

Start crop Point

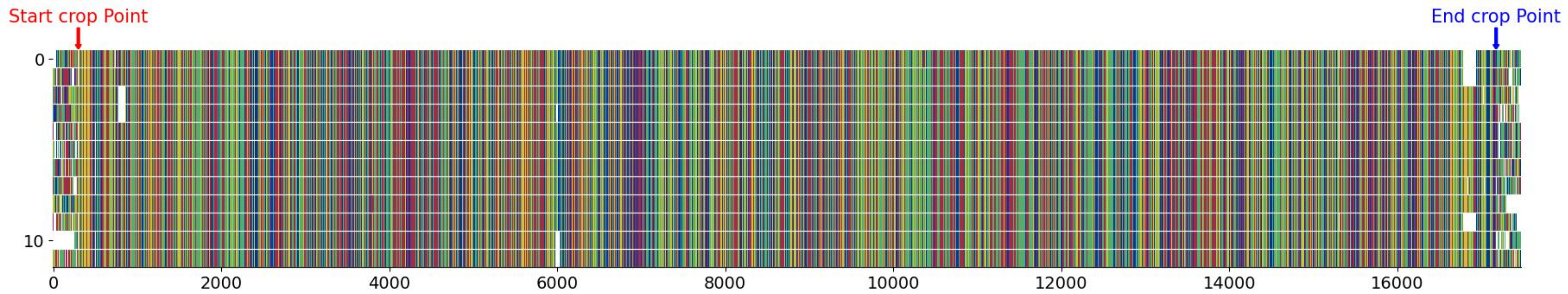


End crop Point



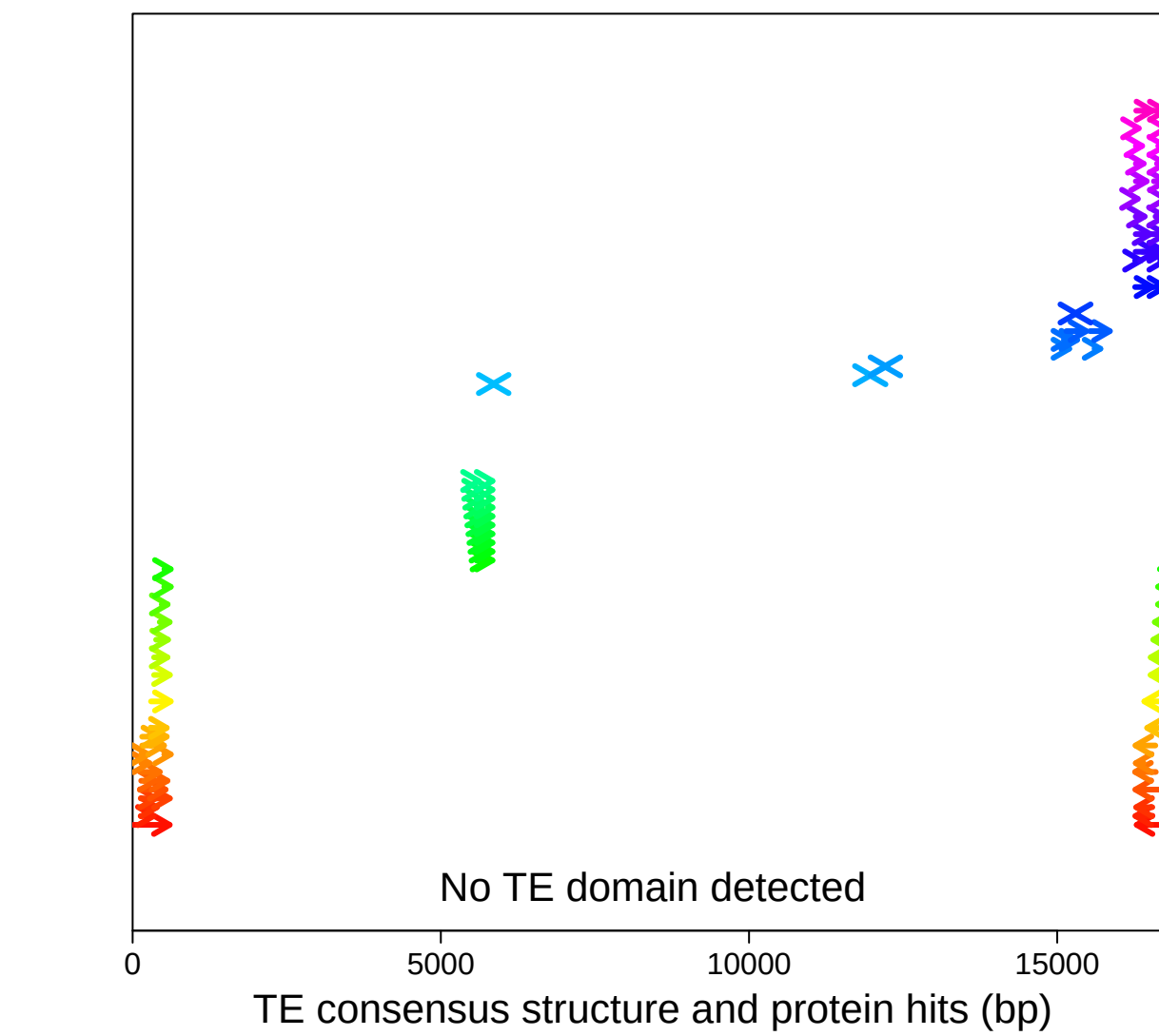
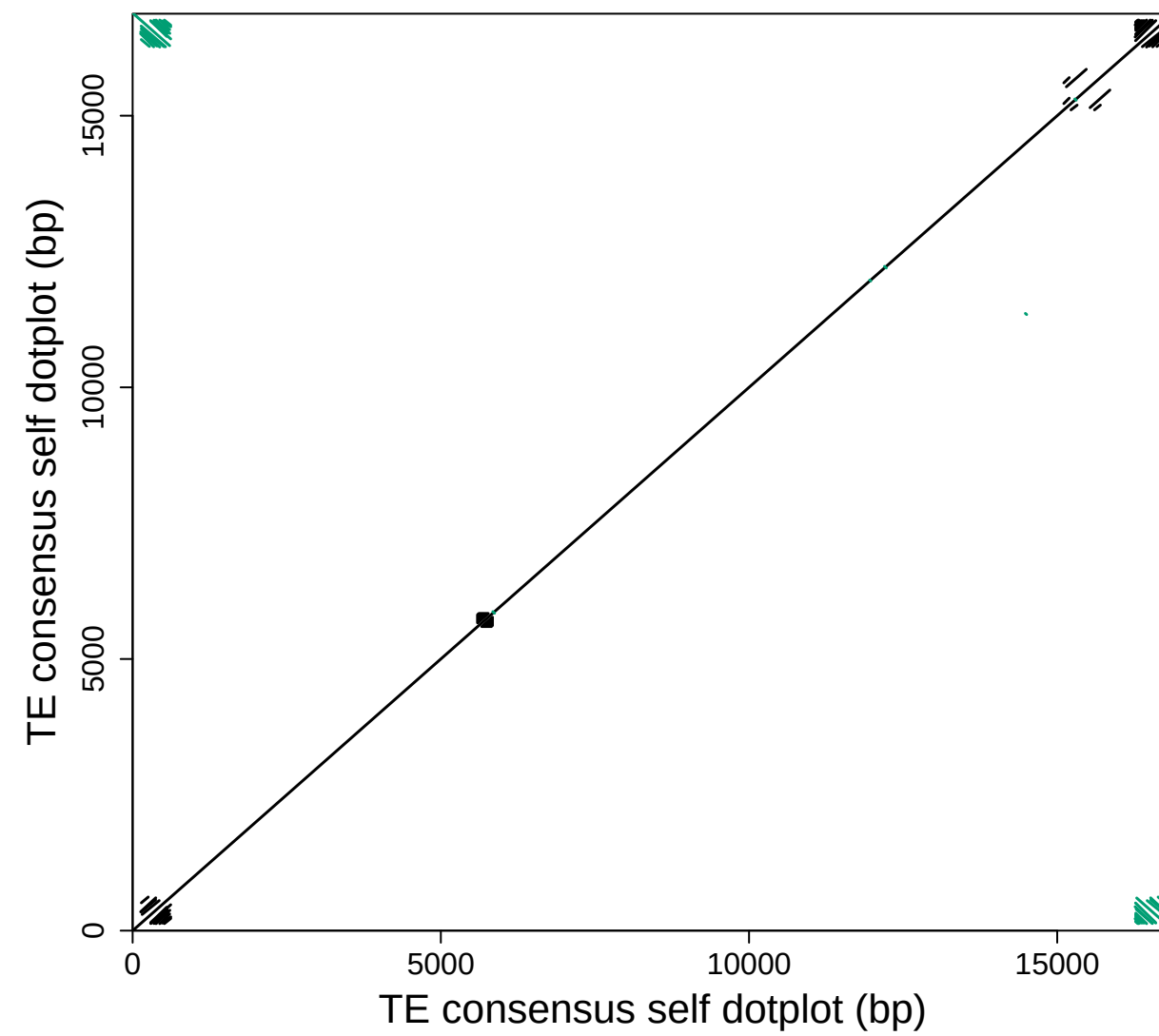
