

Start crop Point

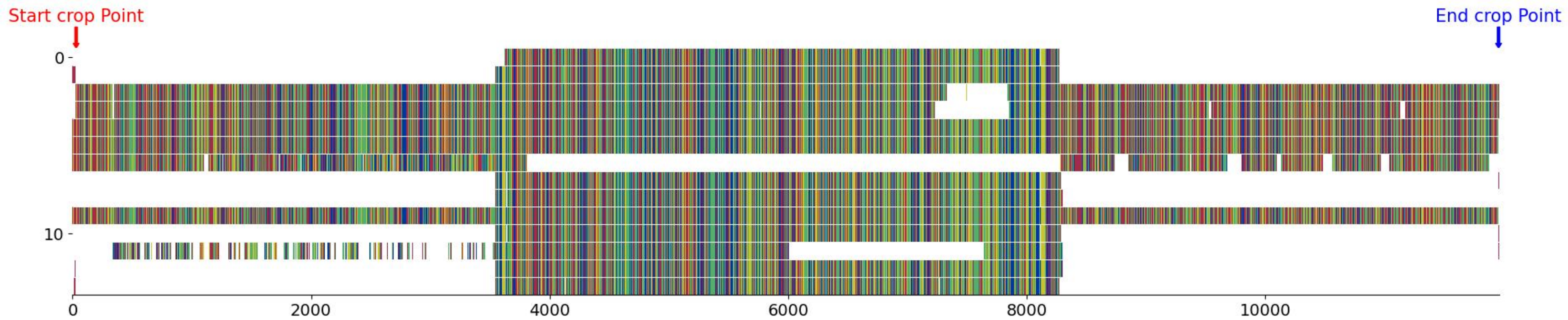


End crop Point



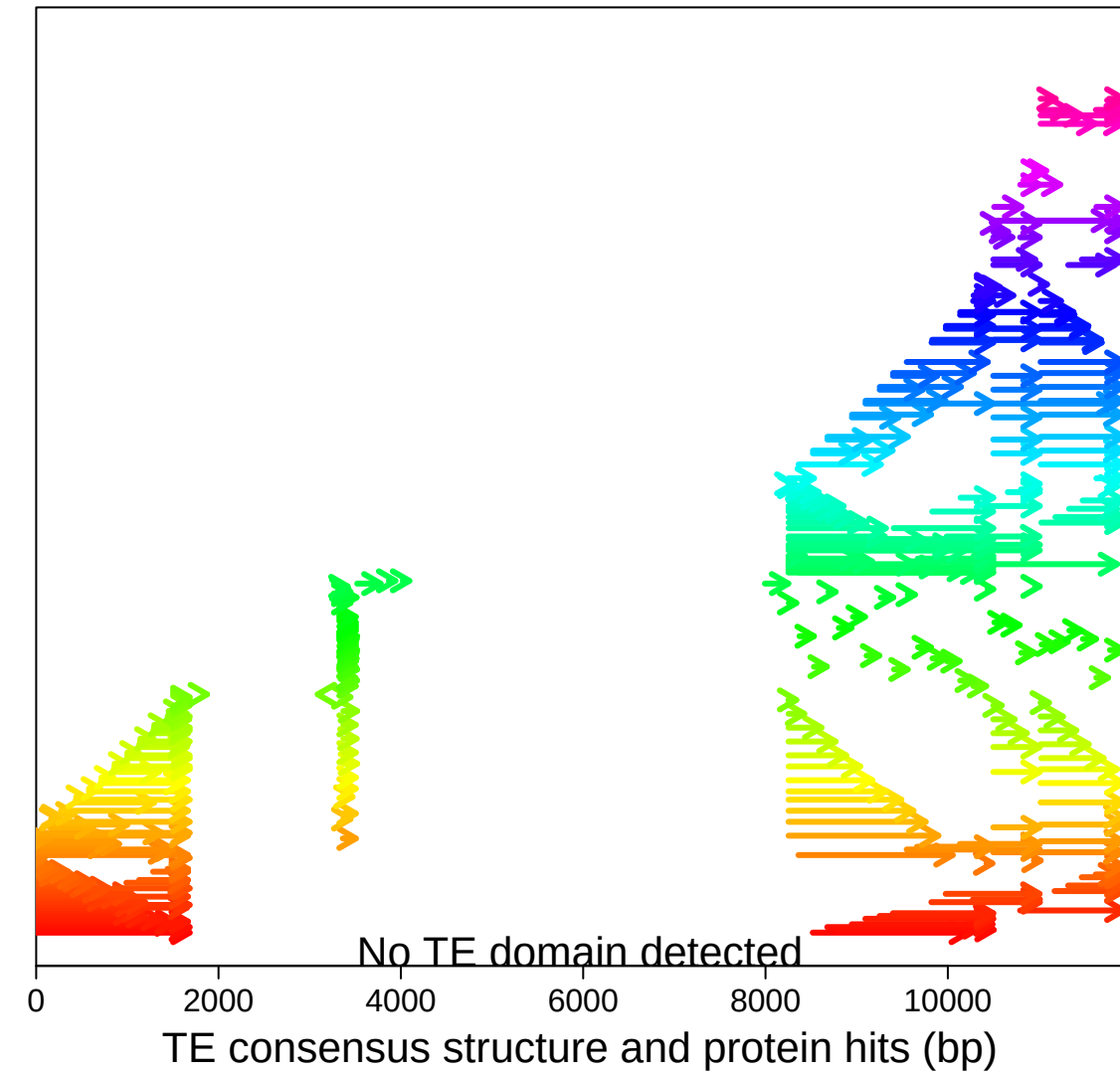
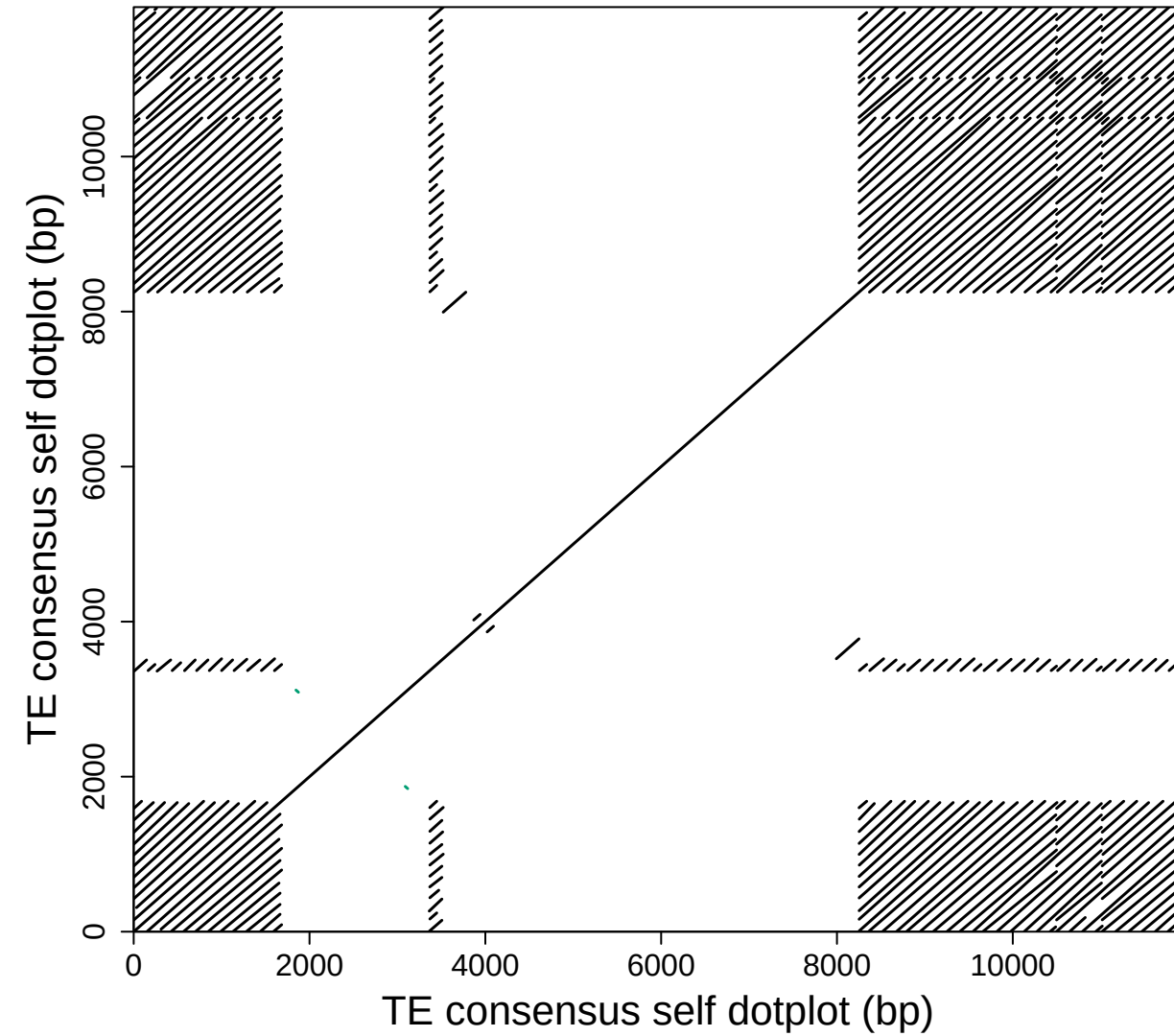
