

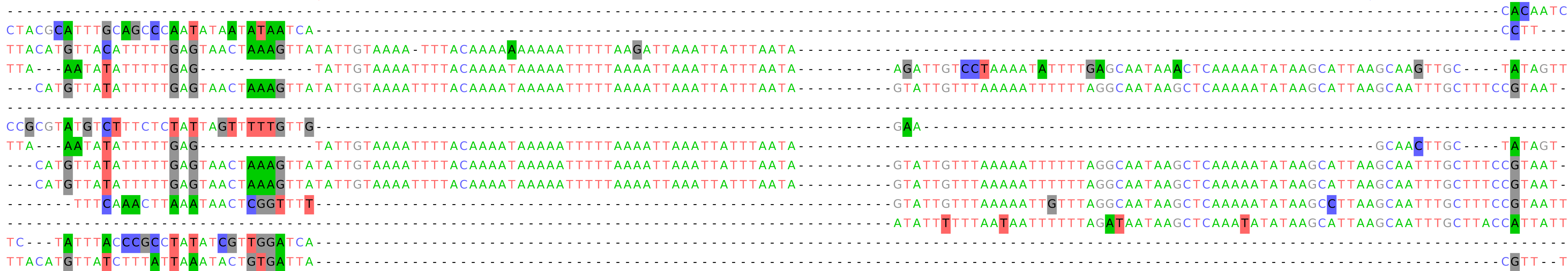
Start crop Point

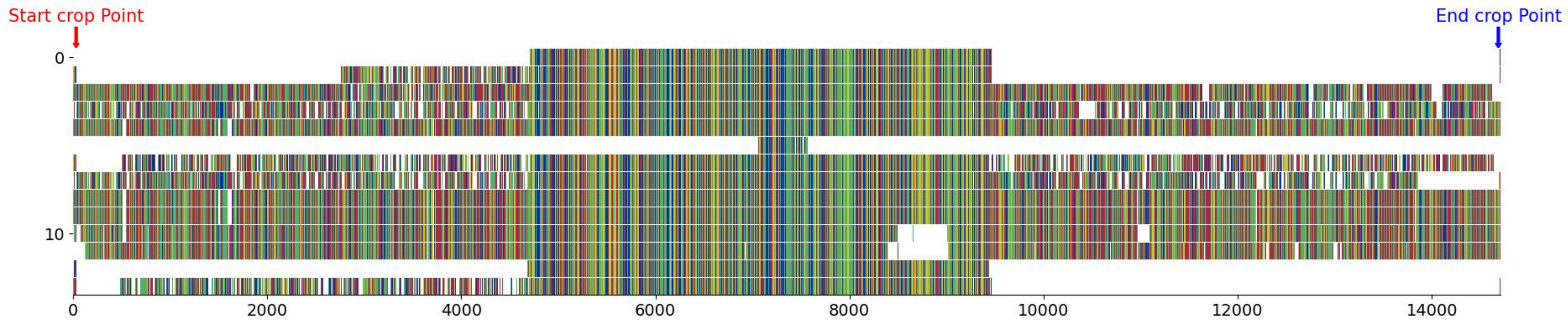


End crop Point



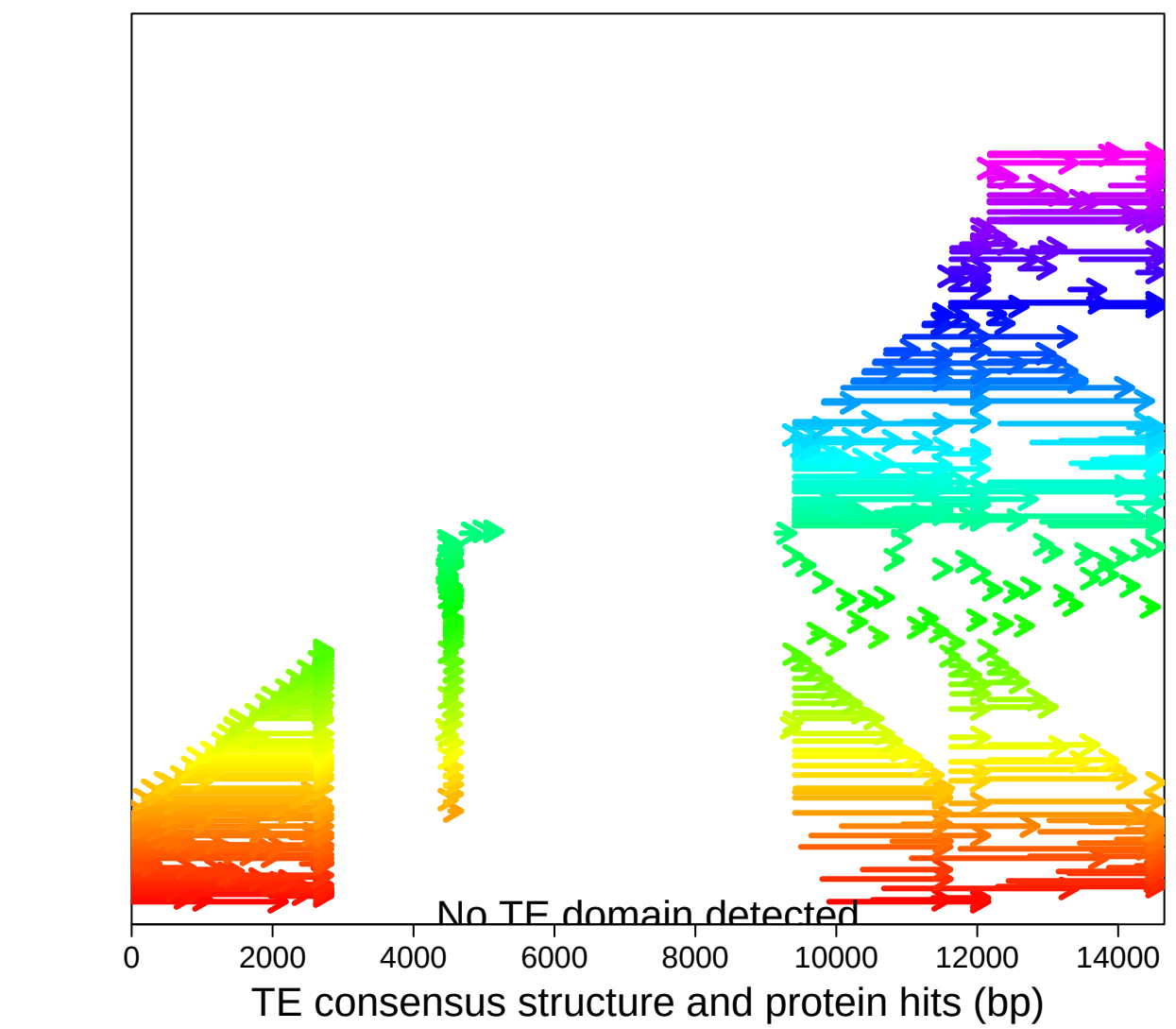
