

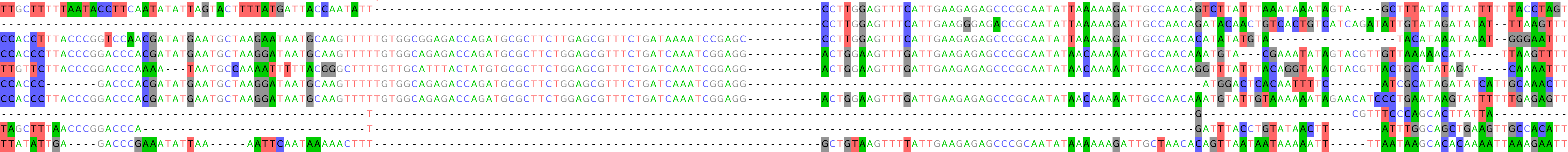
Start Crop Point



End Crop Point

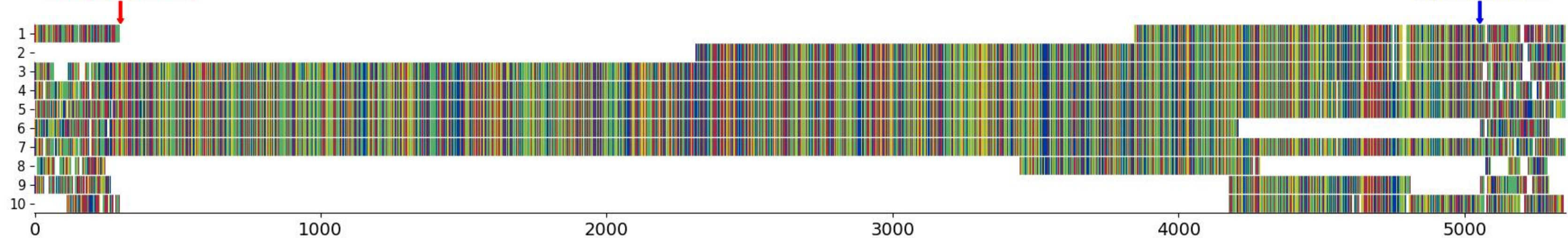


MSA length = 4755