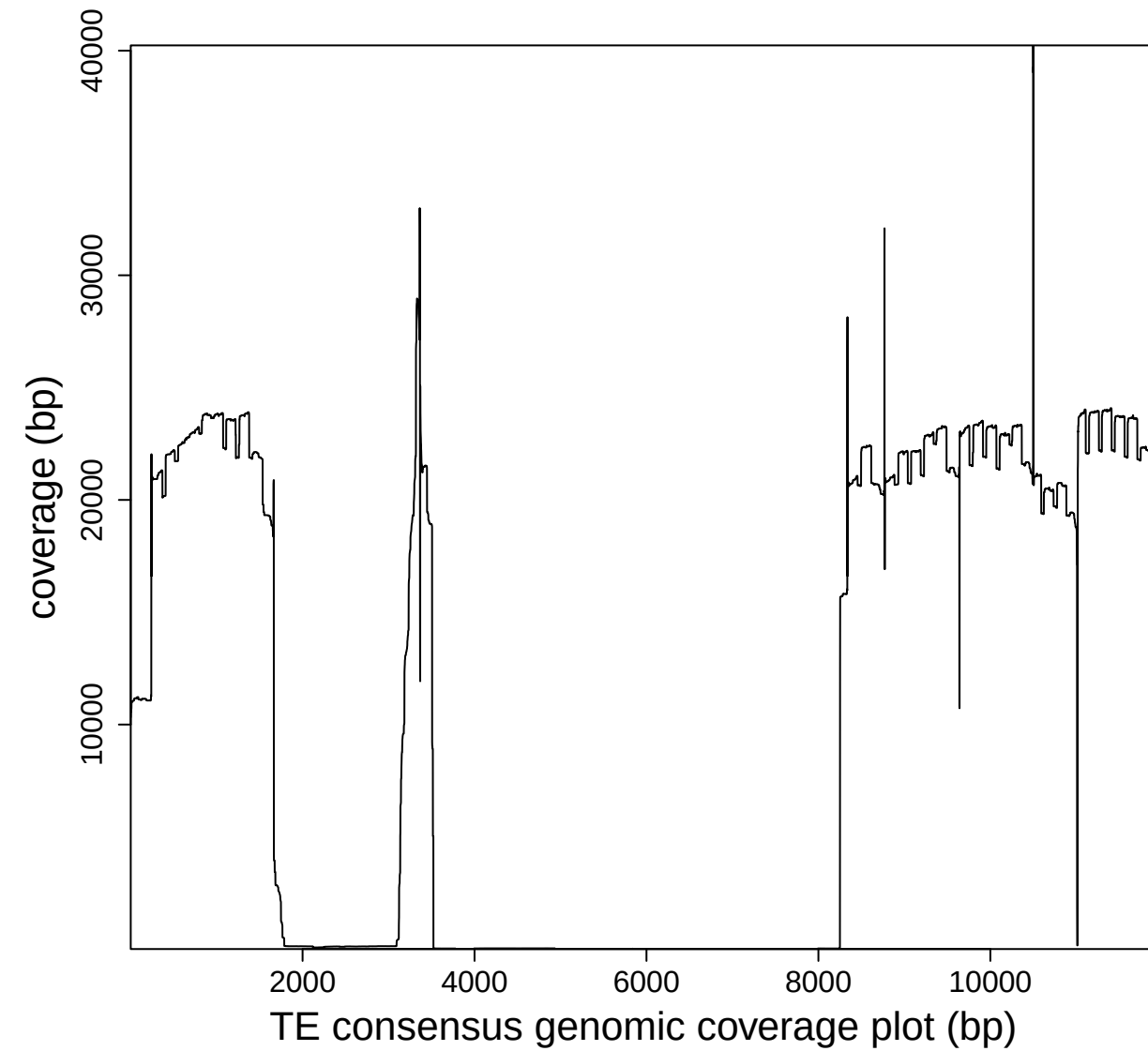
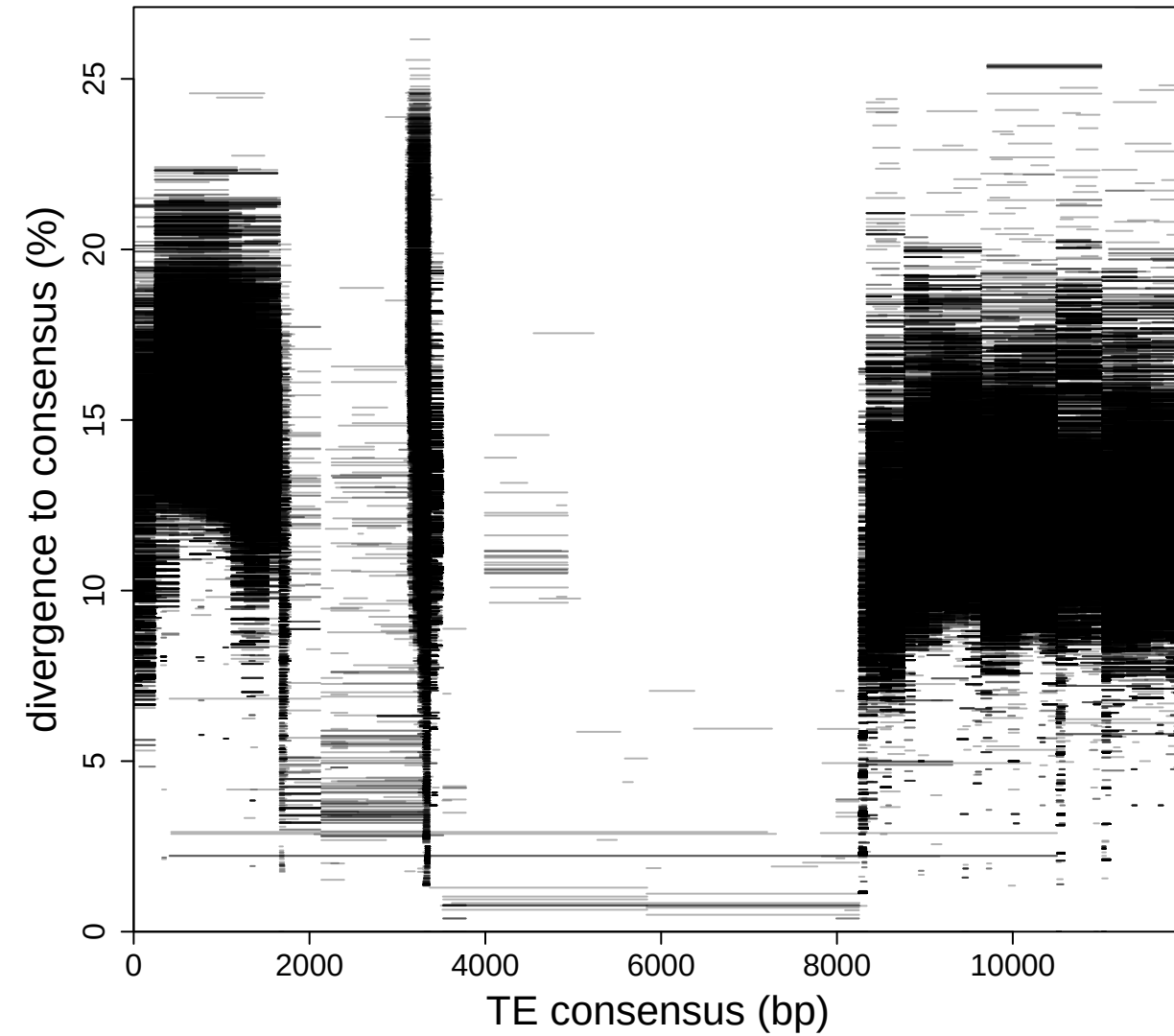
MSA length = 11928





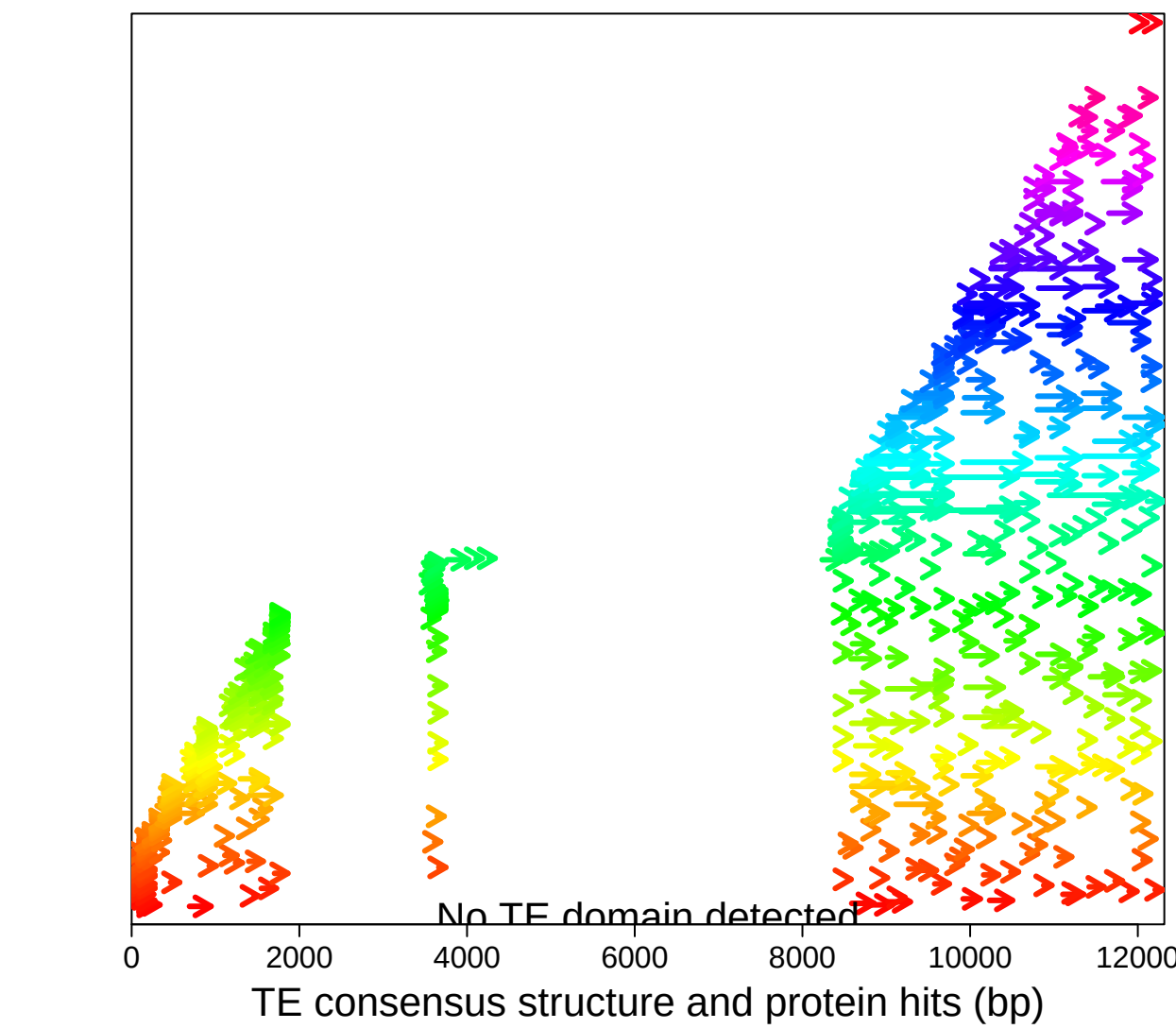