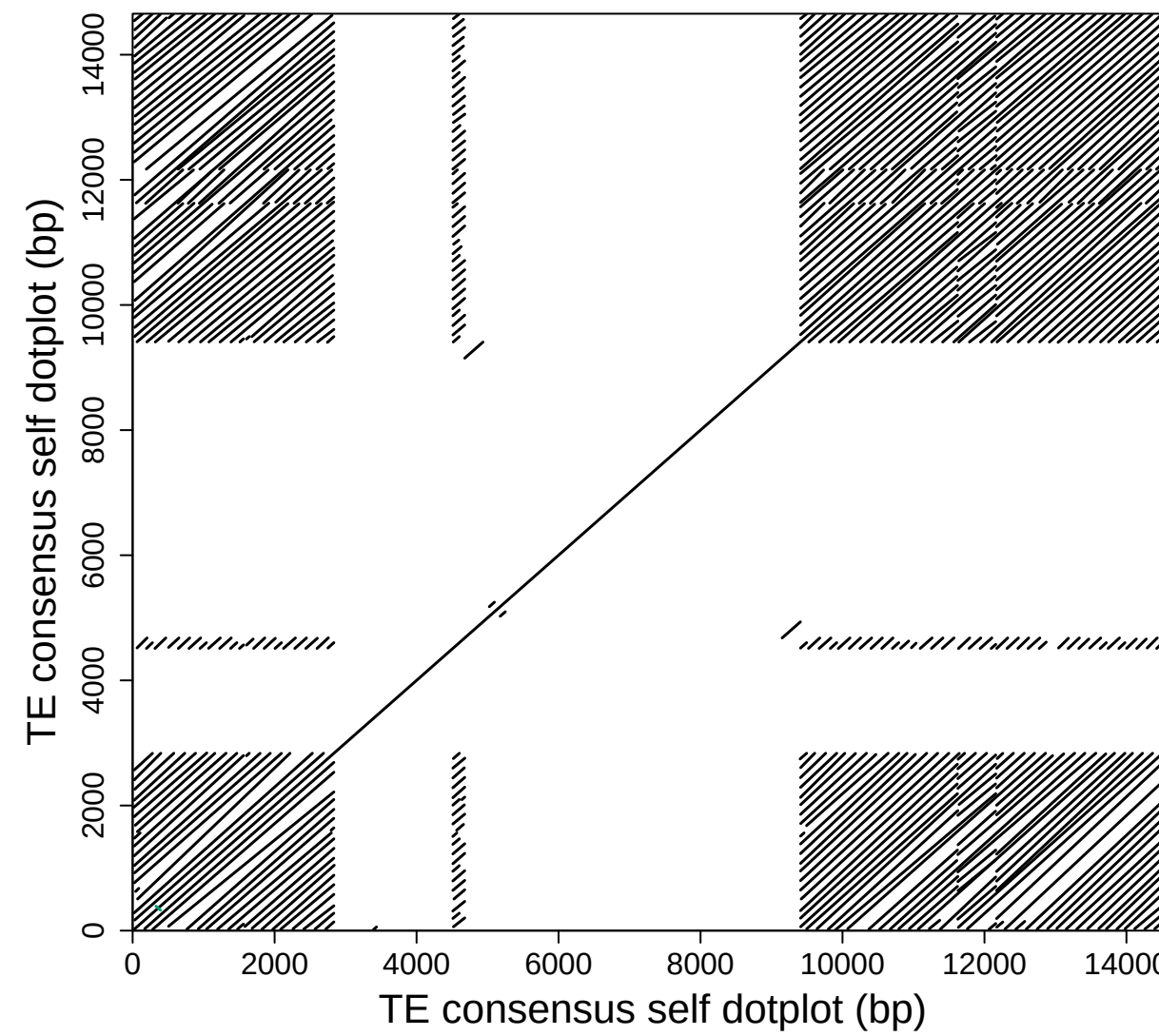
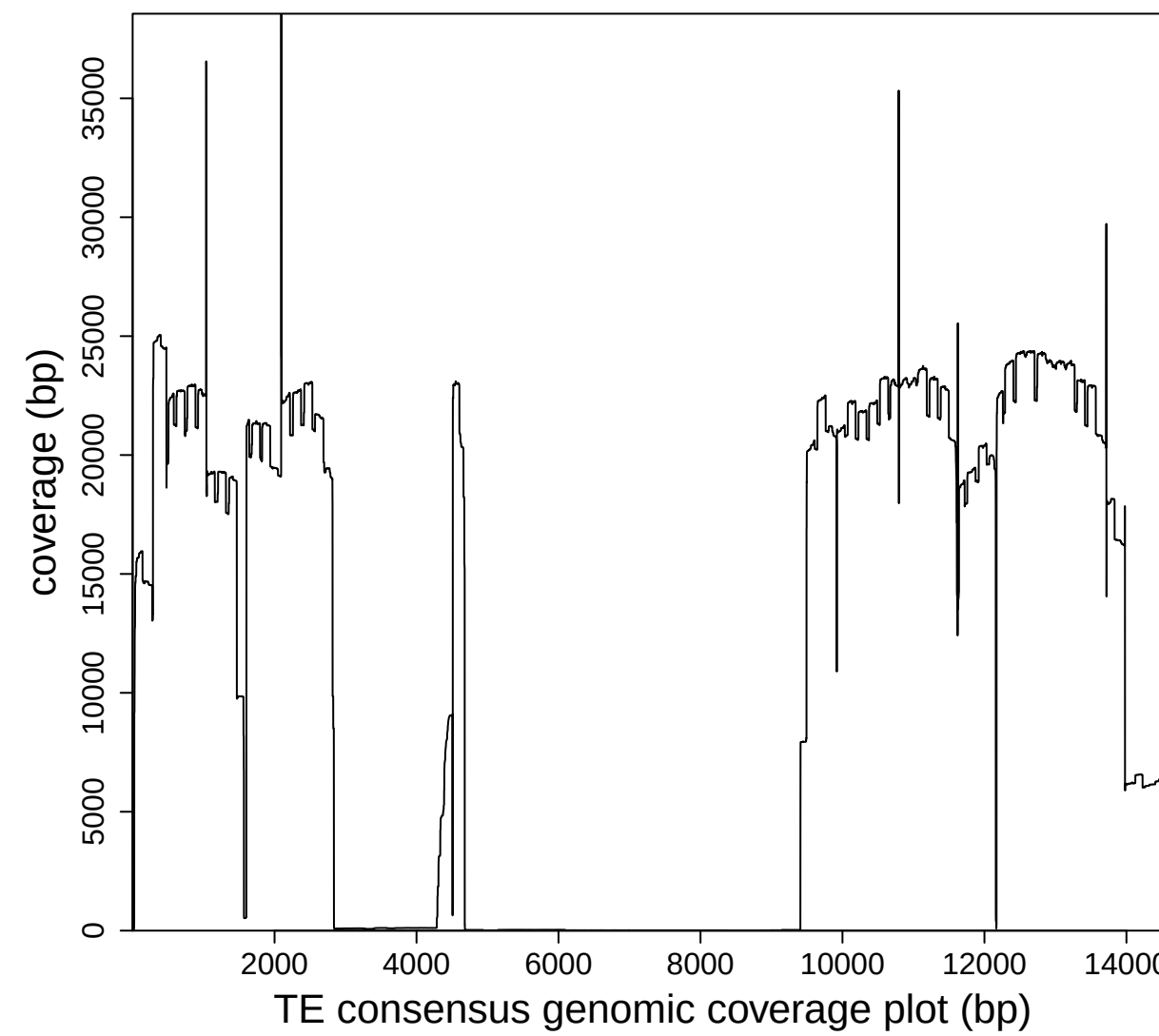
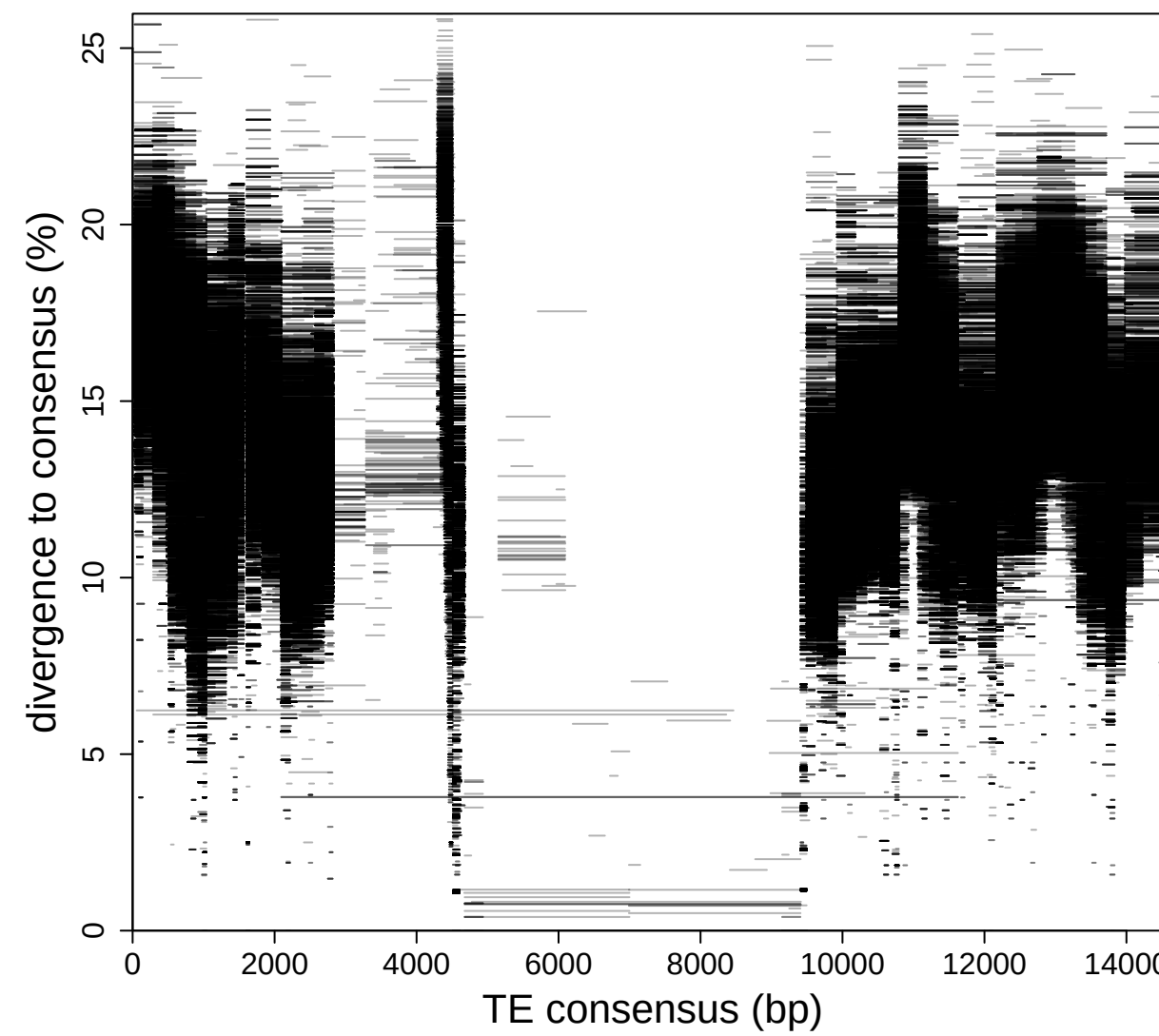
MSA length = 14655