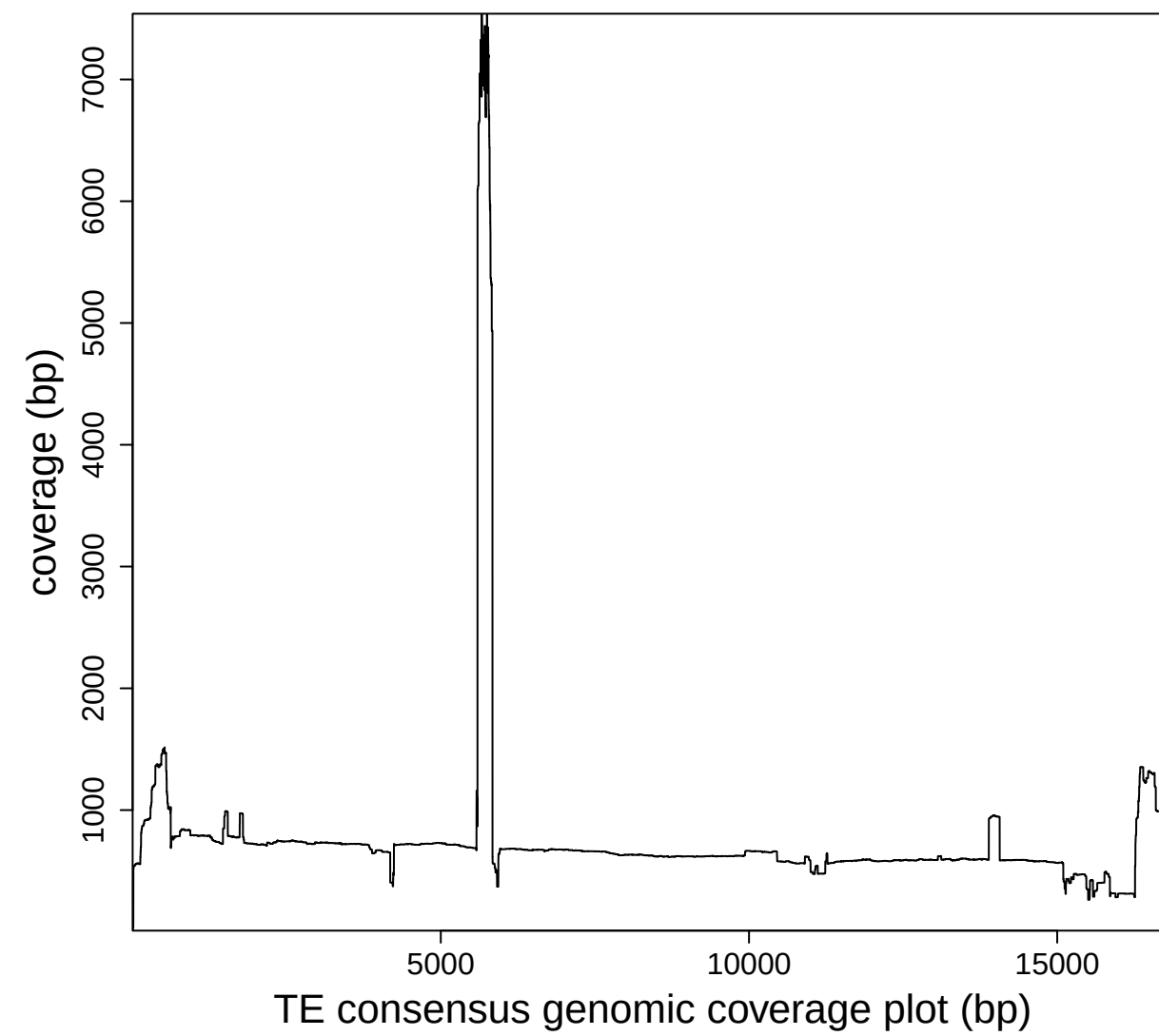
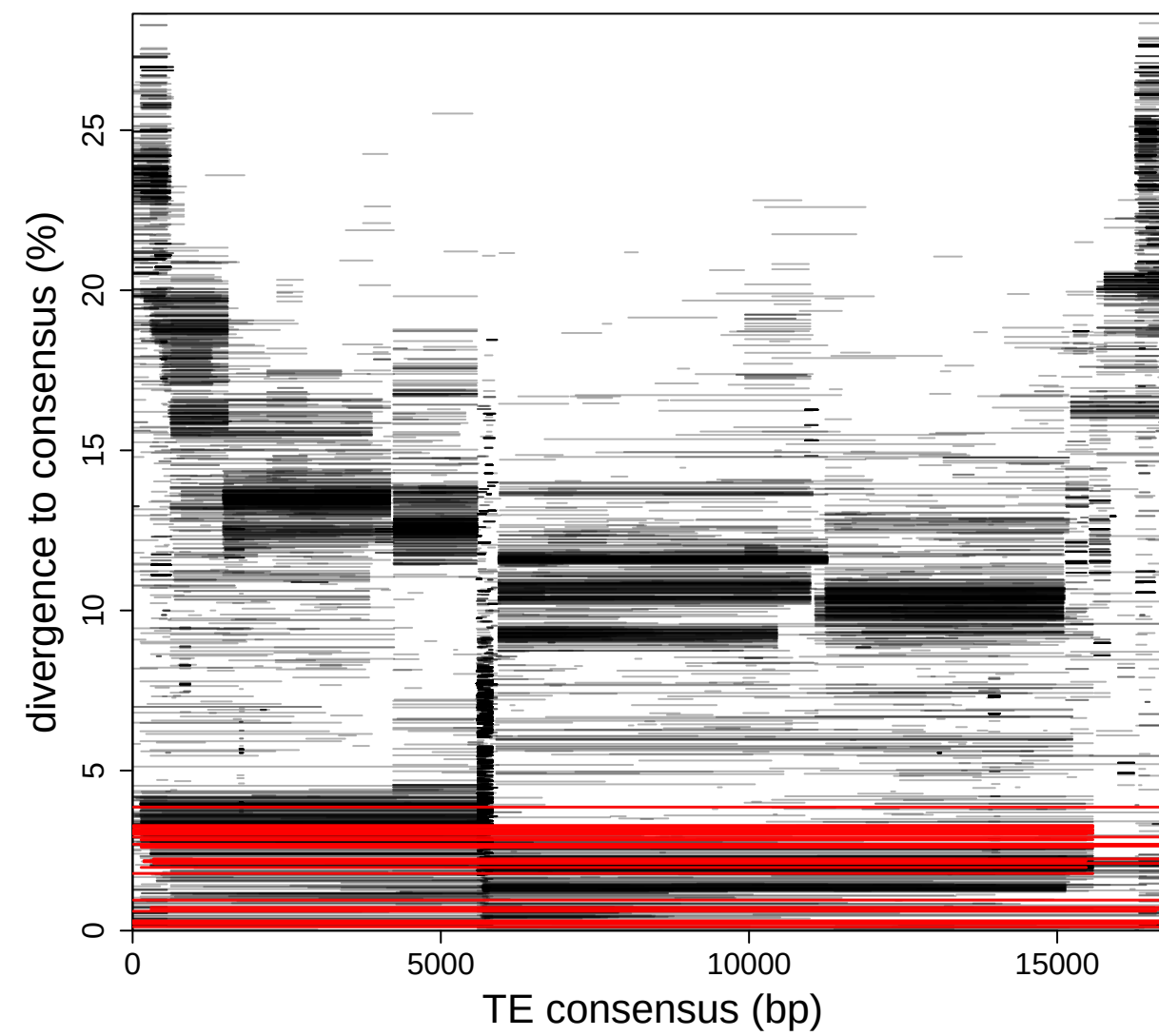
MSA length = 16877





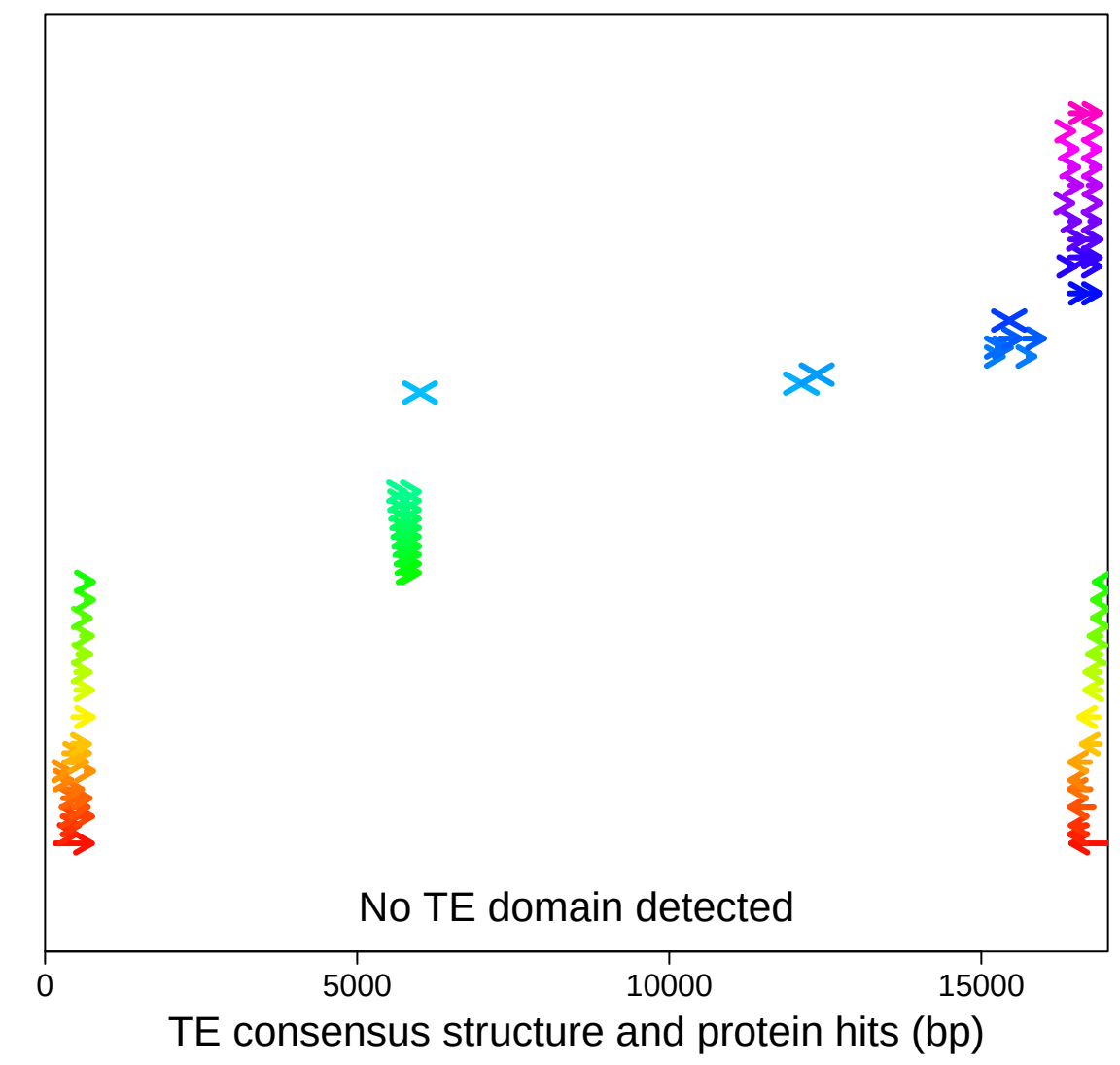