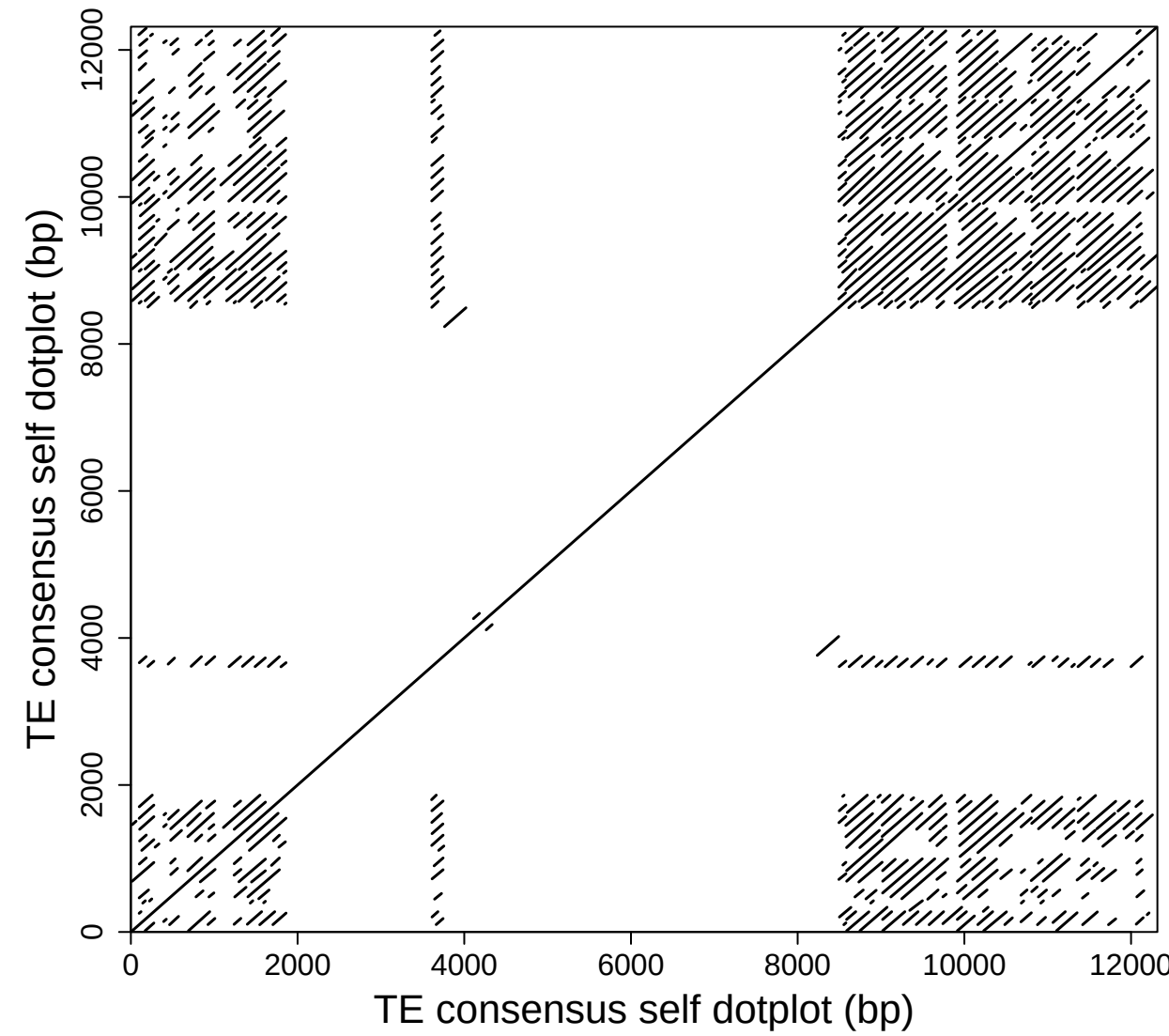
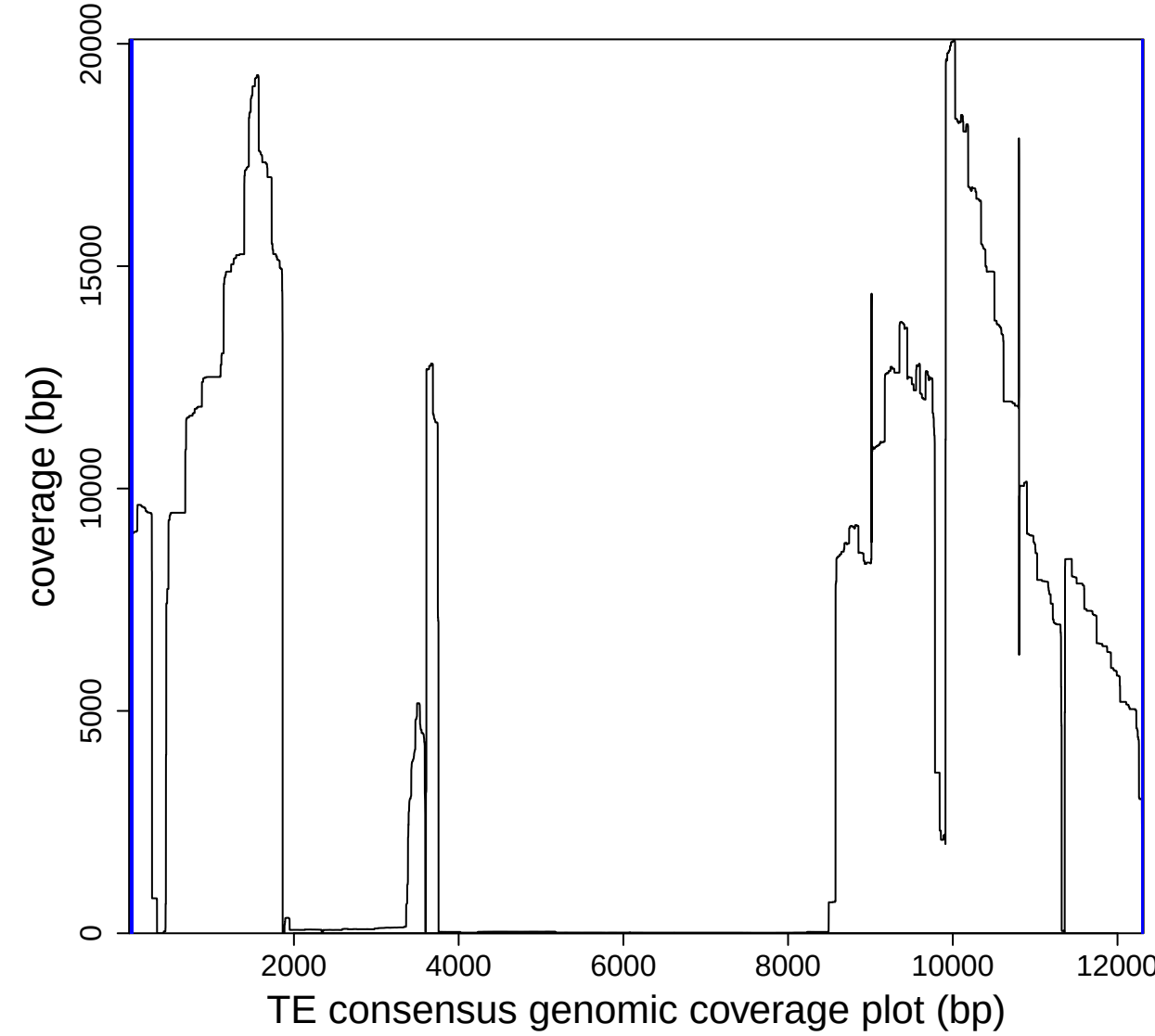
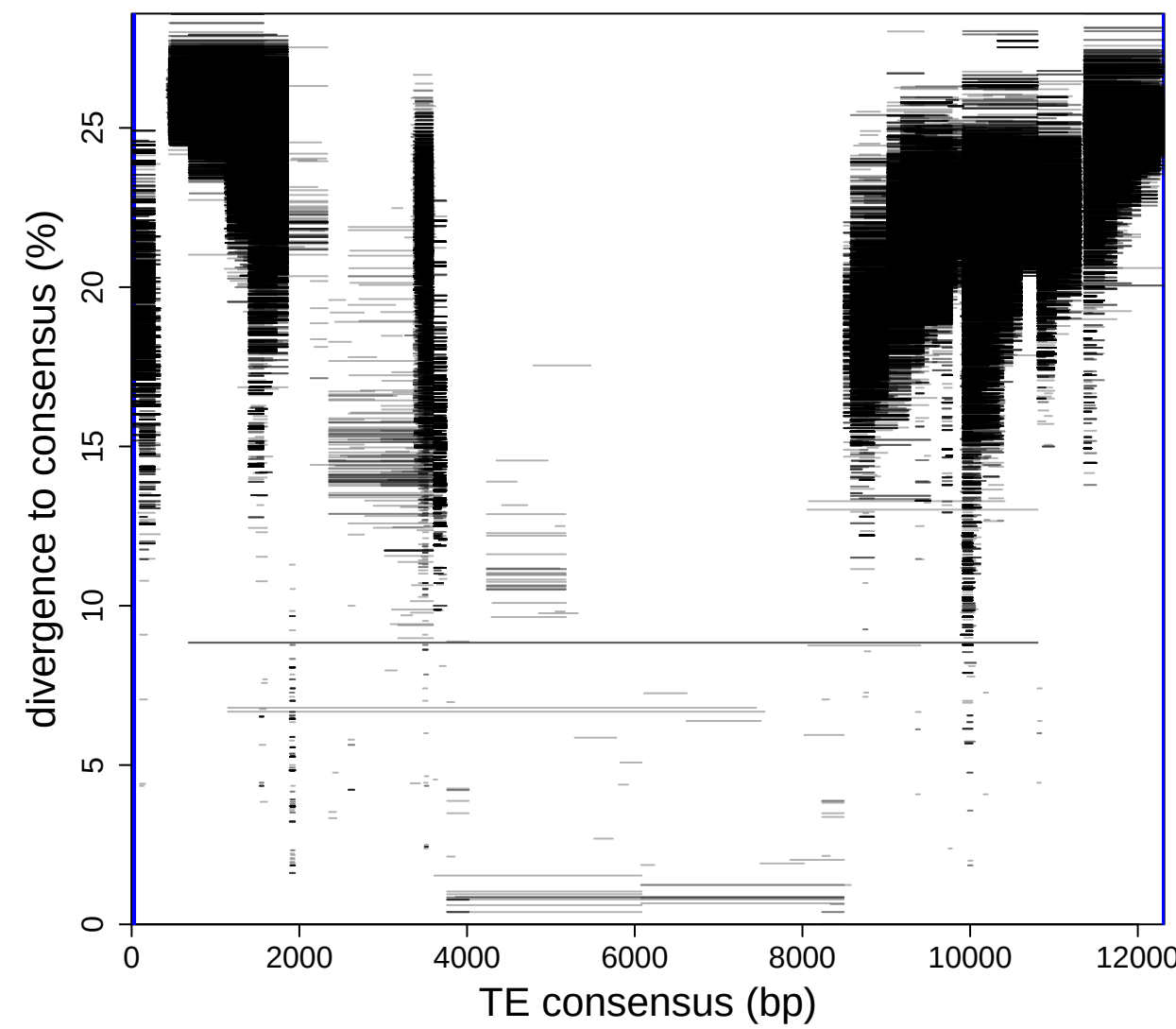
bed_fm_1.bed 0 0 n.bed g 1.bed fm 2.bed 0 0 bcln.fa aln.fa