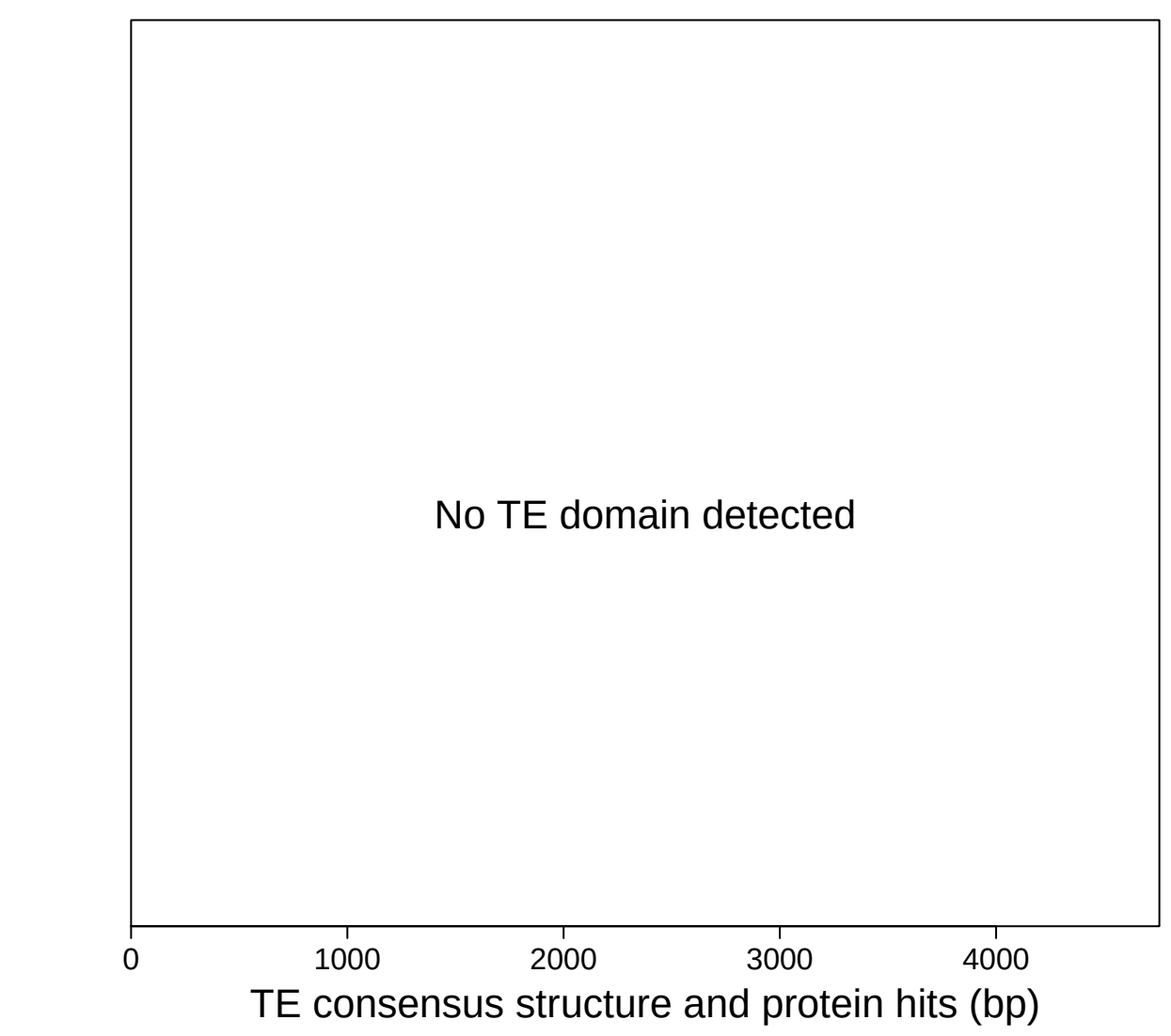
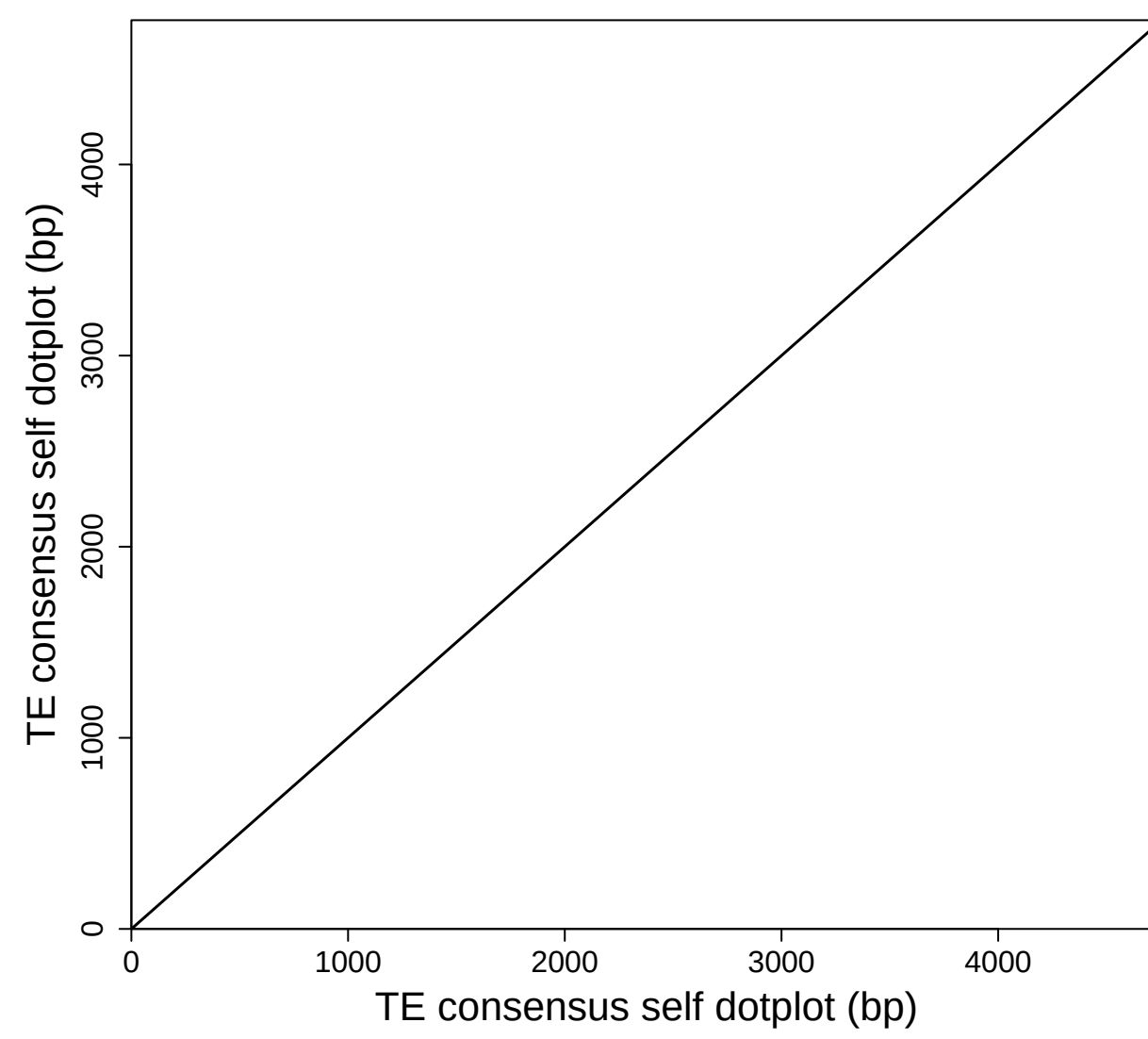
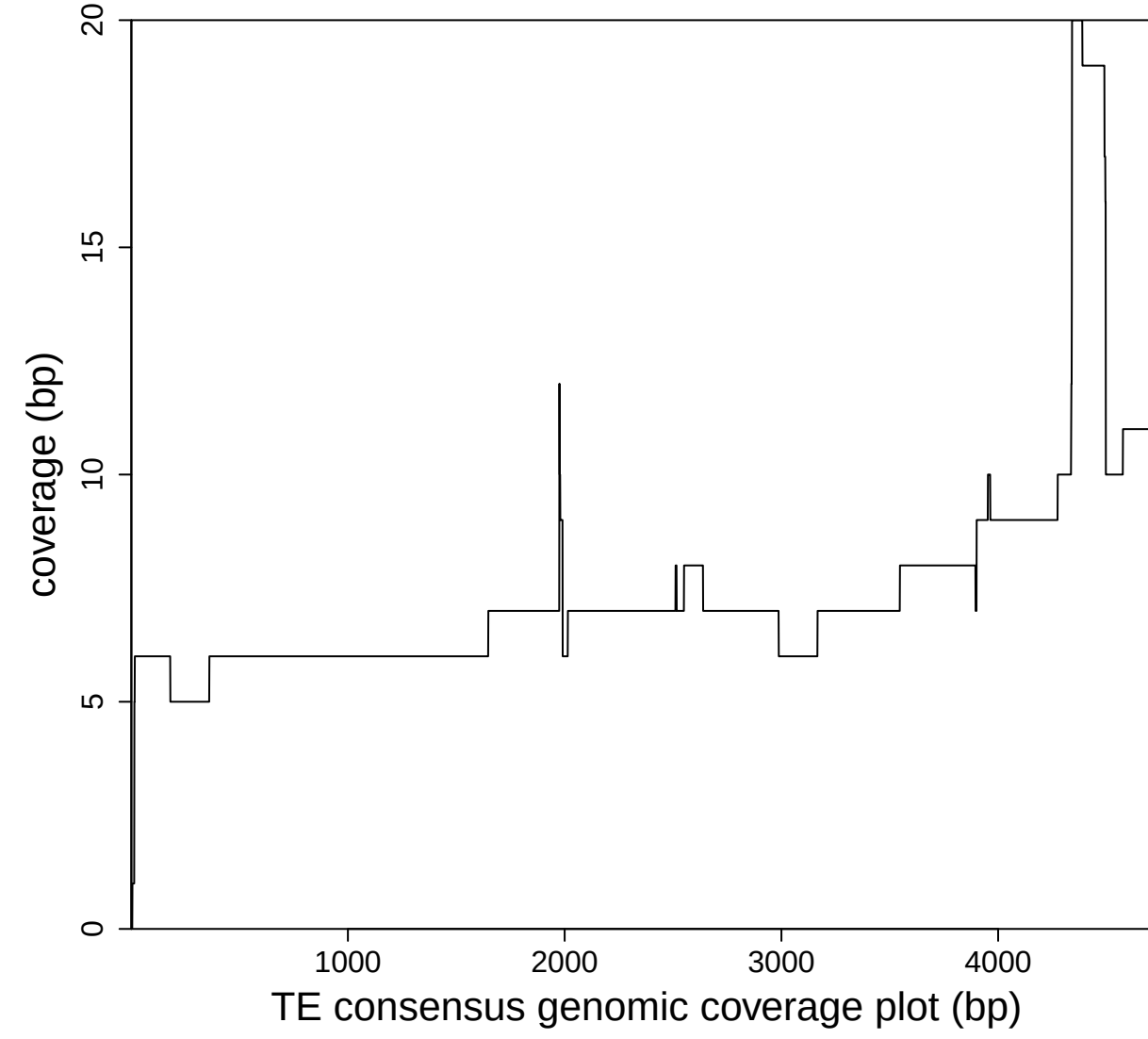
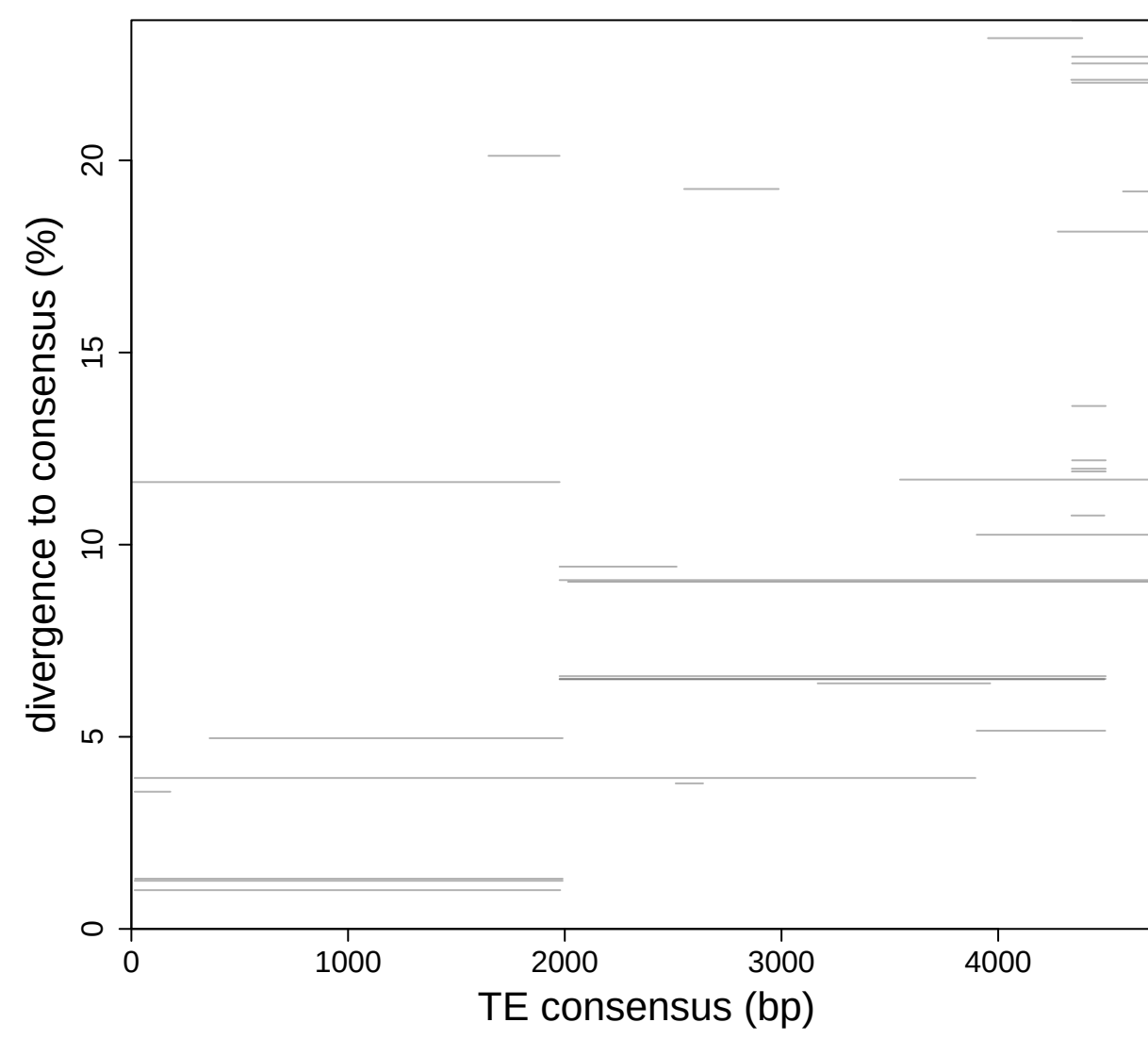
Start crop Point