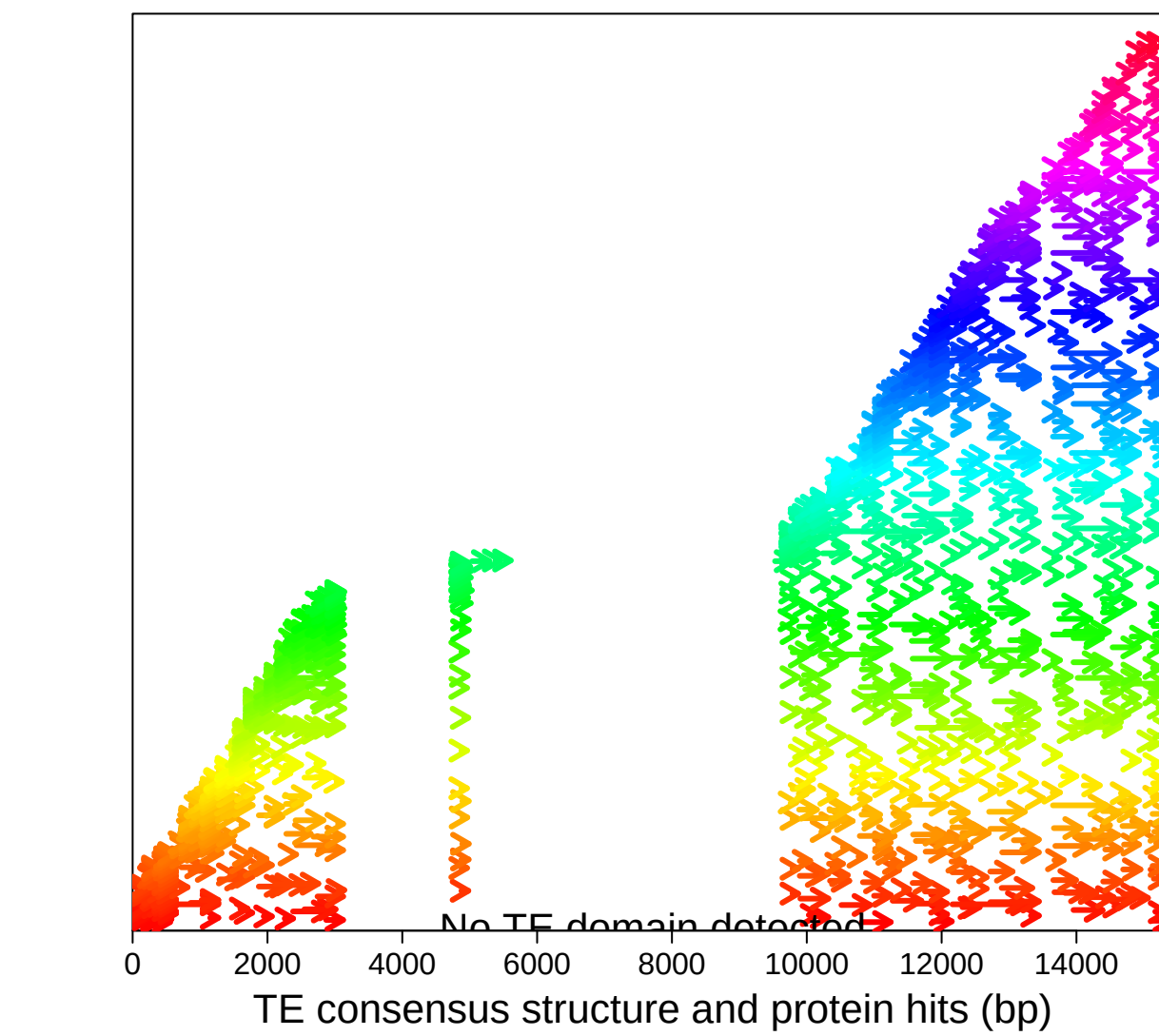
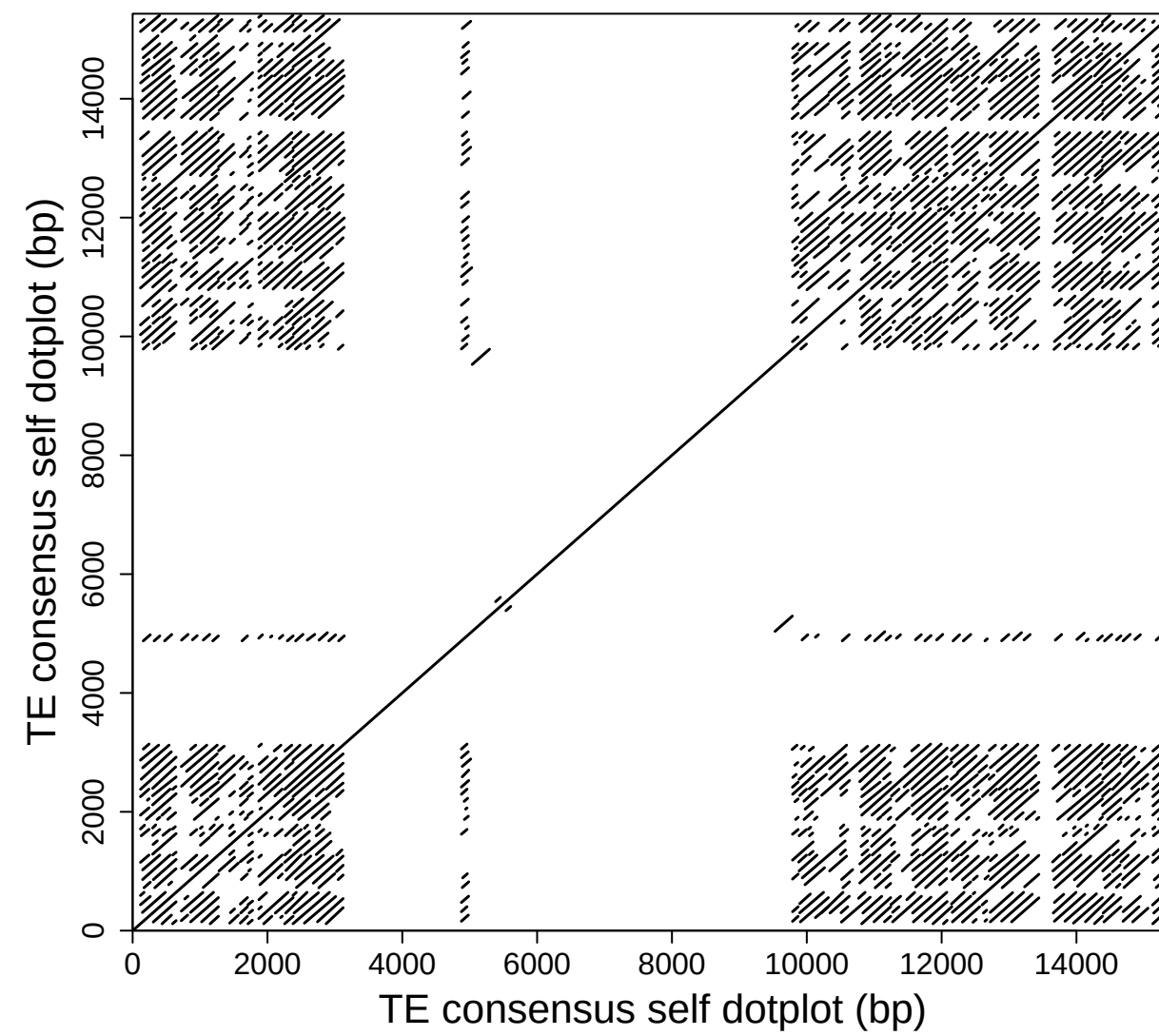
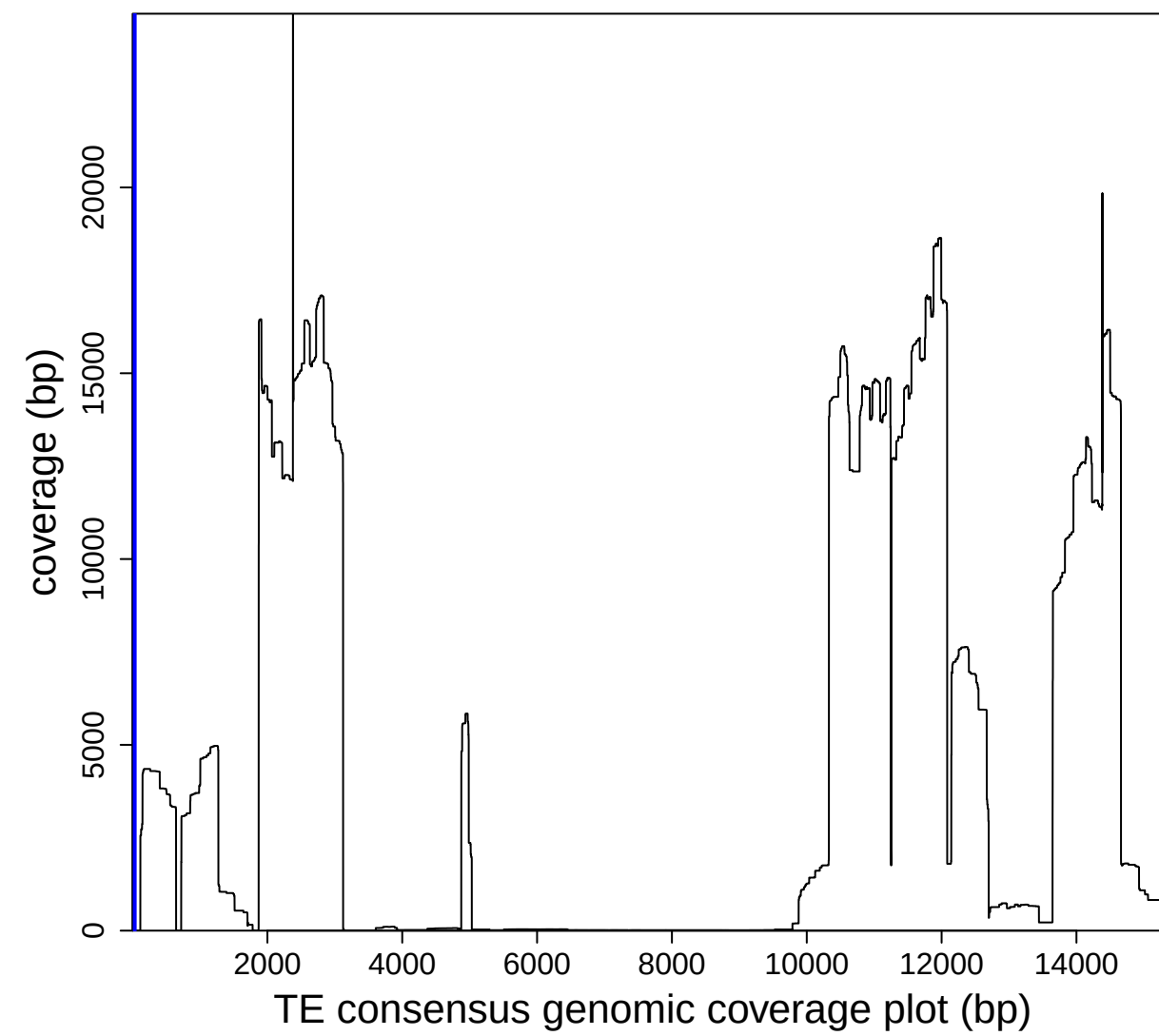
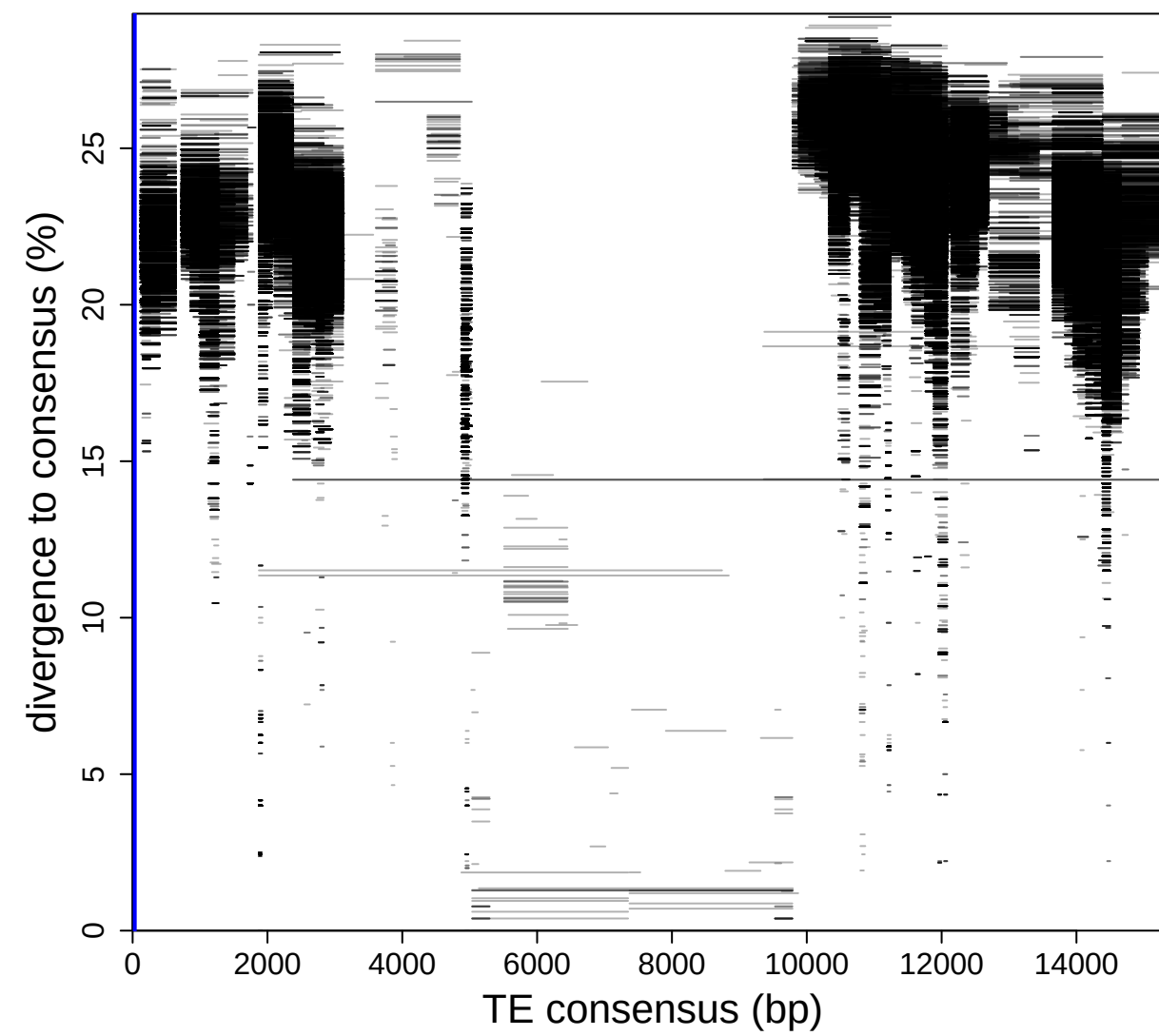