size: 11928bp; fragments: 241099; full length: 0 (>=10735.2bp)

After TEtrimmer 11928 bp



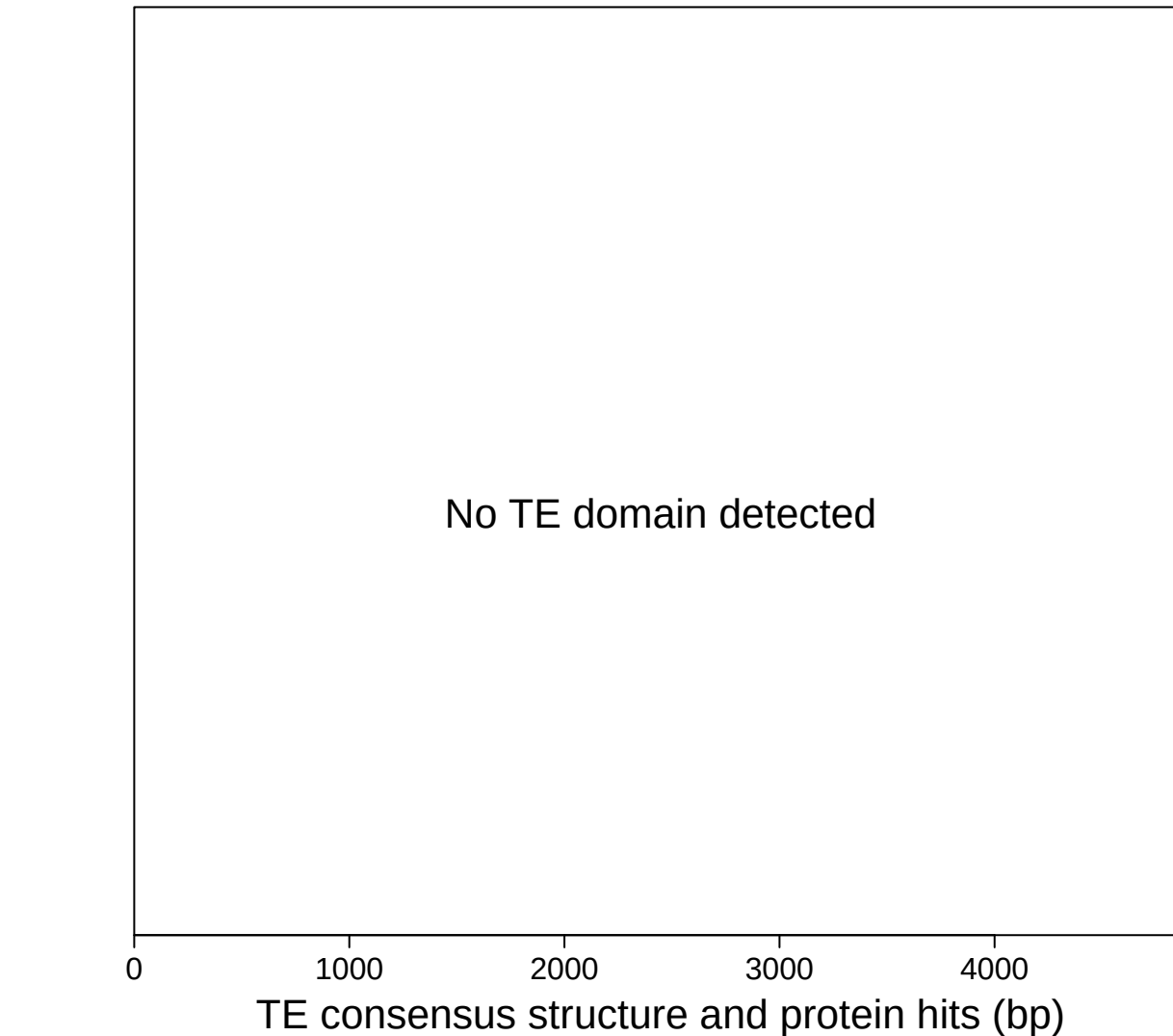