End crop Point



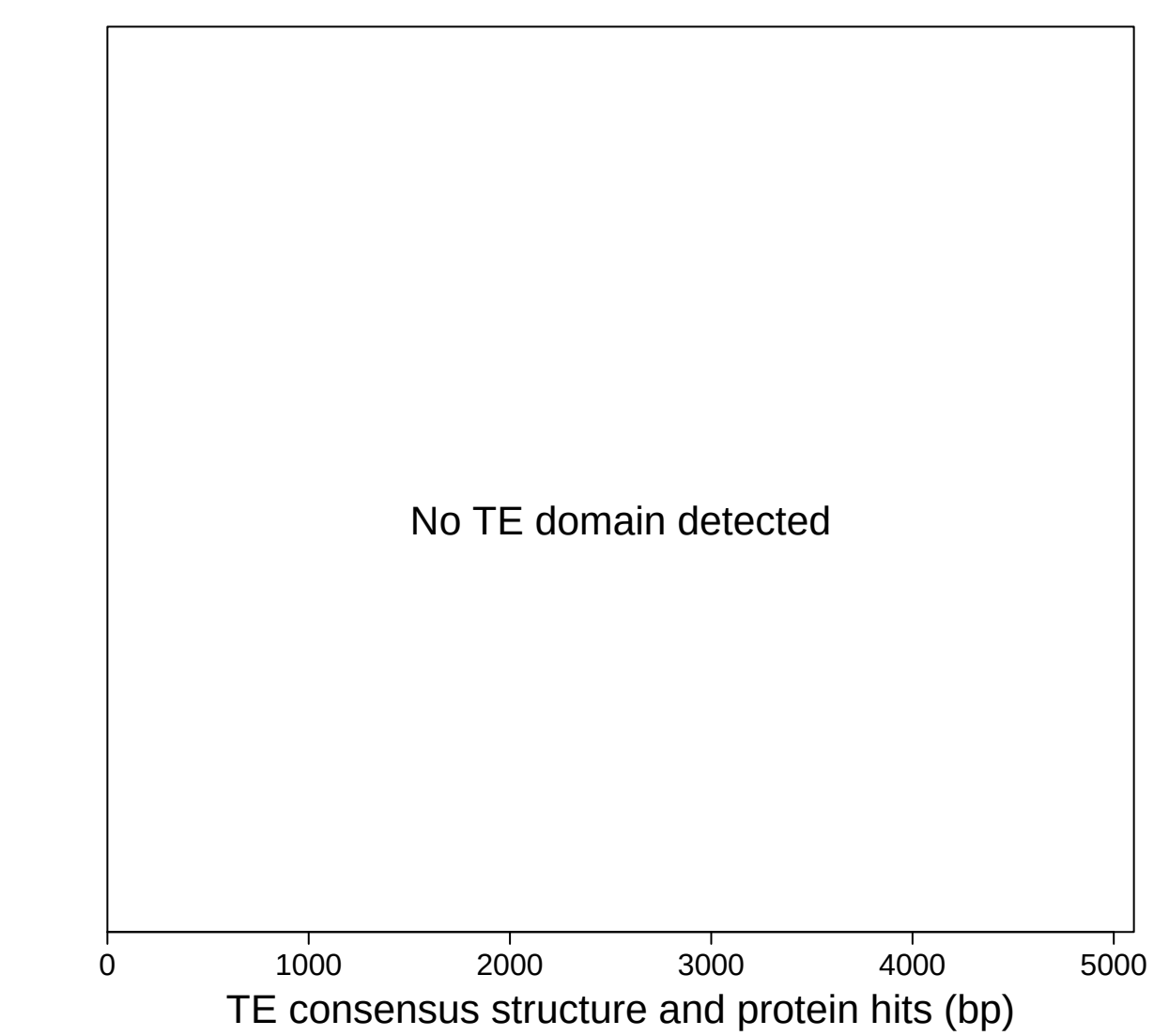
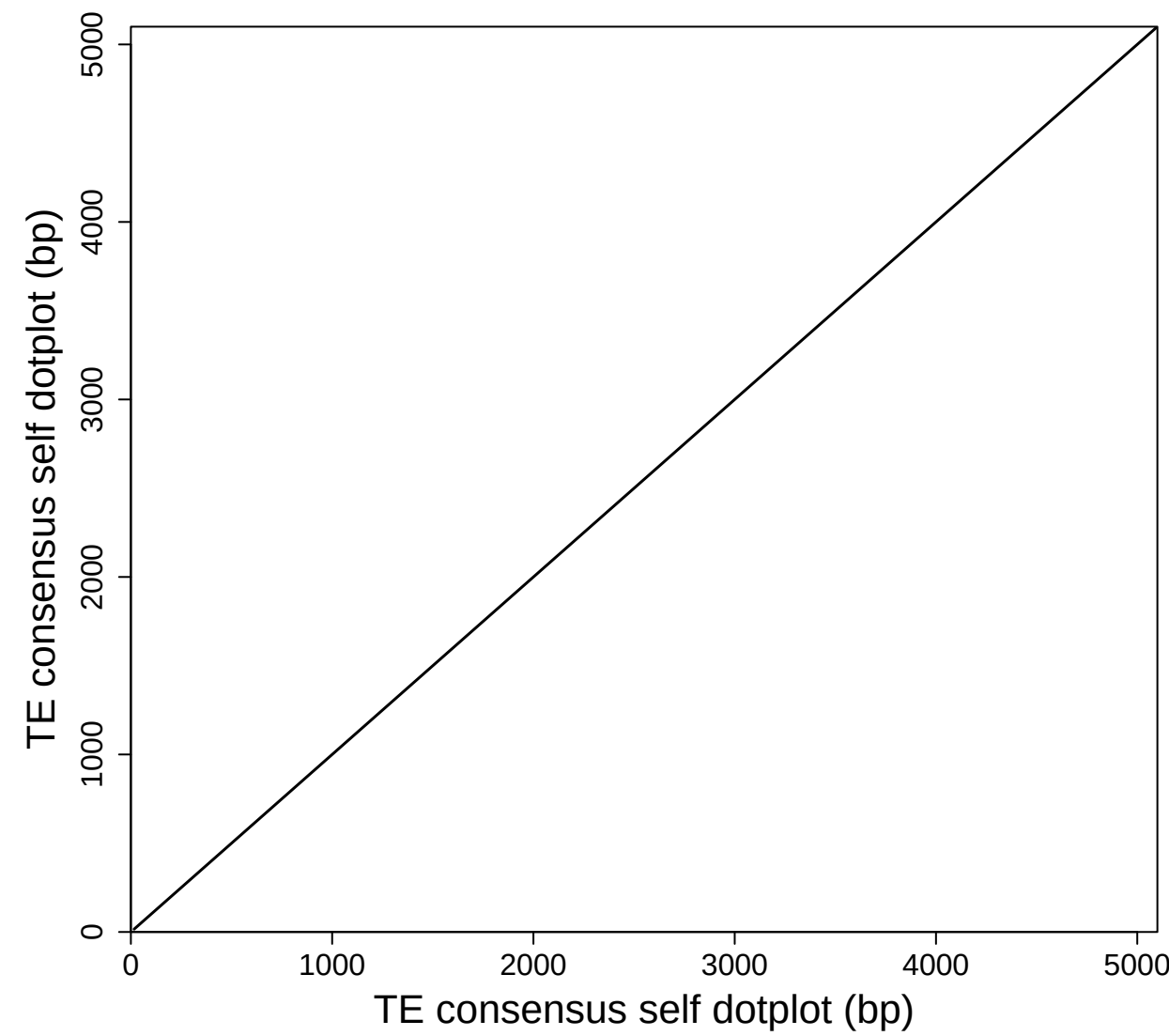
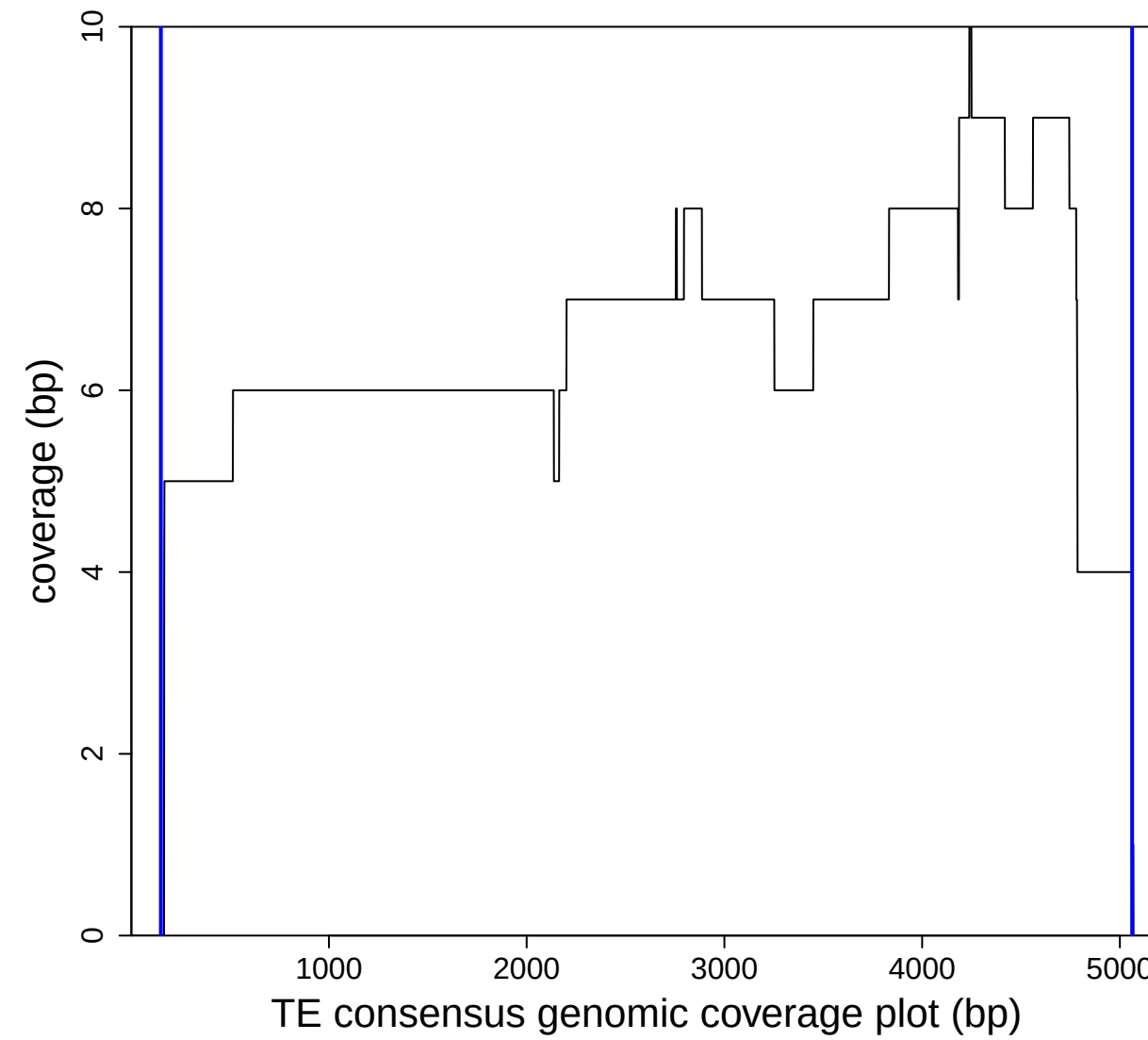