bed uf.bed g 3.bed fm 1.bed 0 0 bcln.fa aln.fa cl.fa gs.fa ce
size: 14655bp; fragments: 304683; full length: 0 (>=13189.5bp)

After TEtrimmer 14655 bp

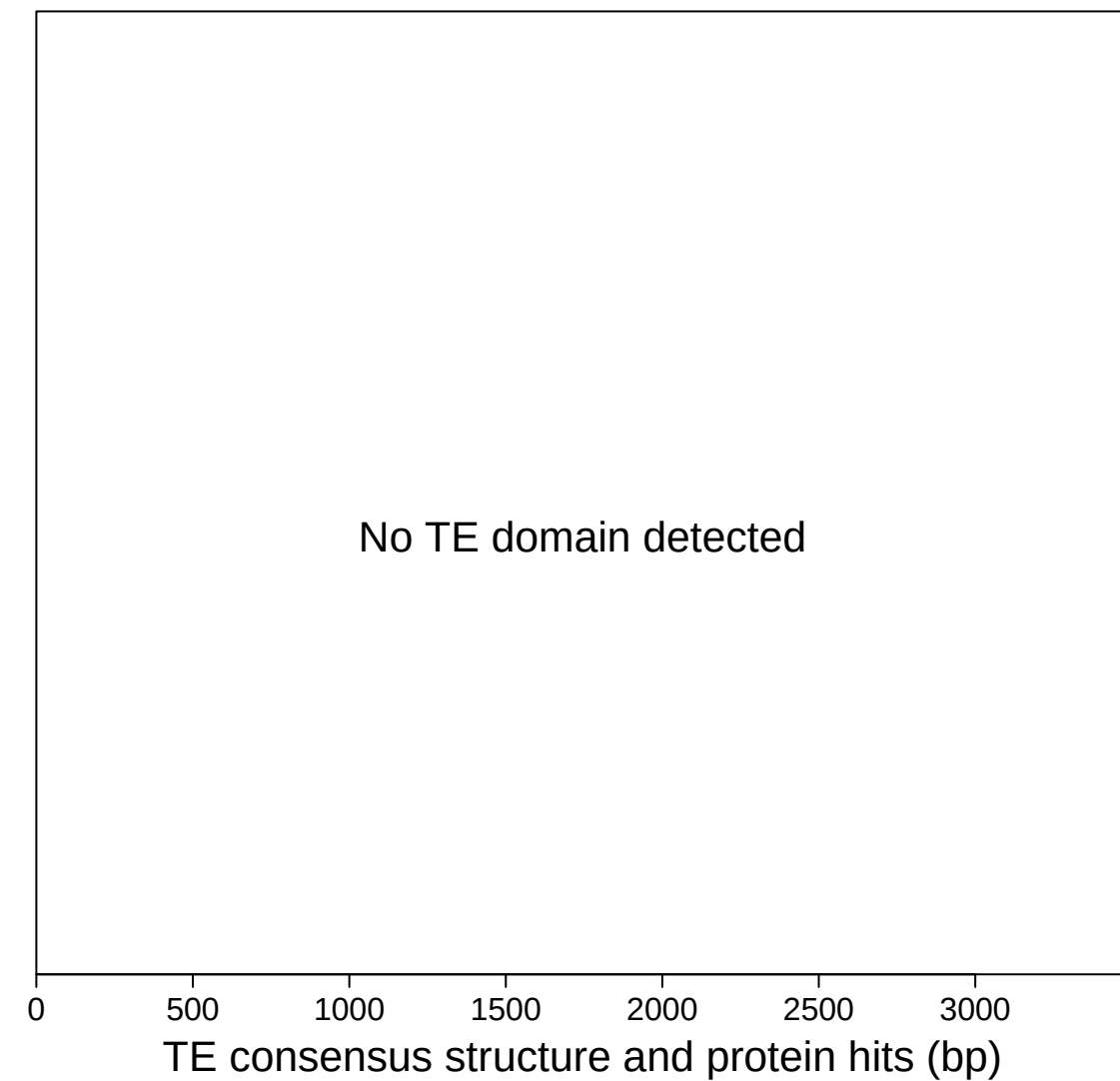
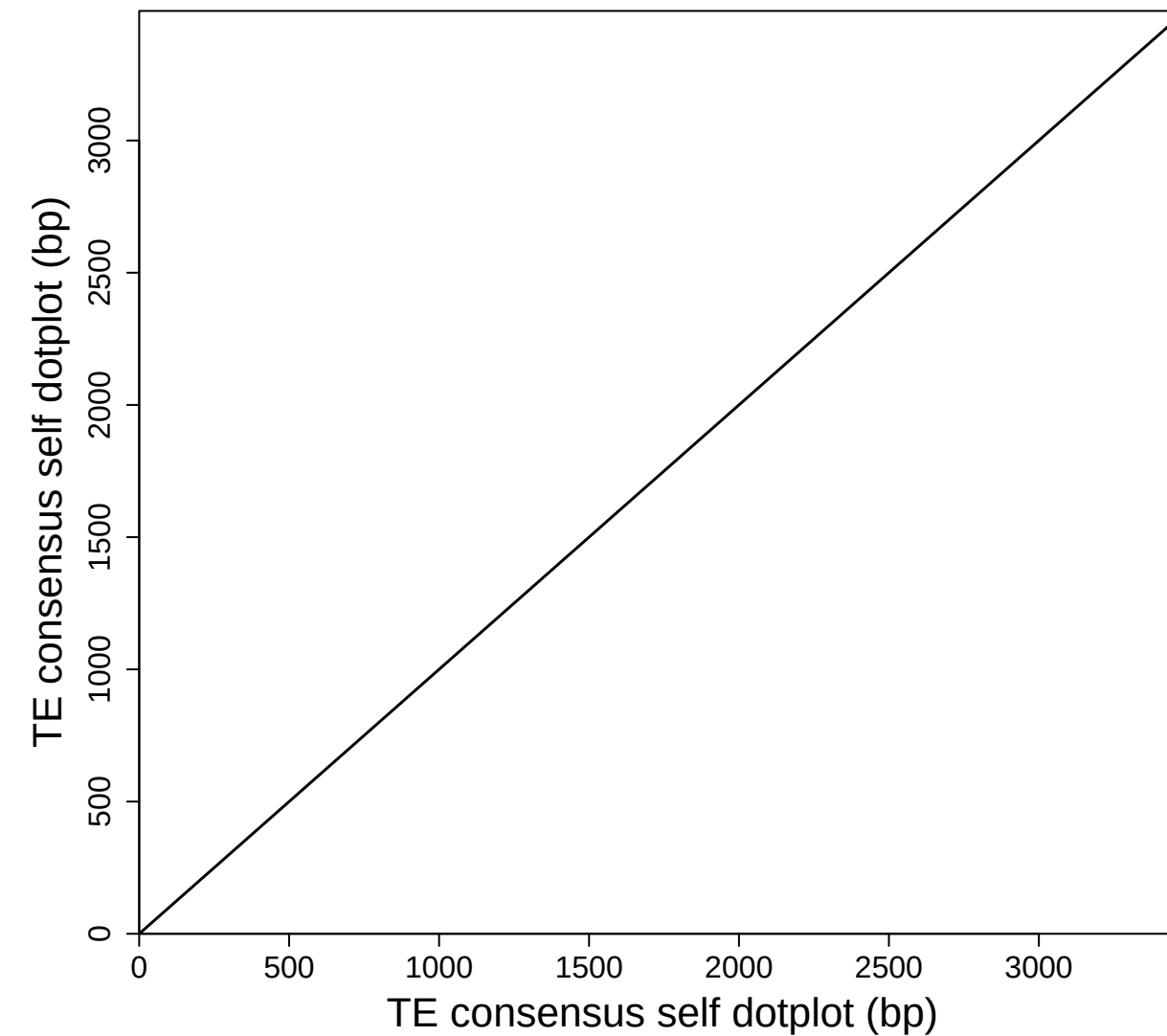
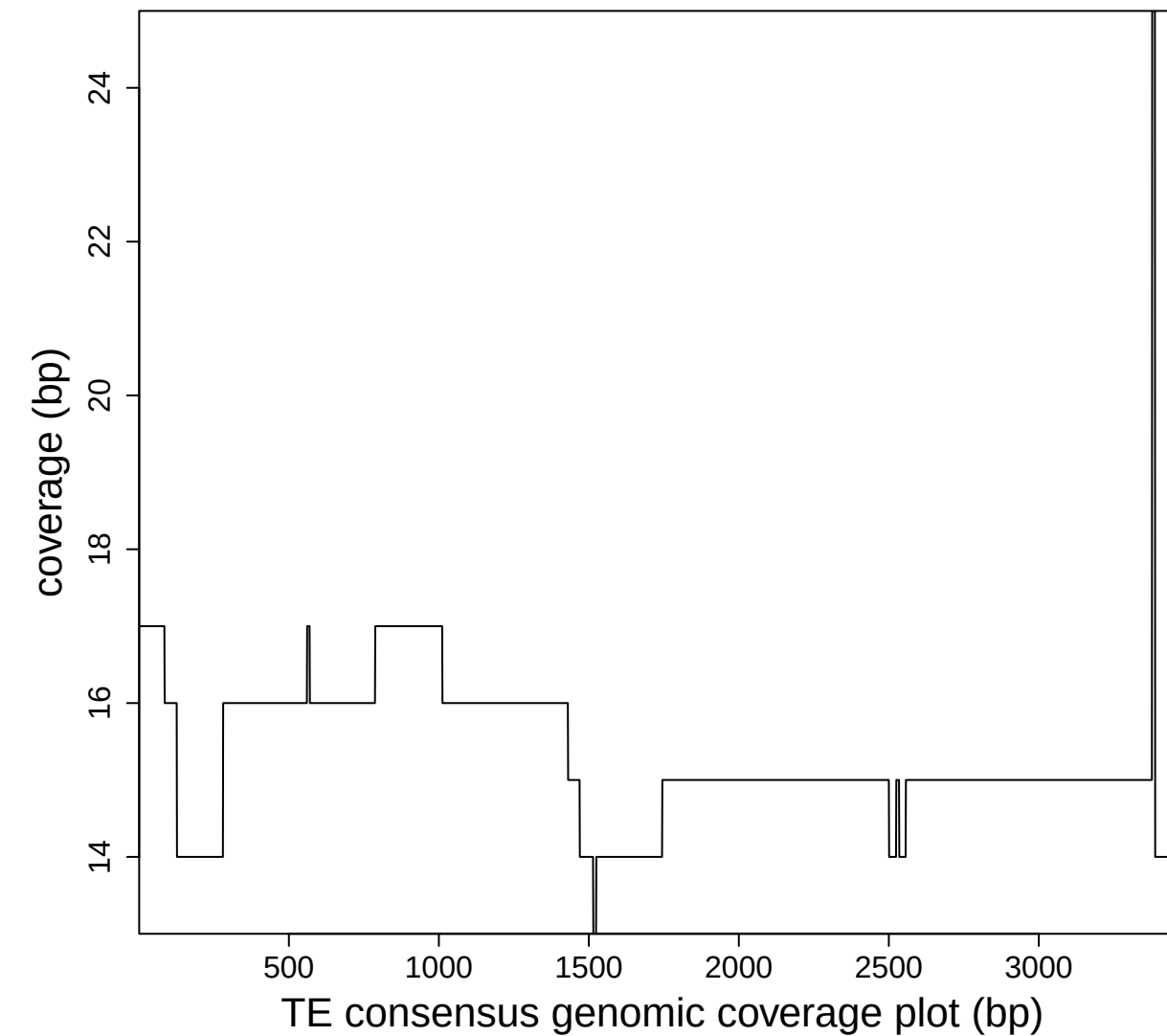
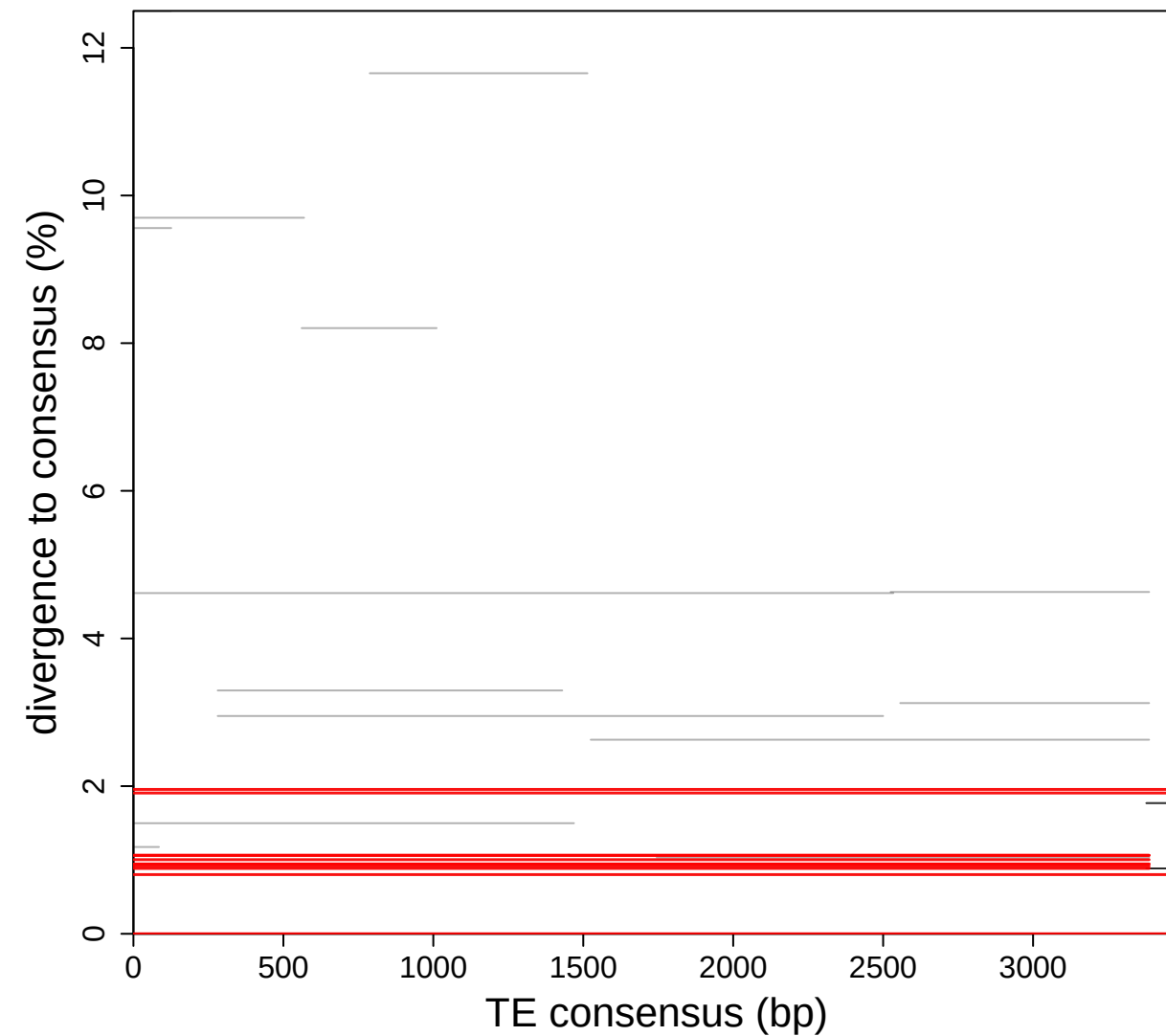