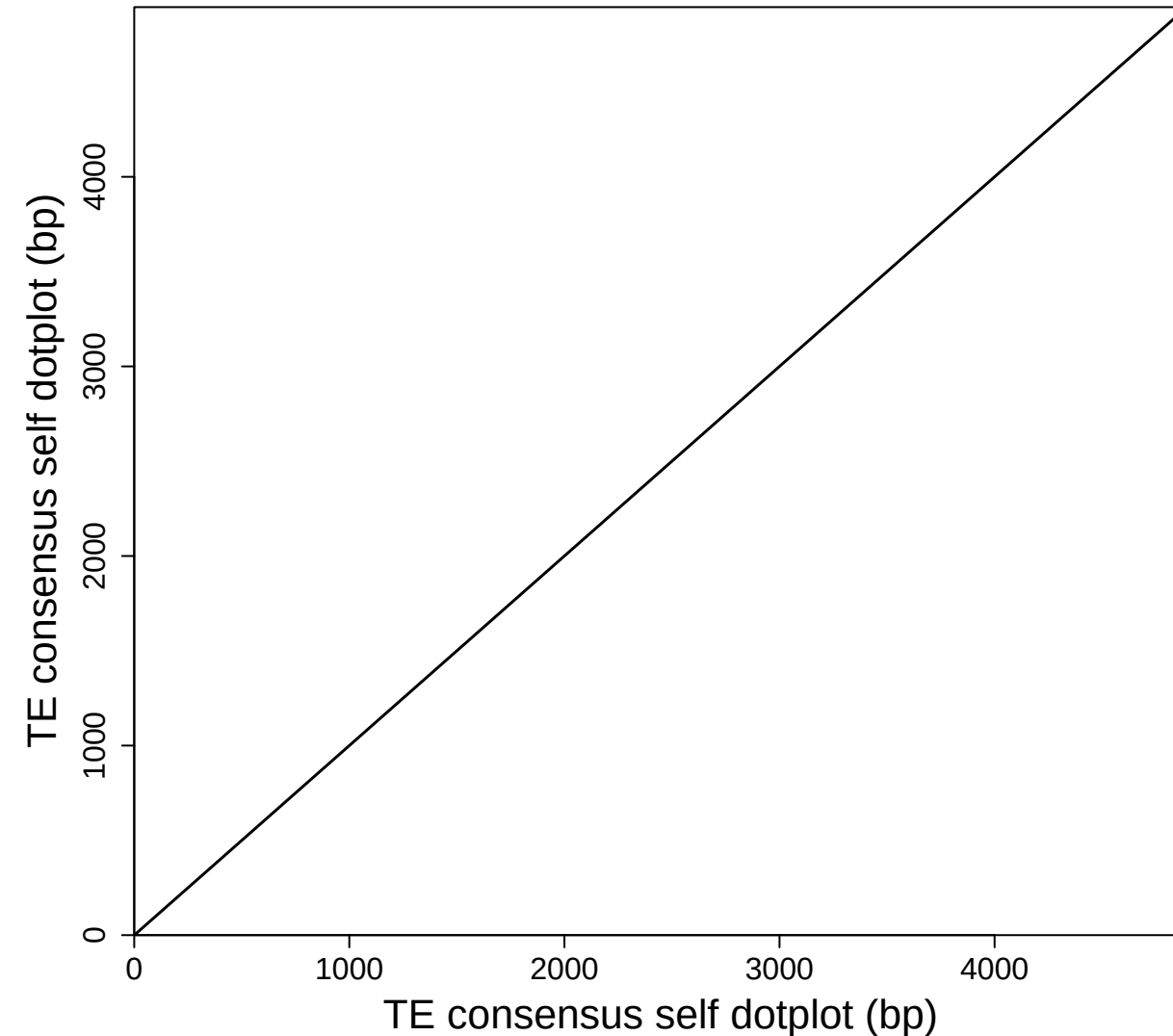
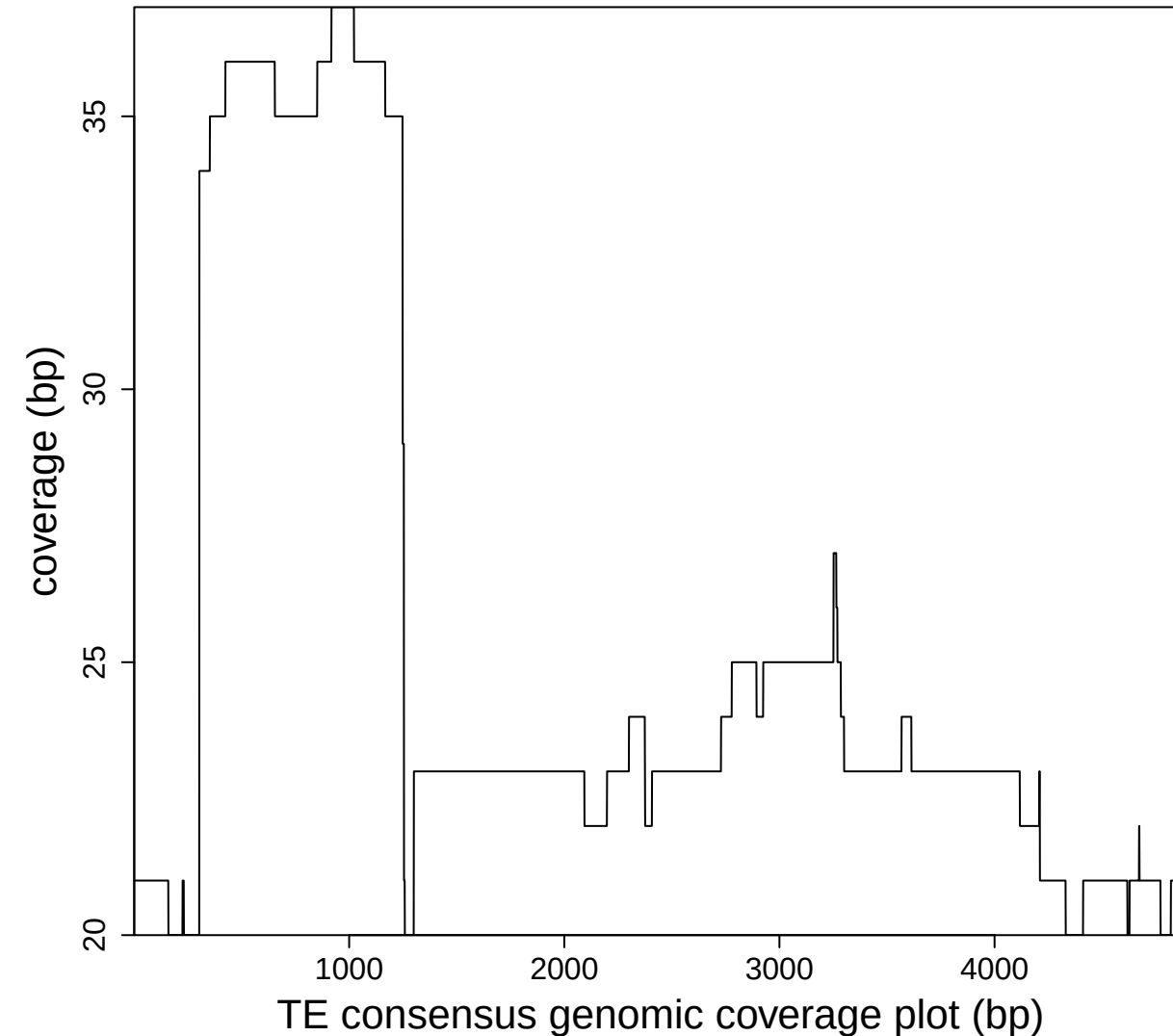
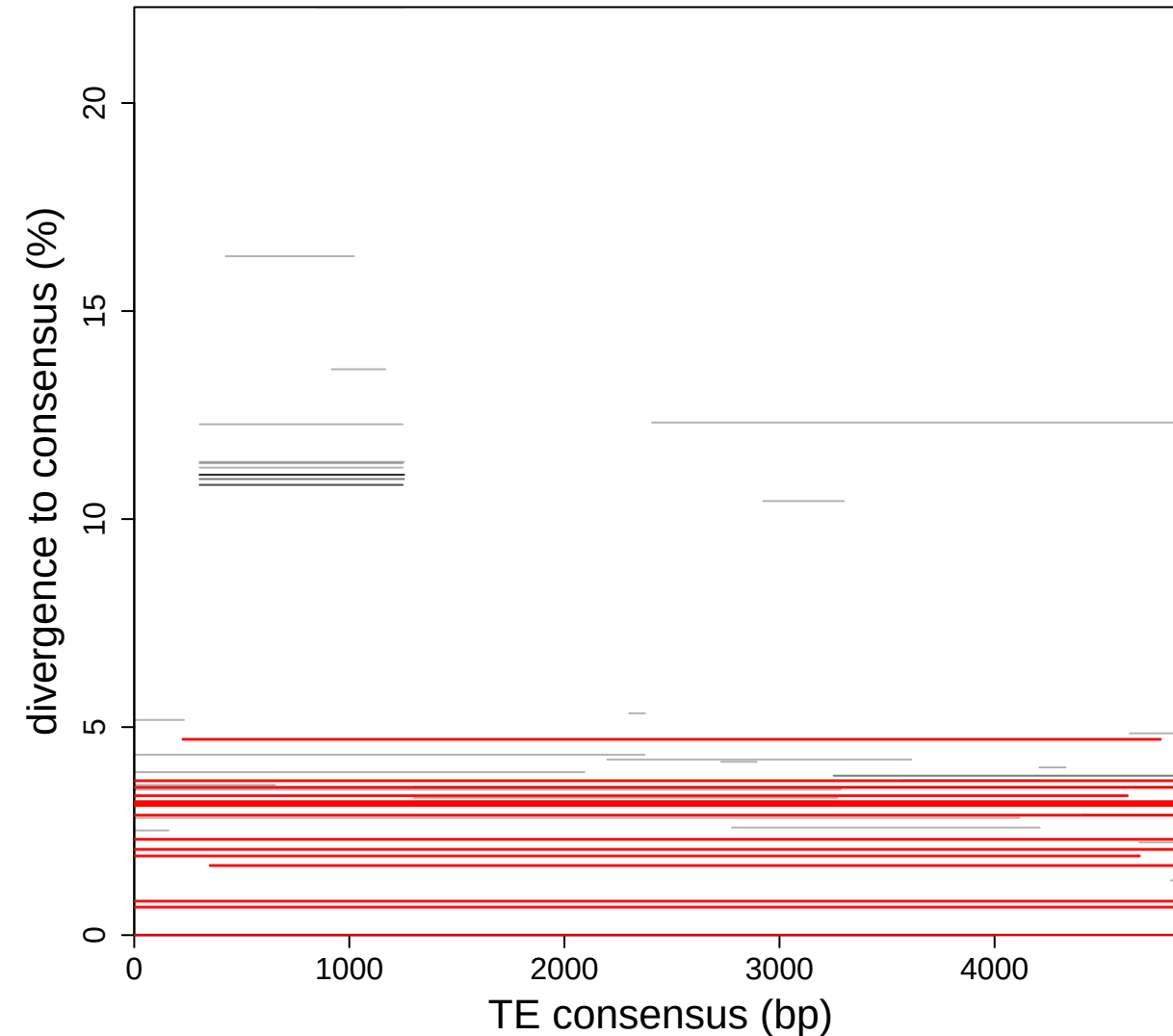
bed uf.bed g 1.bed fm 1.bed 0 0 n.bed g 1.bed fm 2.bed 0