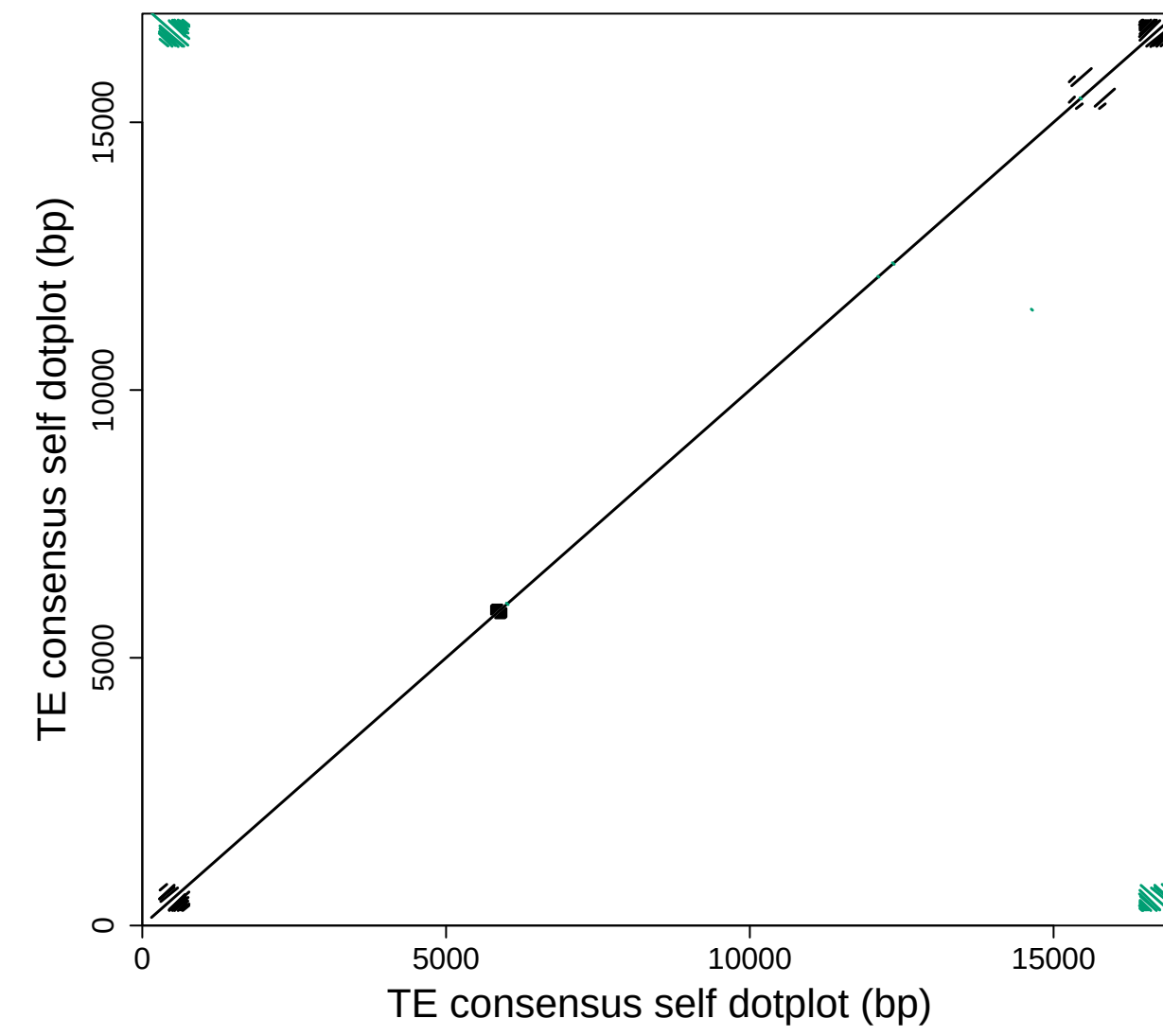
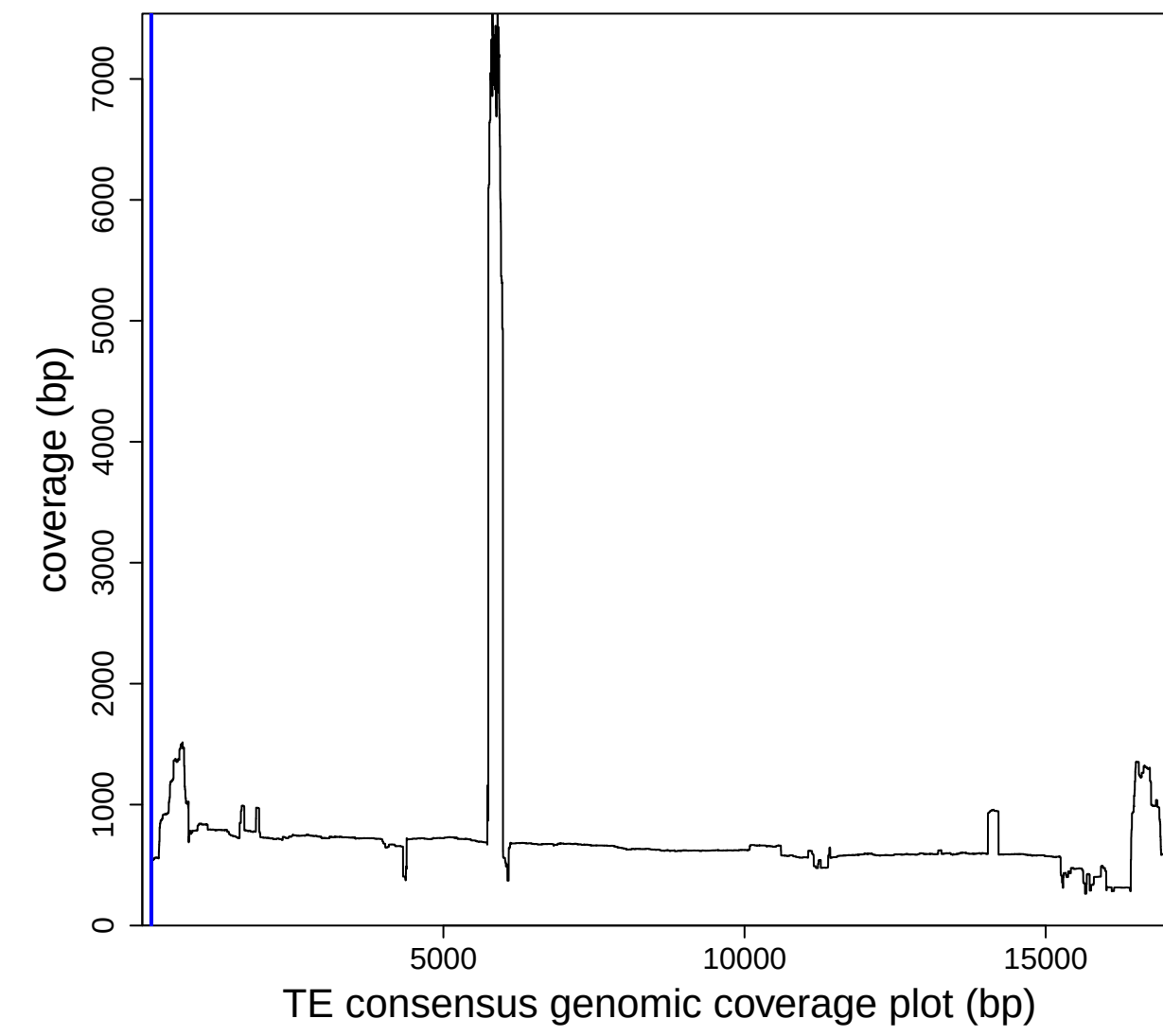
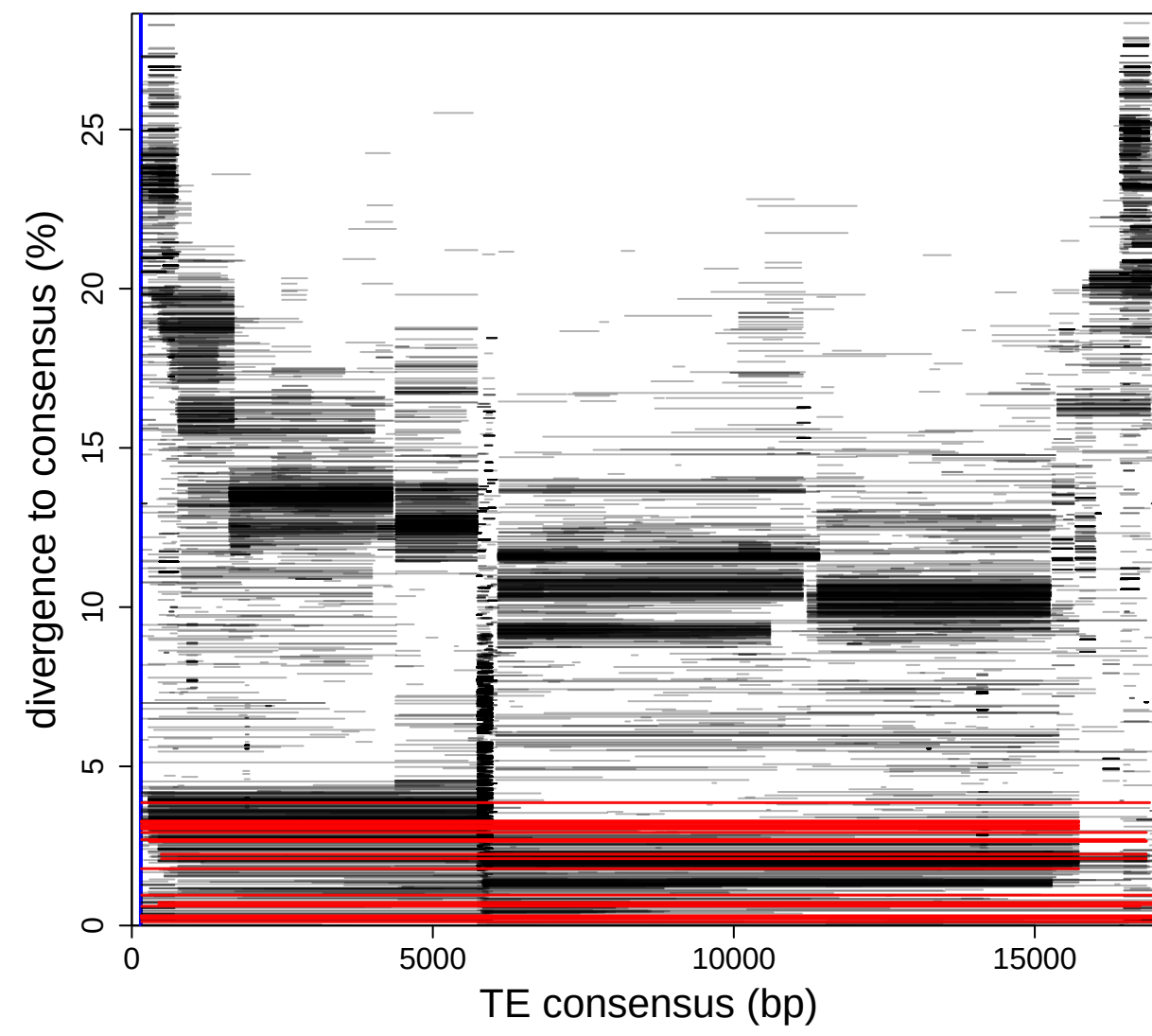
l.b.bed uf.bed g_3.bed fm_1.bed 