y_1021.fasta.b.bed_uf.bed_g_3.bed_fm_1.bed_0_0_bcln.fa_aln.f
size: 15431bp; fragments: 123033; full length: 0 (>=13887.9bp

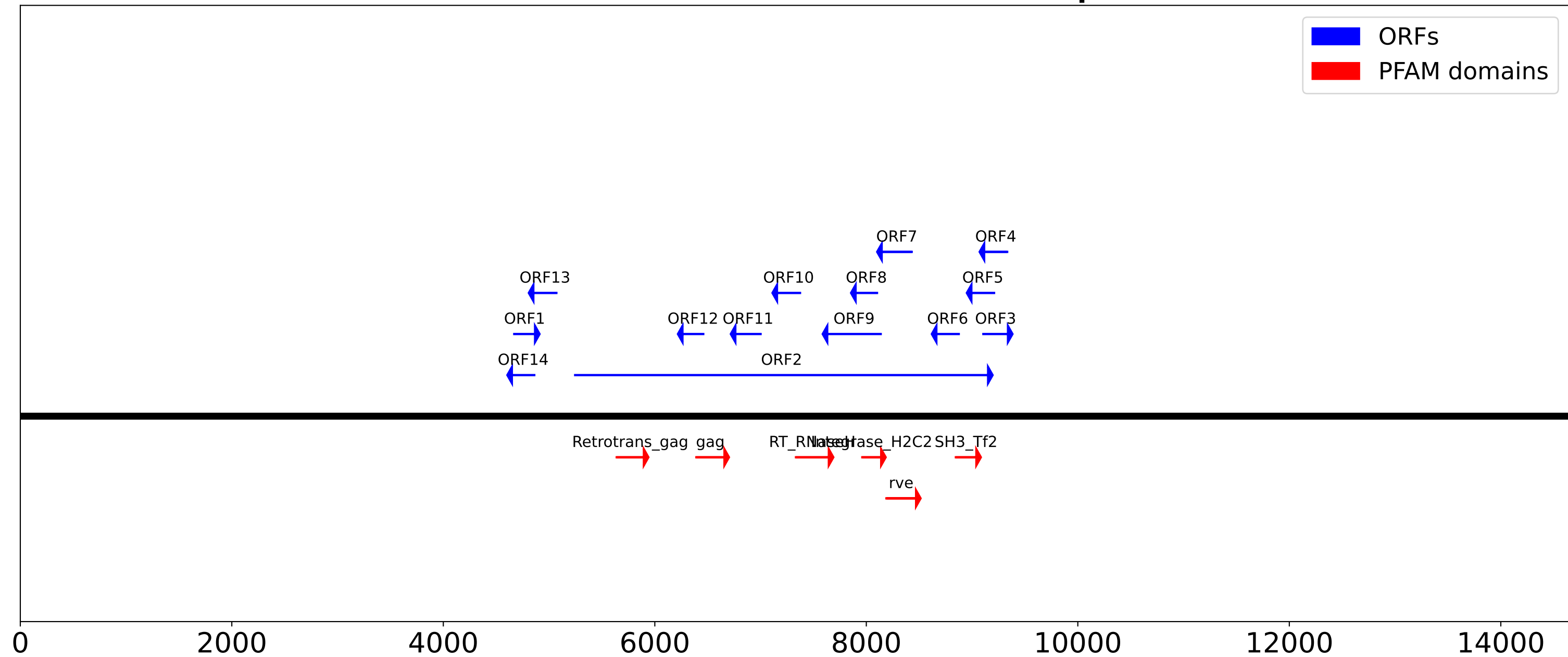