size: 12316bp; fragments: 108156; full length: 0 (>=11084.4bp)

After TEtrimmer Extended plot Blue lines are boundaries

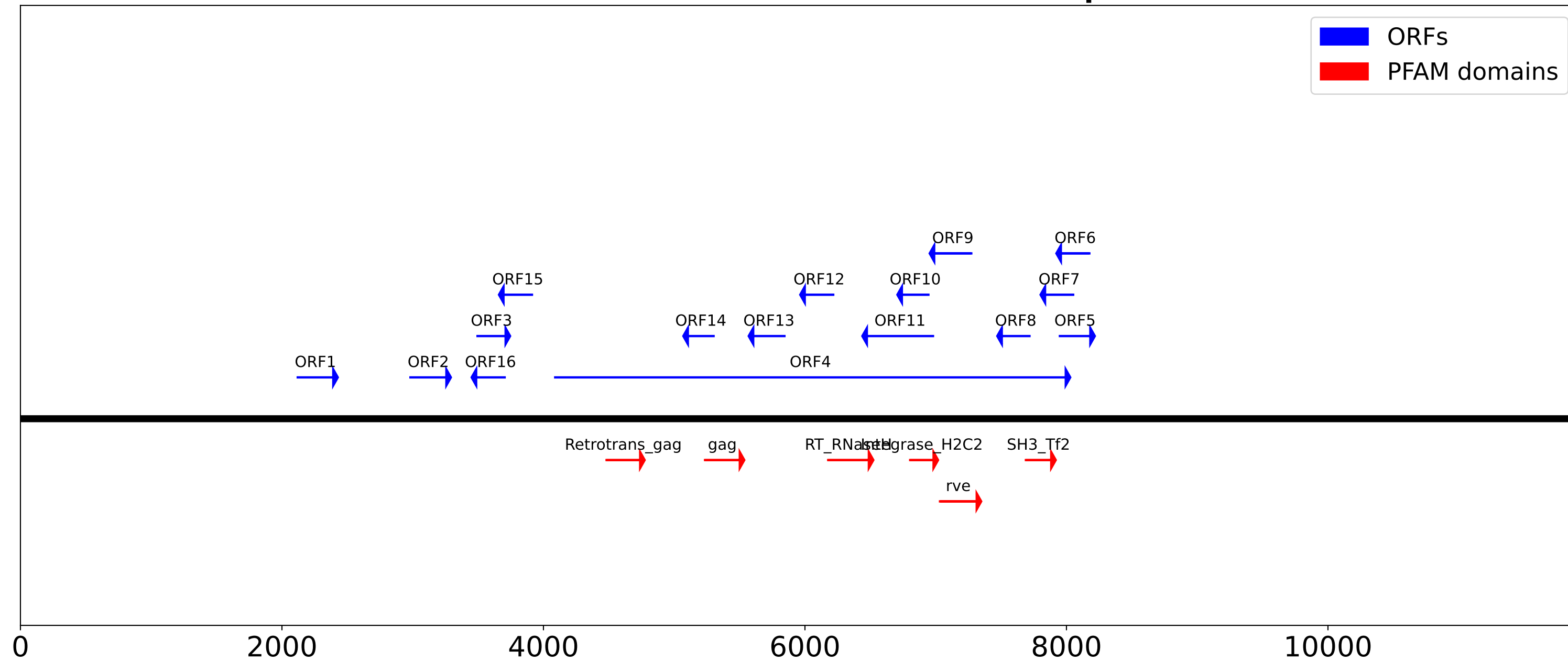


TE: ltr_1_family_1034
size: 4895bp; fragments: 57; full length: 16 (≥ 4405.5 bp)

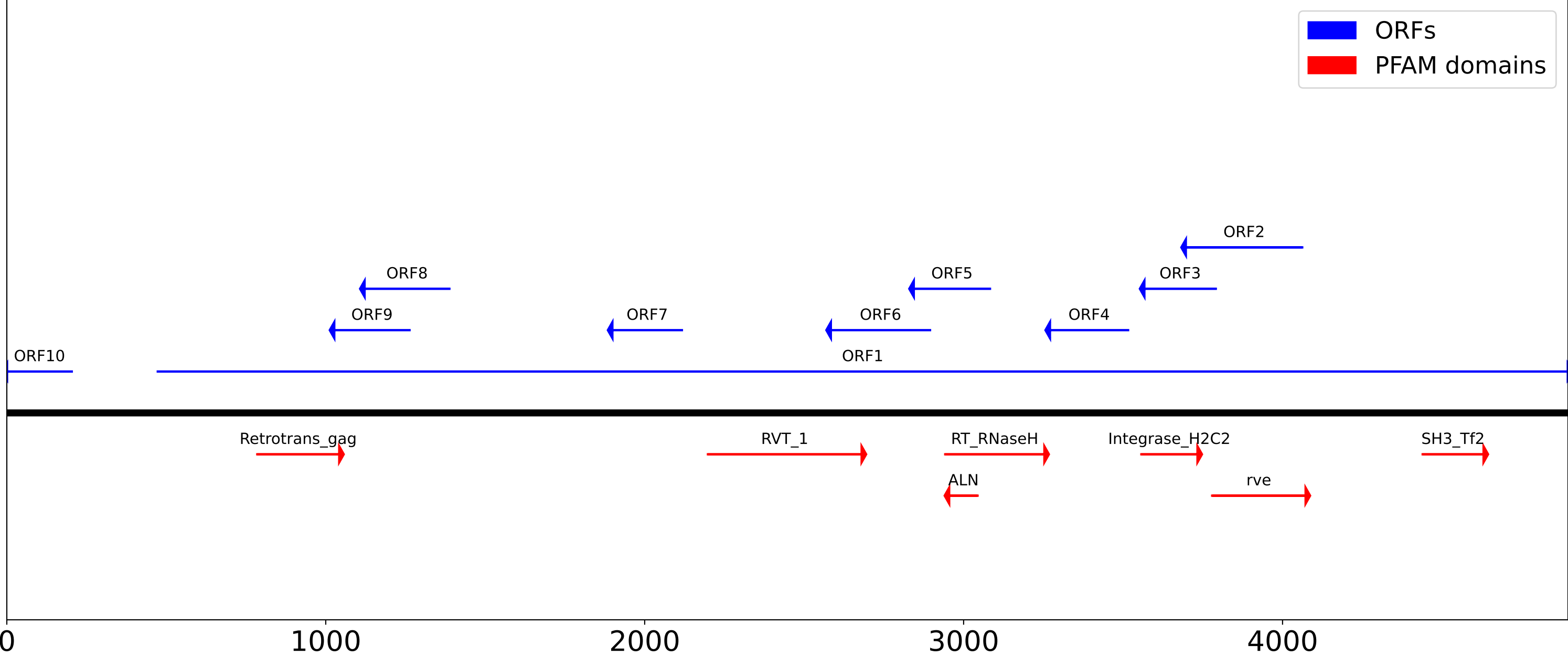