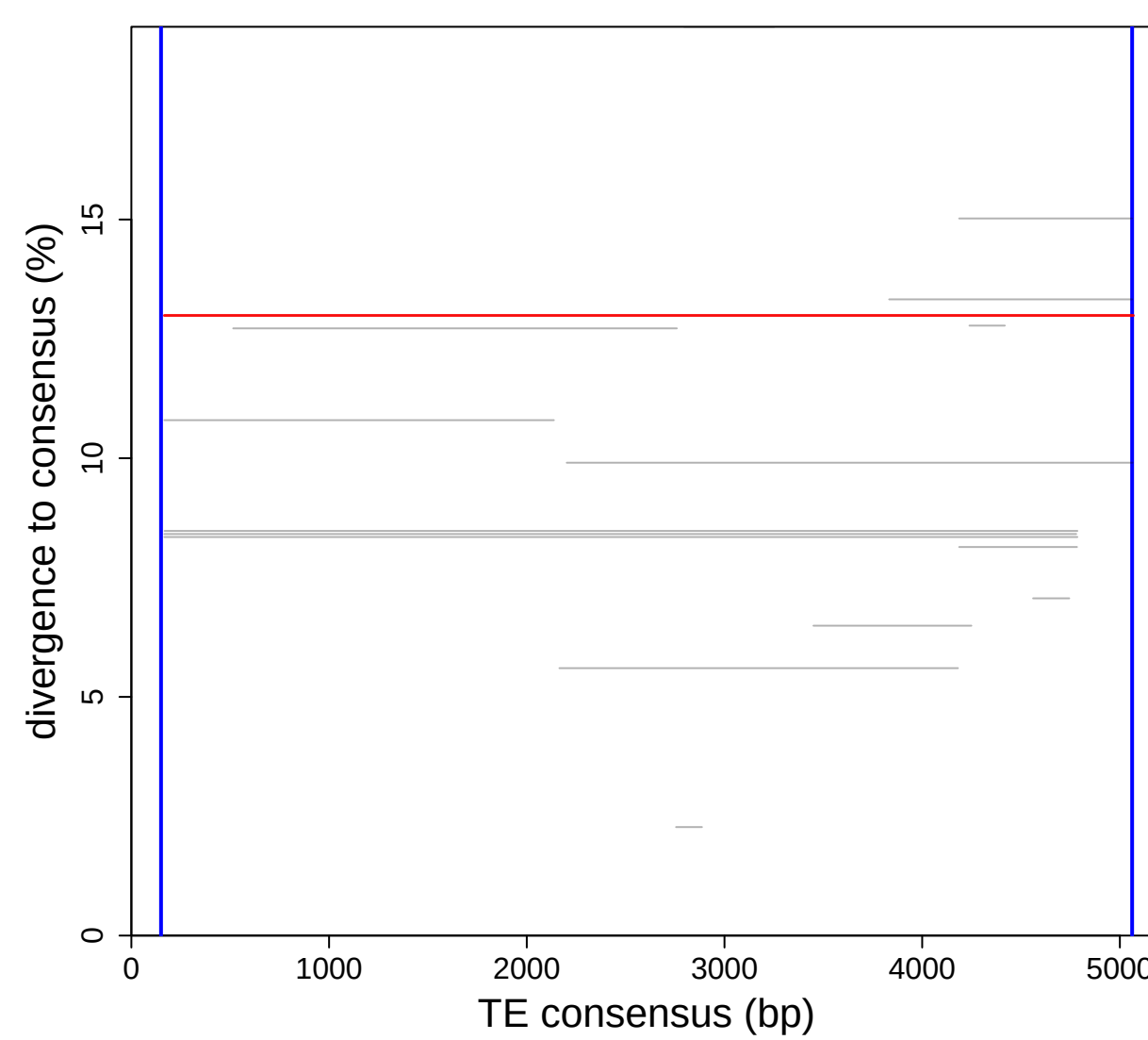
b.bed uf.bed g 1.bed fm 1.bed 0 0 bcln.fa aln.fa cl.fa gs.fa
size: 4755bp; fragments: 33; full length: 0 (>=4279.5bp)