0 0 bcln.fa aln.fa cl.fa gs.fa
size: 16877bp; fragments: 17669; full length: 43 (>=15189.3bp)

After TEtrimmer 16877 bp



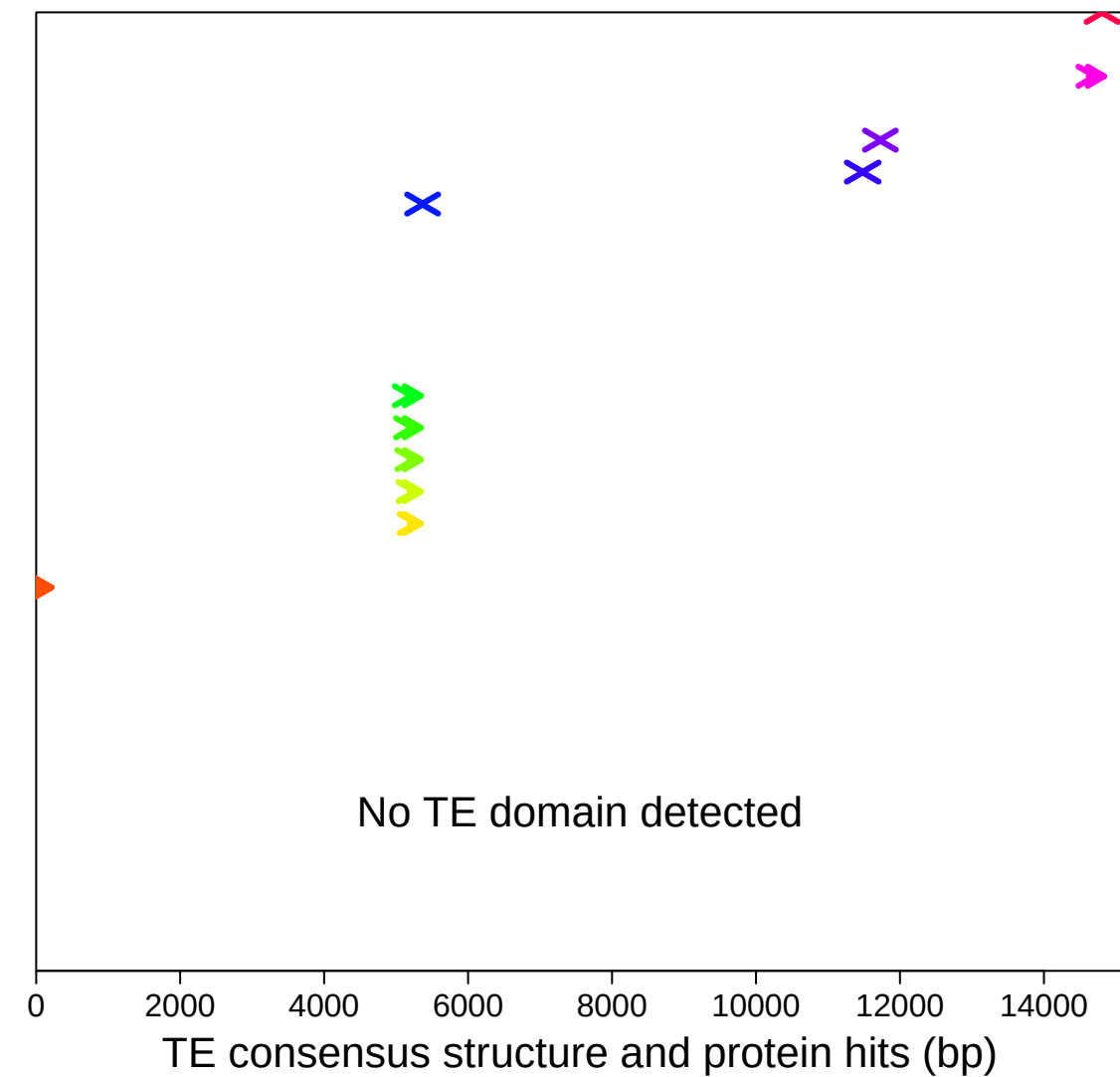