After TEtrimmer Extended plot Blue lines are boundaries



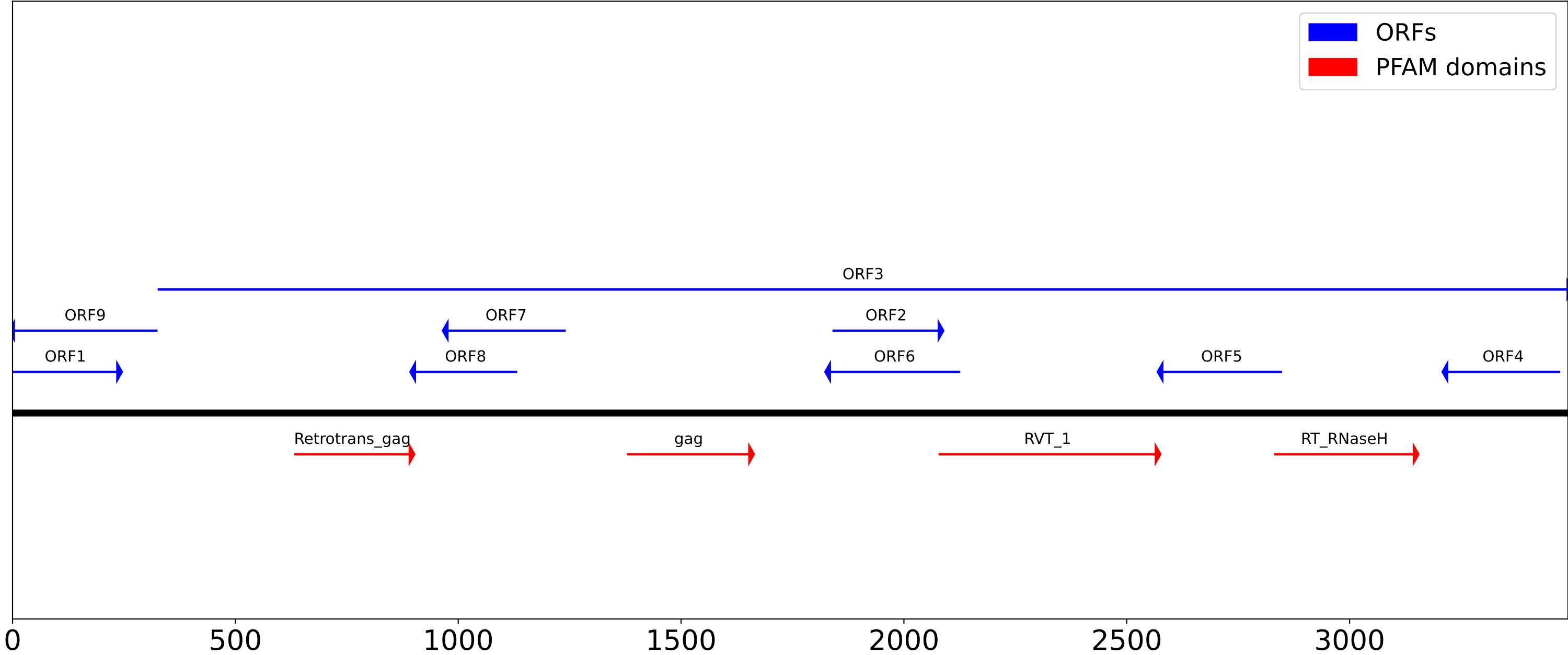
TE: ltr_1_family_1021
size: 3490bp; fragments: 35; full length: 11 (≥ 3141 bp)