After TEtrimmer 4755 bp

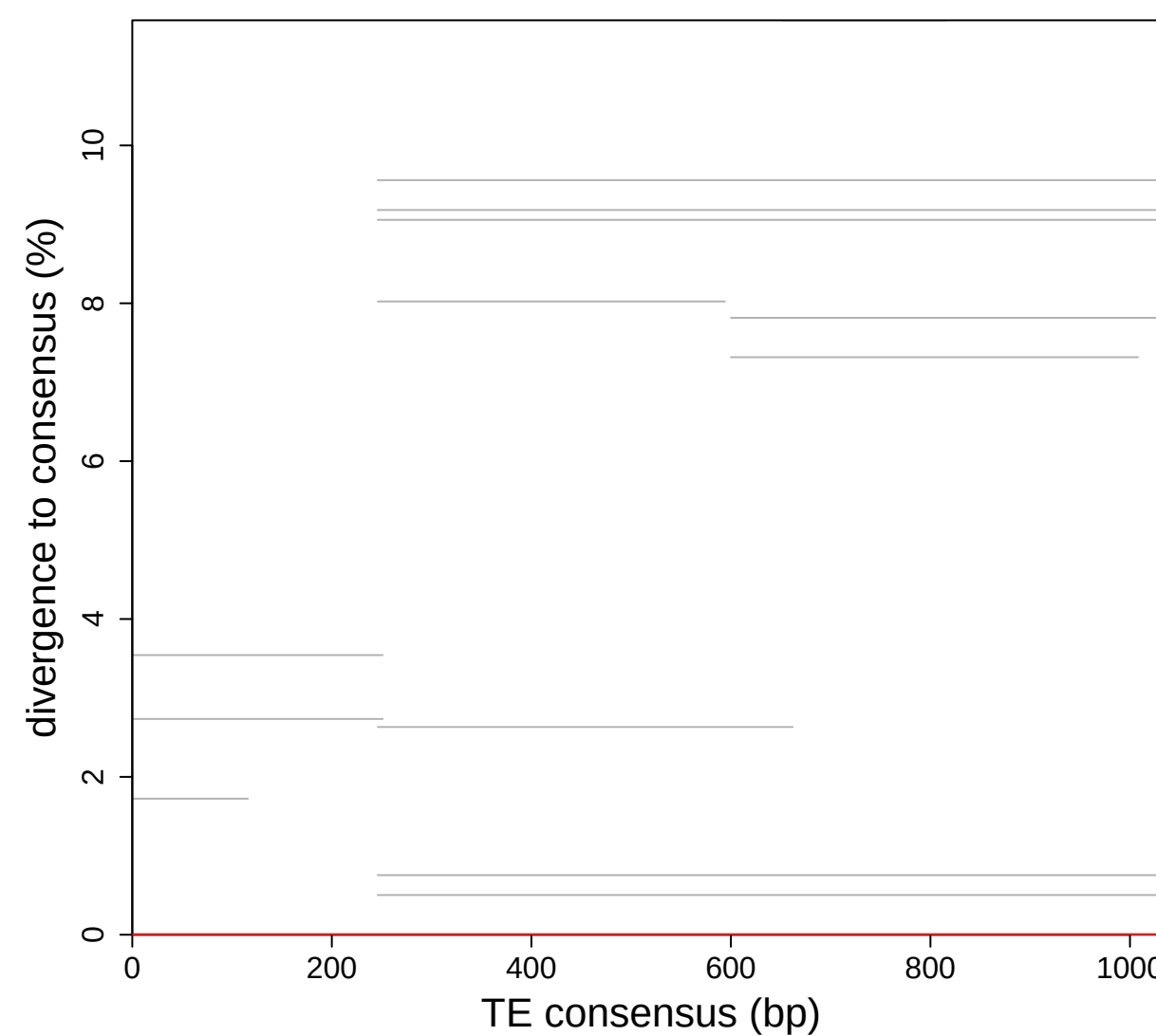


l.fasta.b.bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa
size: 5212bp; fragments: 16; full length: 1 (>=4690.8bp)