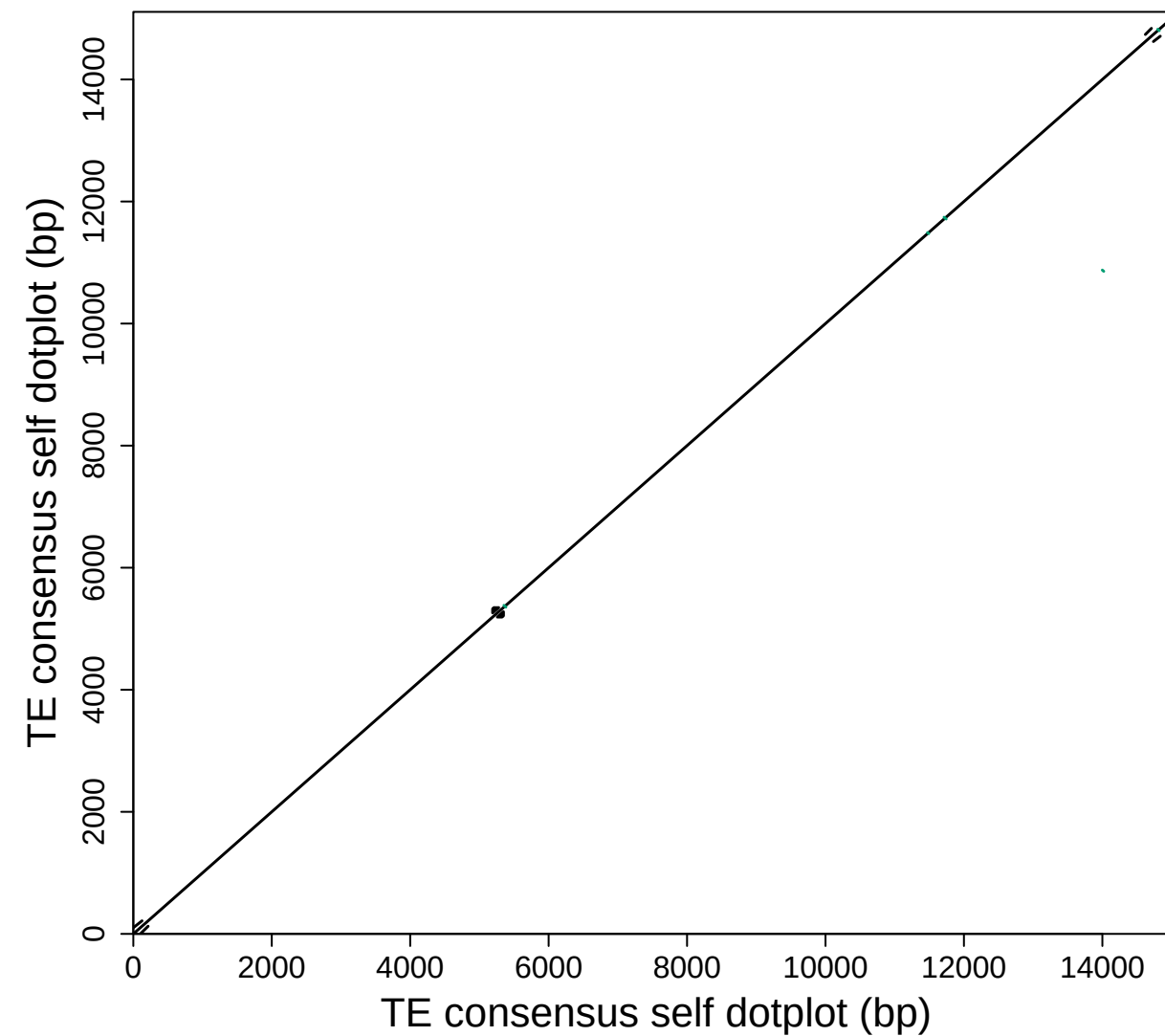
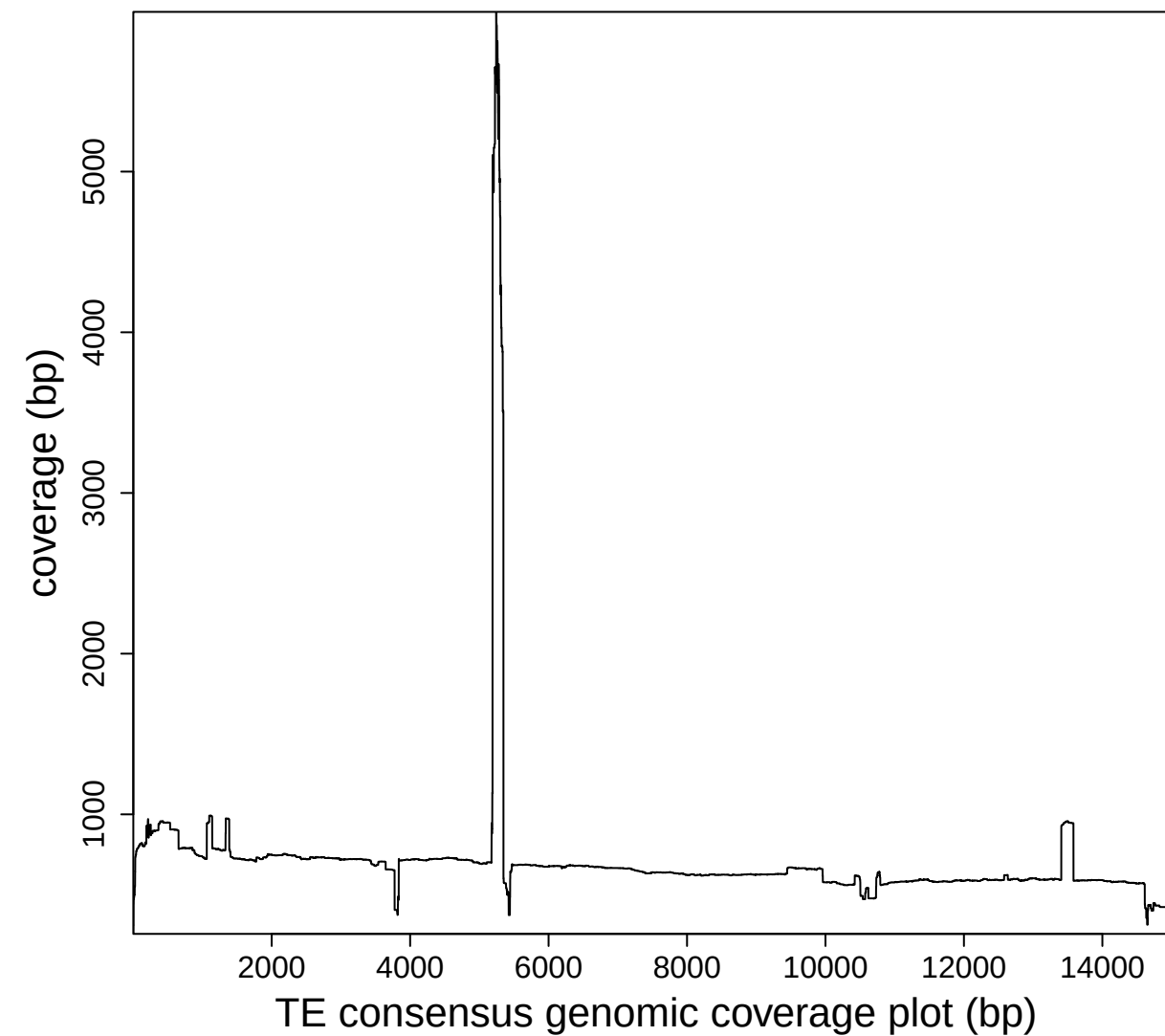
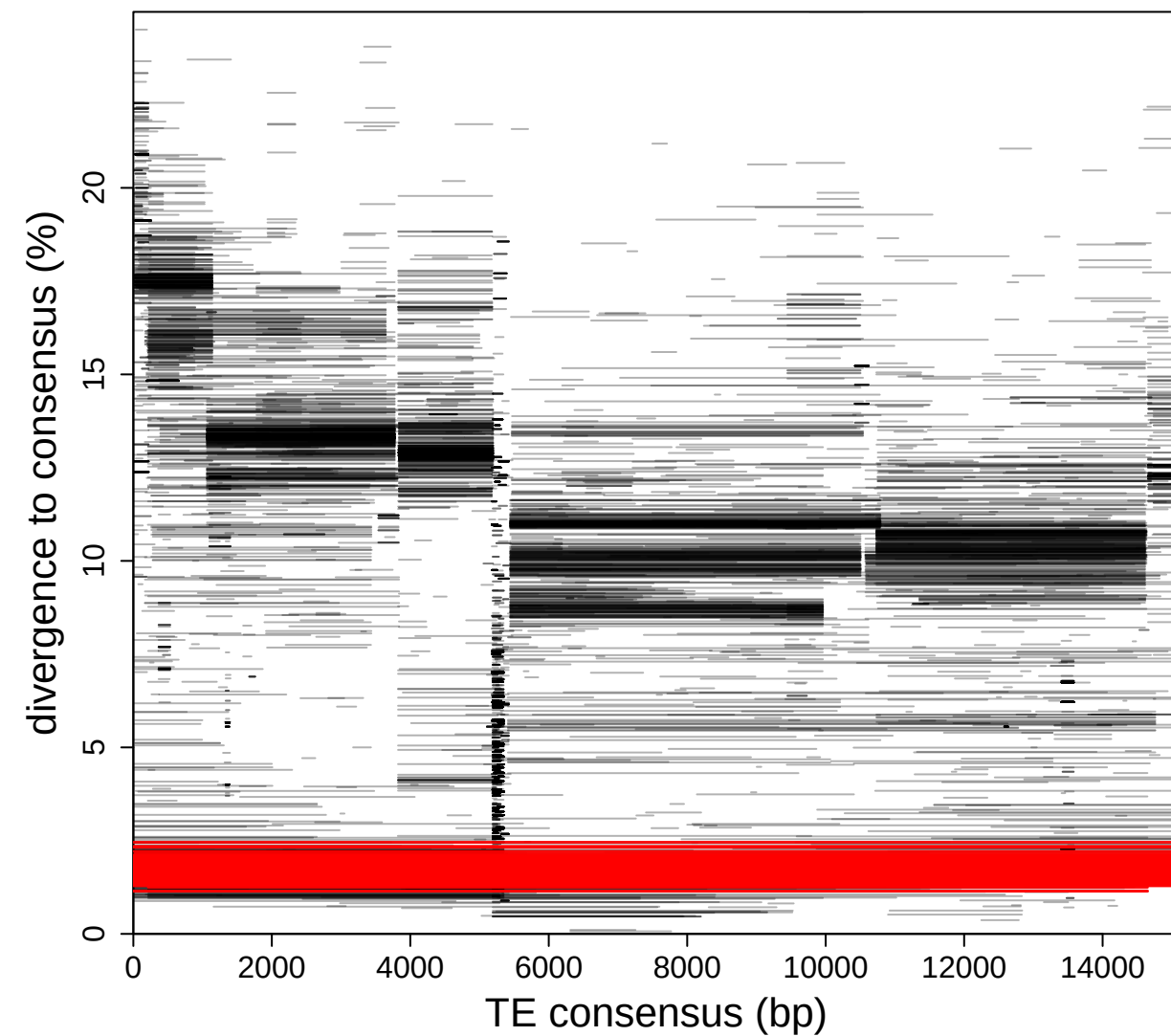