After TEtrimmer ORF and PFAM domain plot



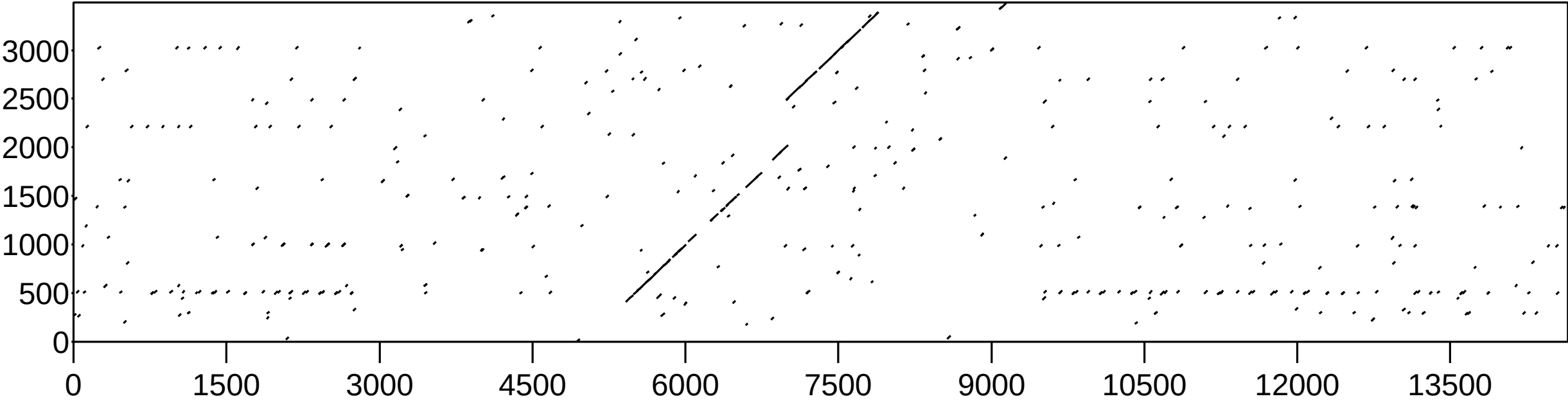
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)