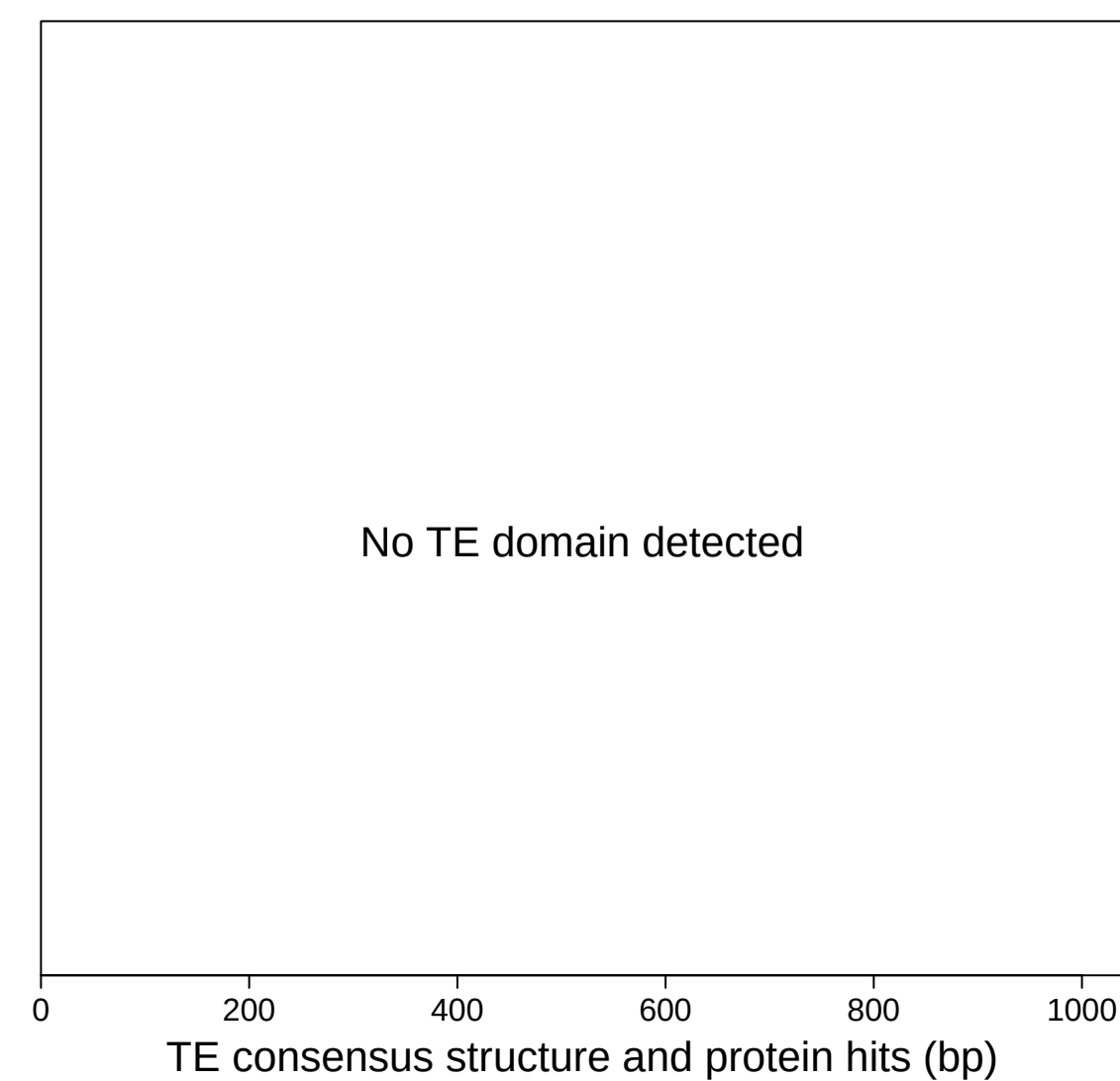
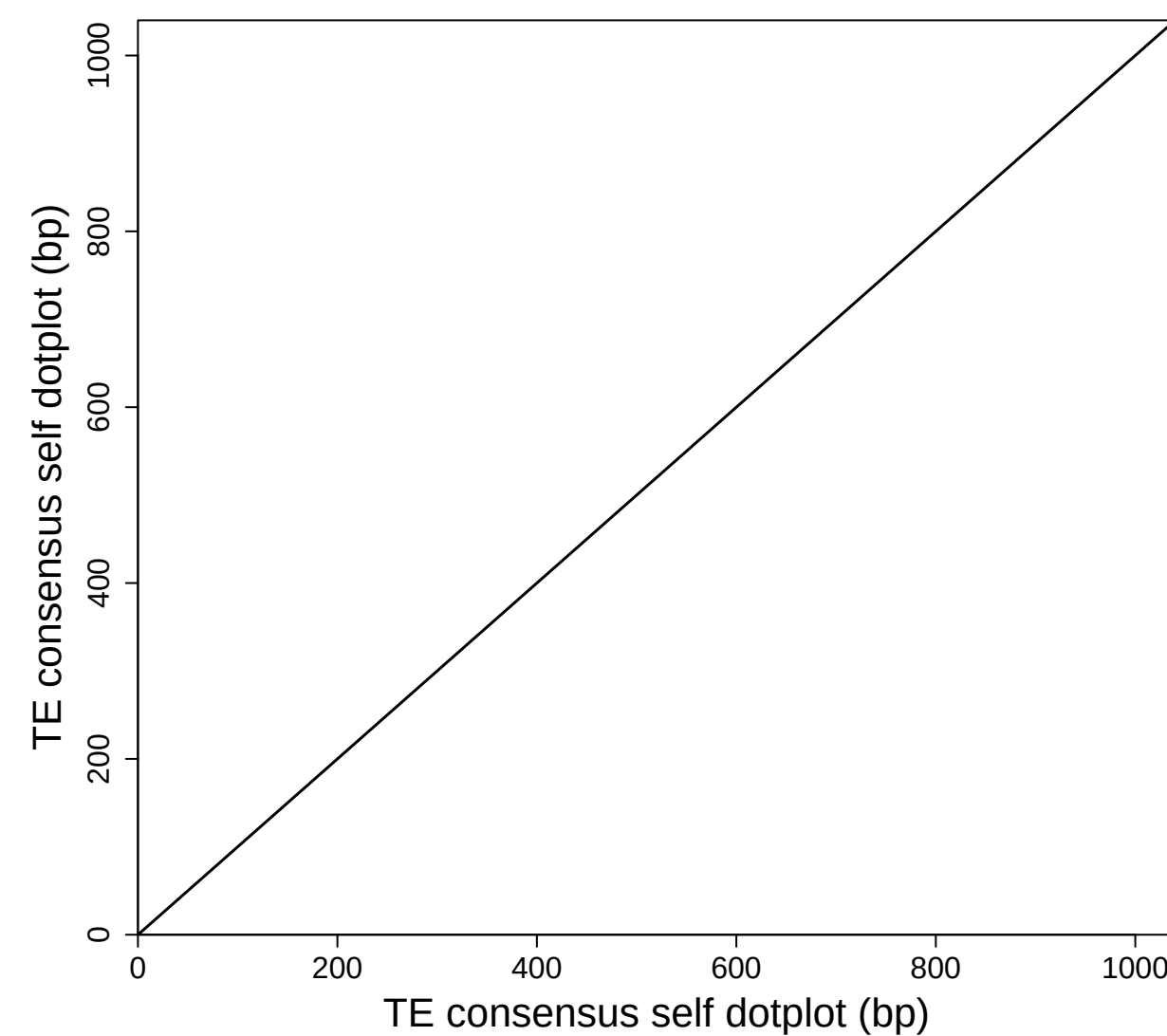