5.fasta.b.bed uf.bed g_3.bed fm_1.bed_0_0 bcln.fa aln.fa cl.fa
size: 17177bp; fragments: 17669; full length: 29 (>=15459.3bp)

After TEtrimmer Extended plot Blue lines are boundaries

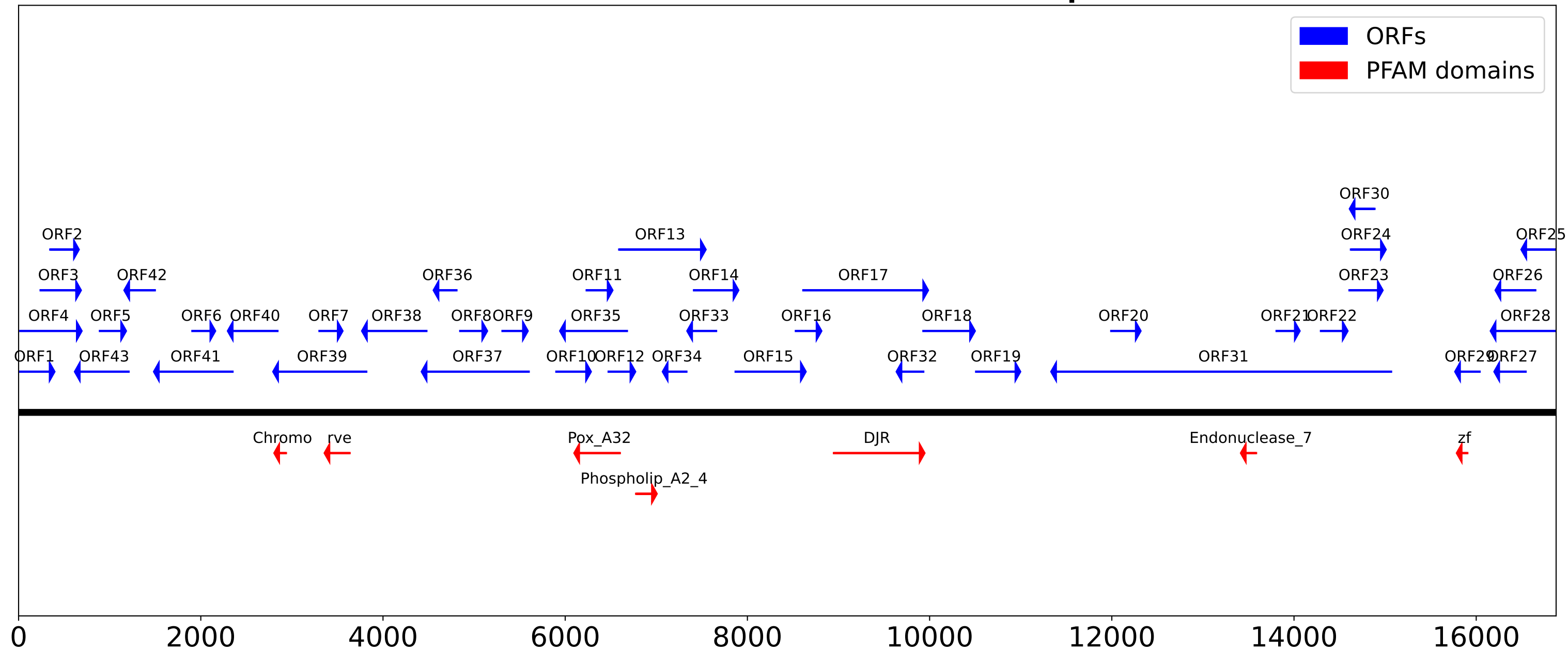


TE: rnd_4_family_735
size: 15107bp; fragments: 11047; full length: 128 (≥ 13596.3 b

