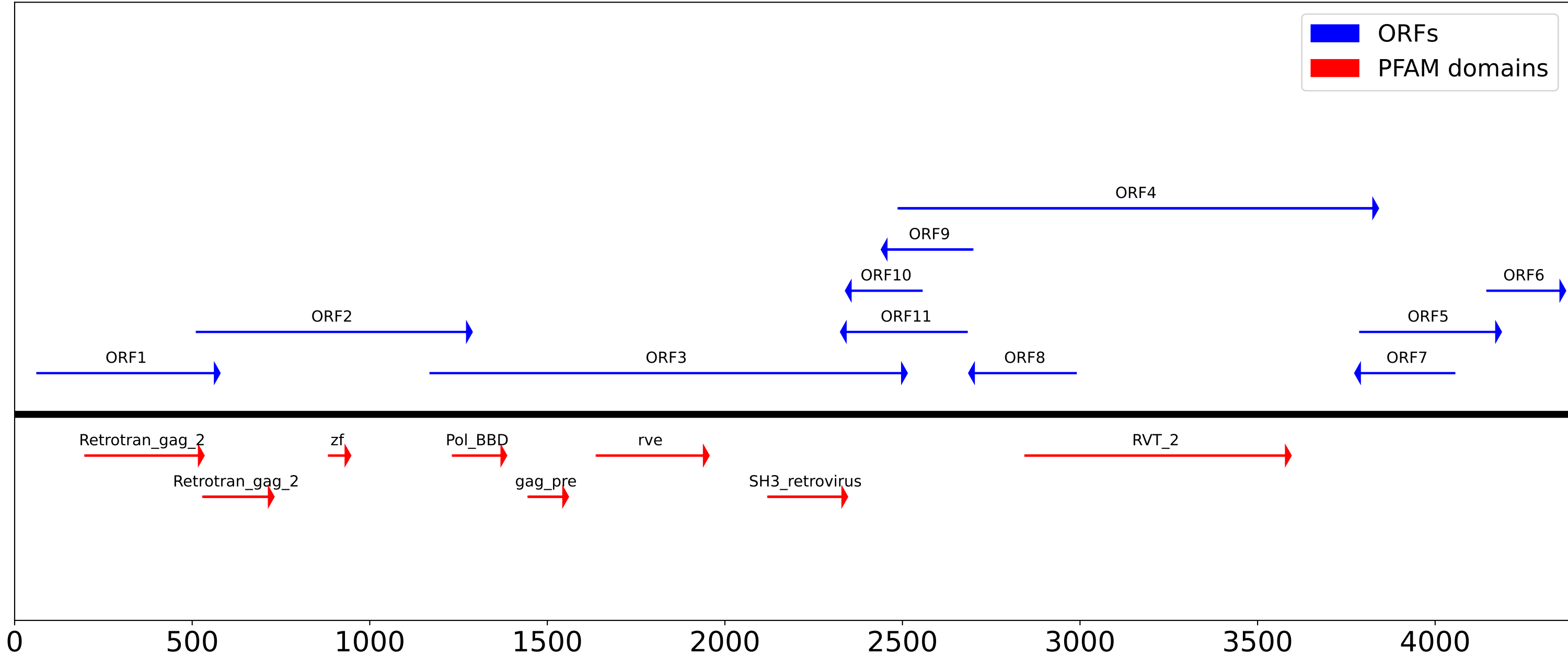
Before TEtrimmer 4380 bp



After TEtrimmer ORF and PFAM domain plot



Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