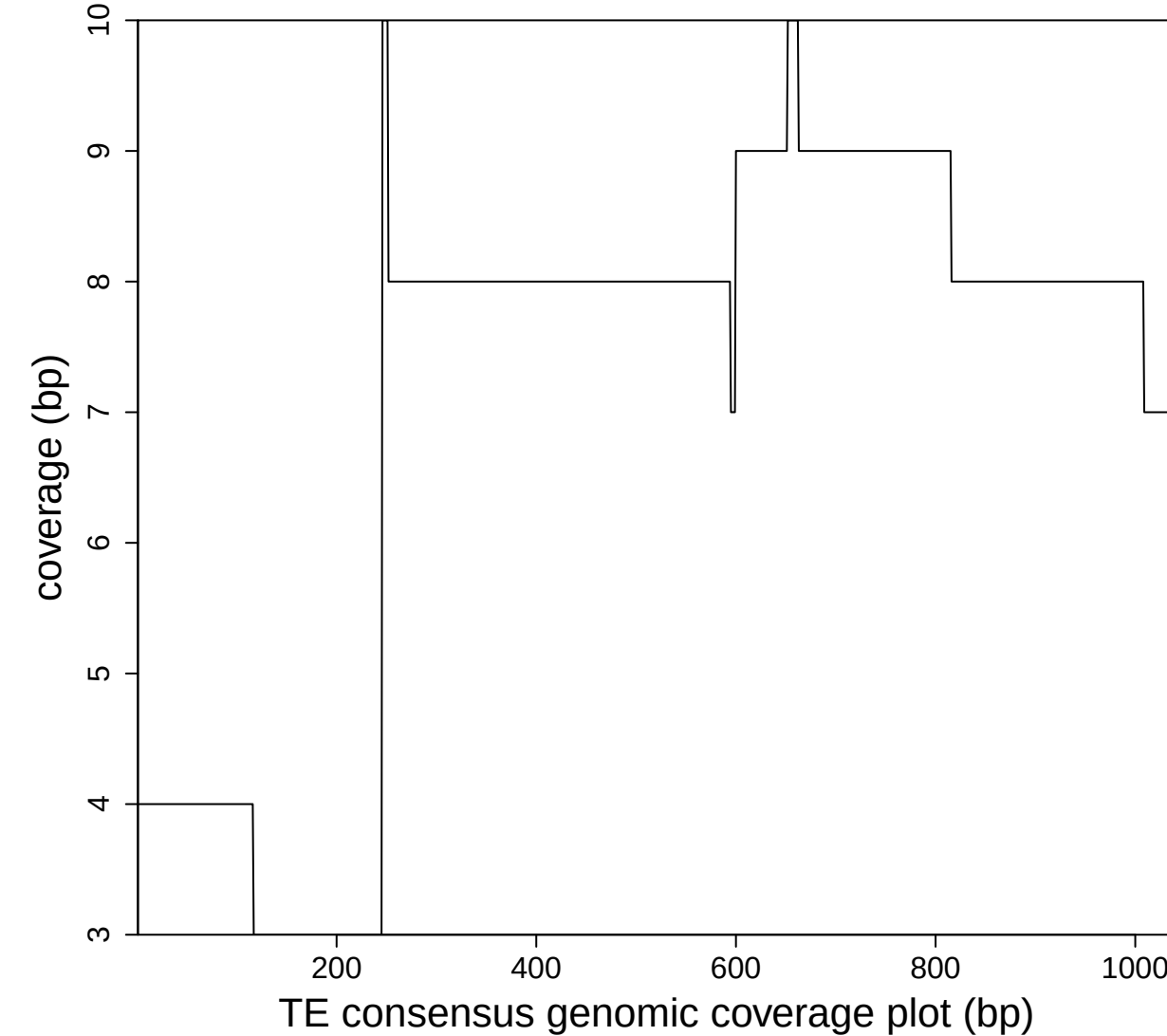
After TEtrimmer Extended plot Blue lines are boundaries