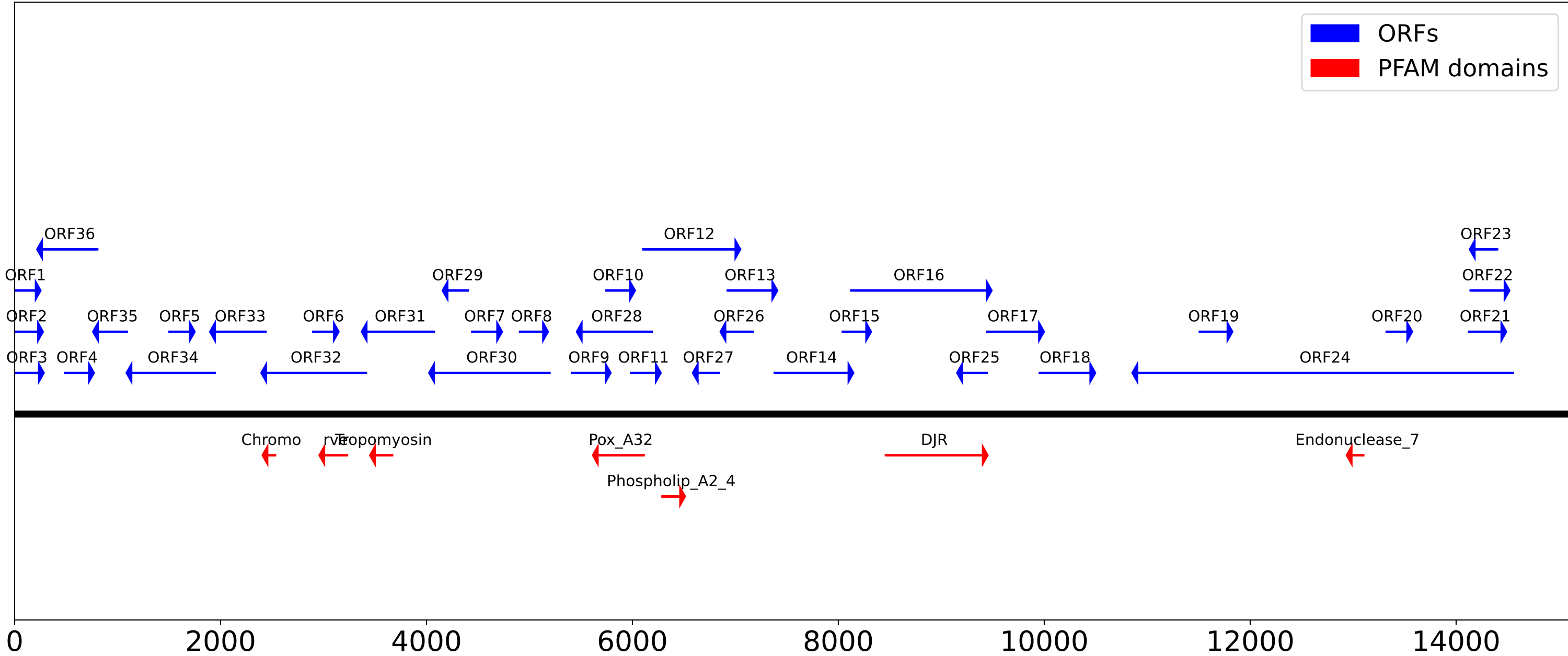
Before TEtrimmer 15107 bp



After TETrimmer ORF and PFAM domain plot



Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