Before TEtrimmer 4895 bp



After TEtrimmer ORF and PFAM domain plot



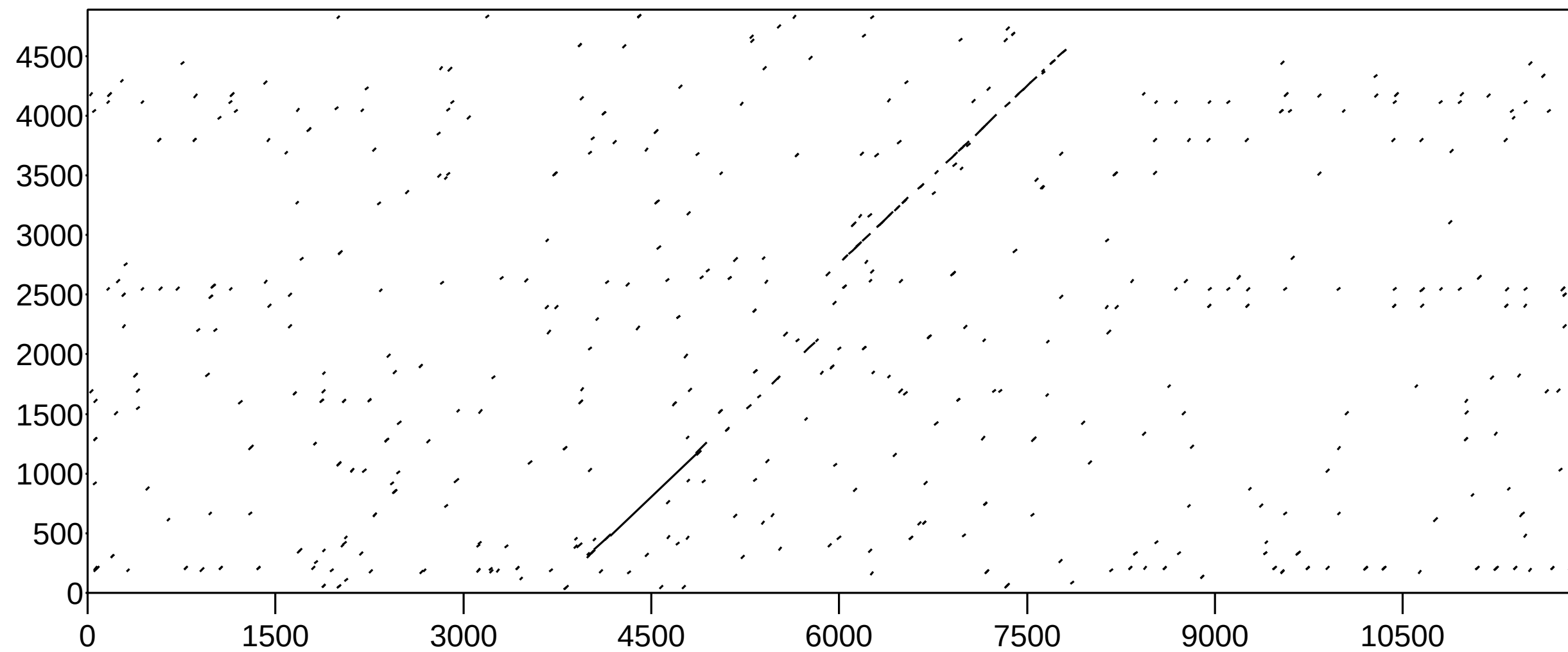
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)