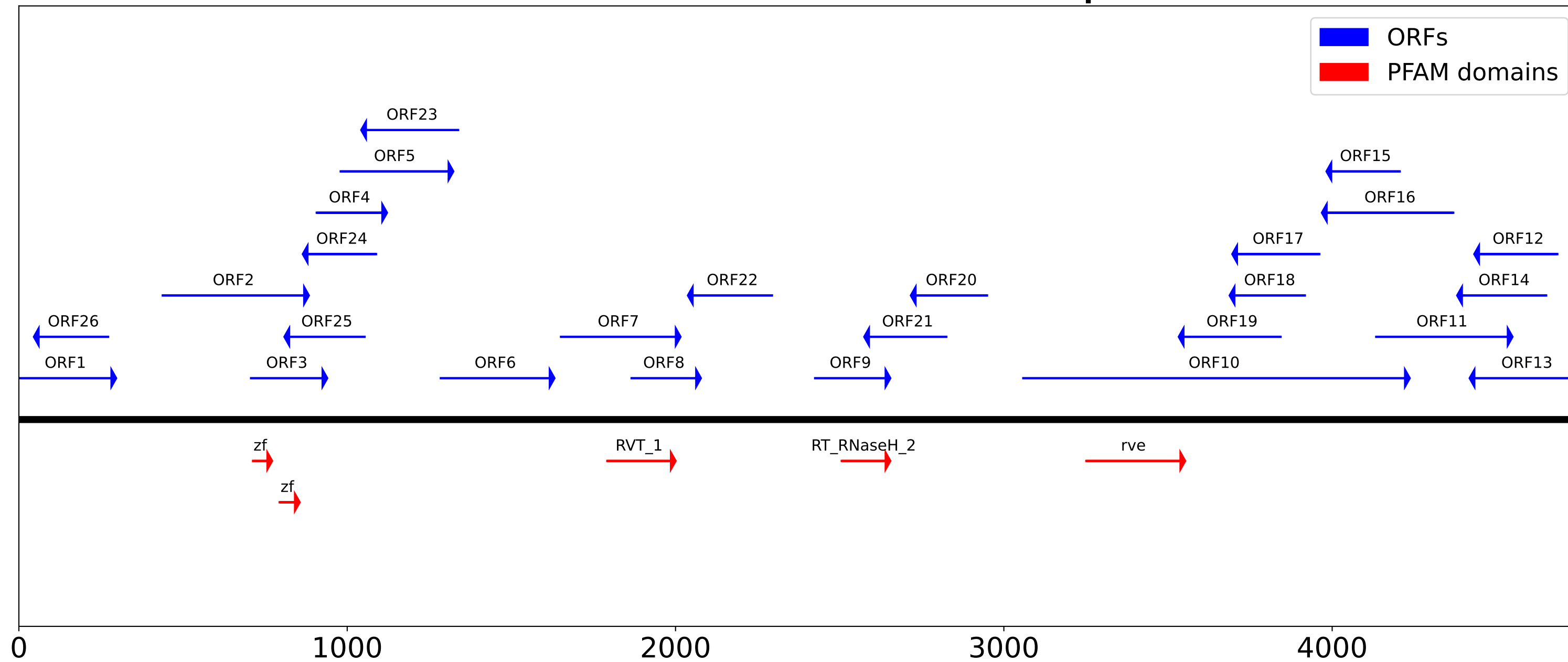
TE: ltr_1_family_884
size: 1040bp; fragments: 14; full length: 1 (≥ 936 bp)



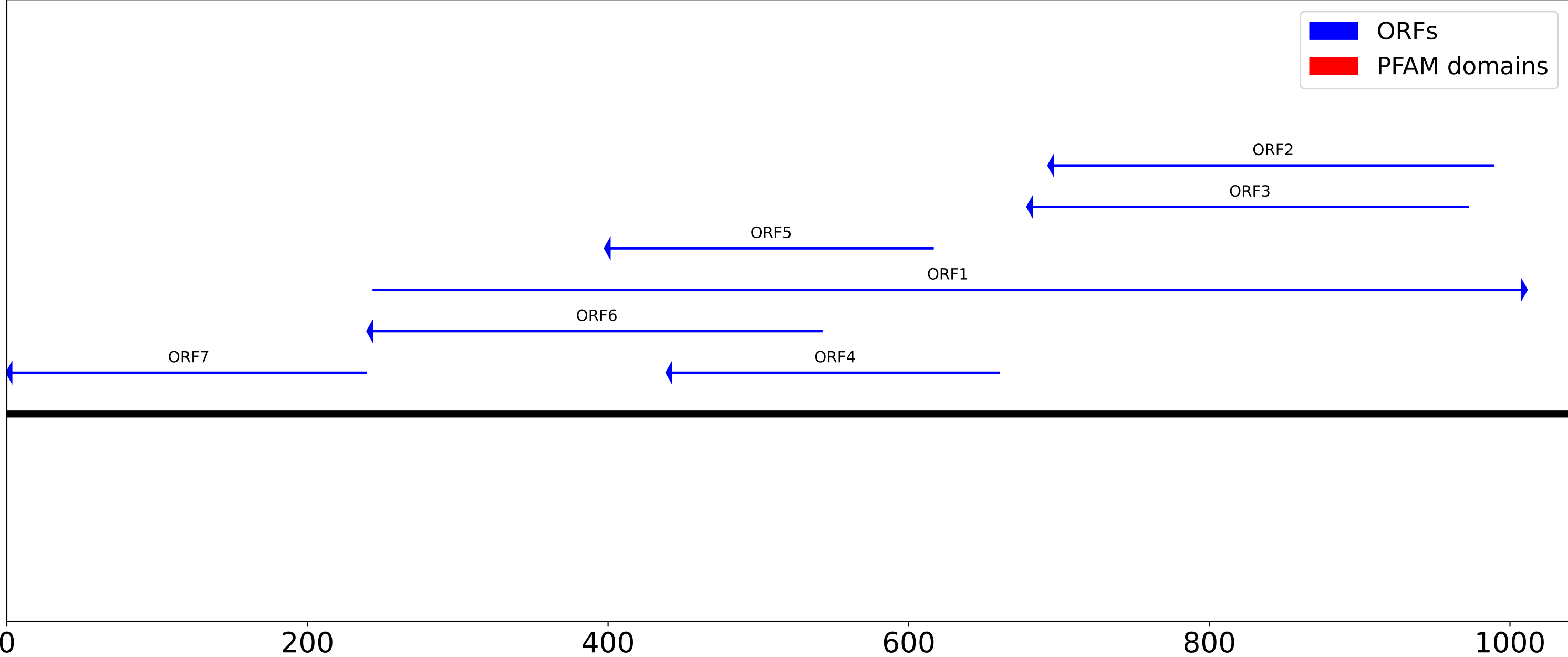
Before TEtrimmer 1040 bp



After TEtrimmer ORF and PFAM domain plot



Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

onsensus before TEtrimmer (bp)

