

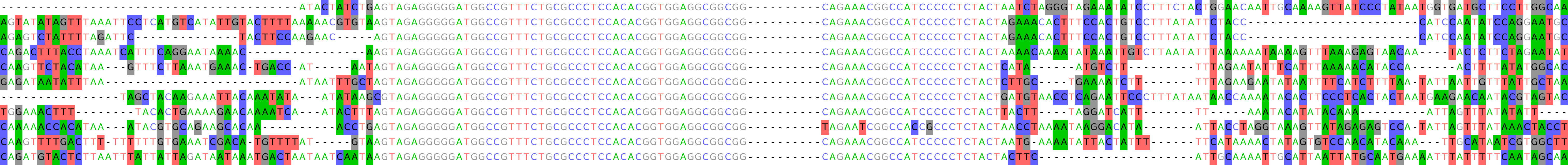
Start crop Point

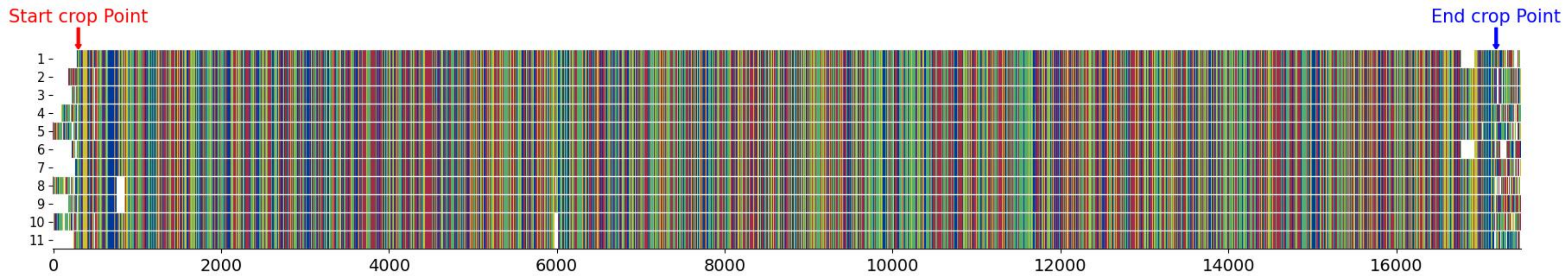


End crop Point



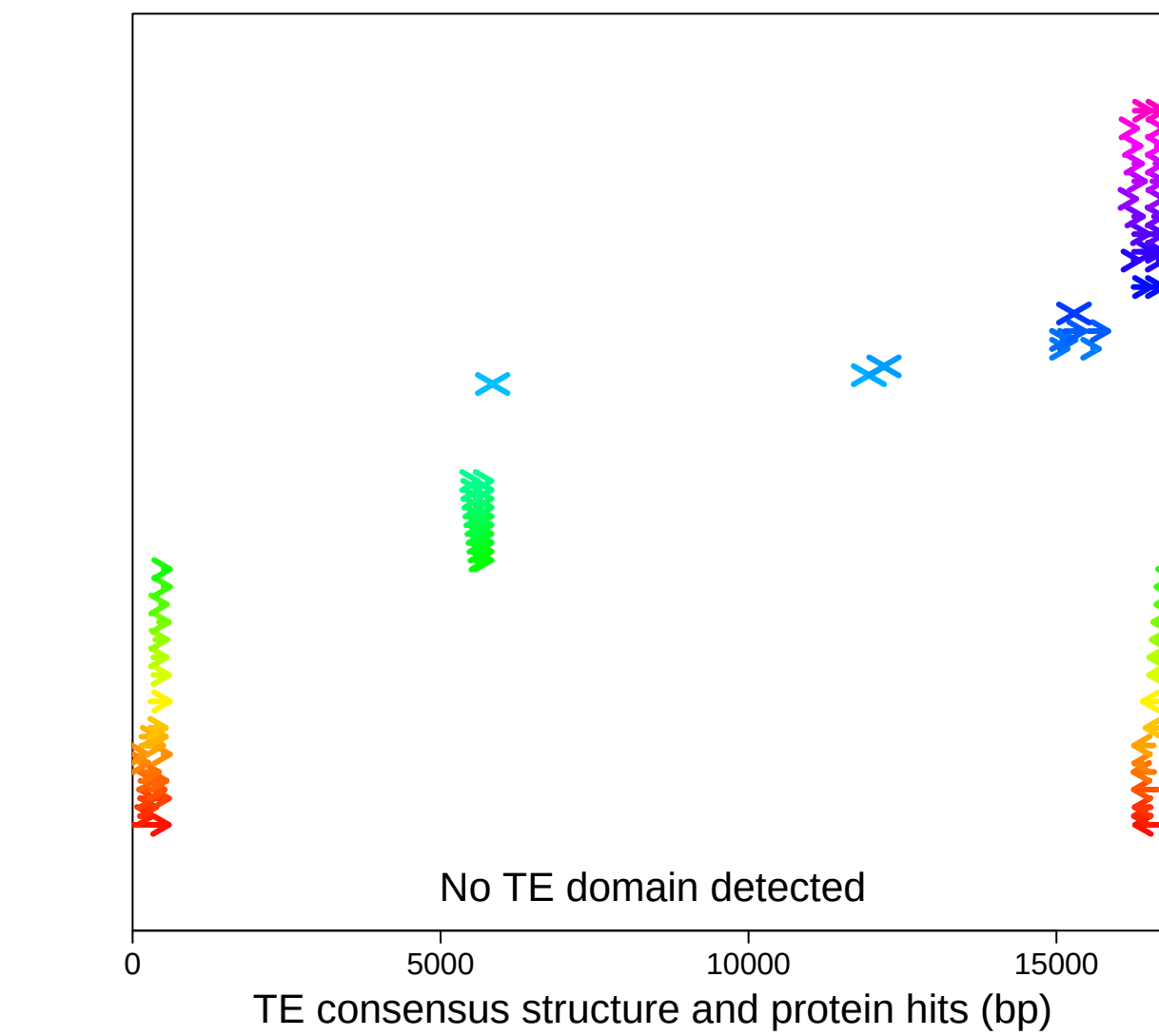
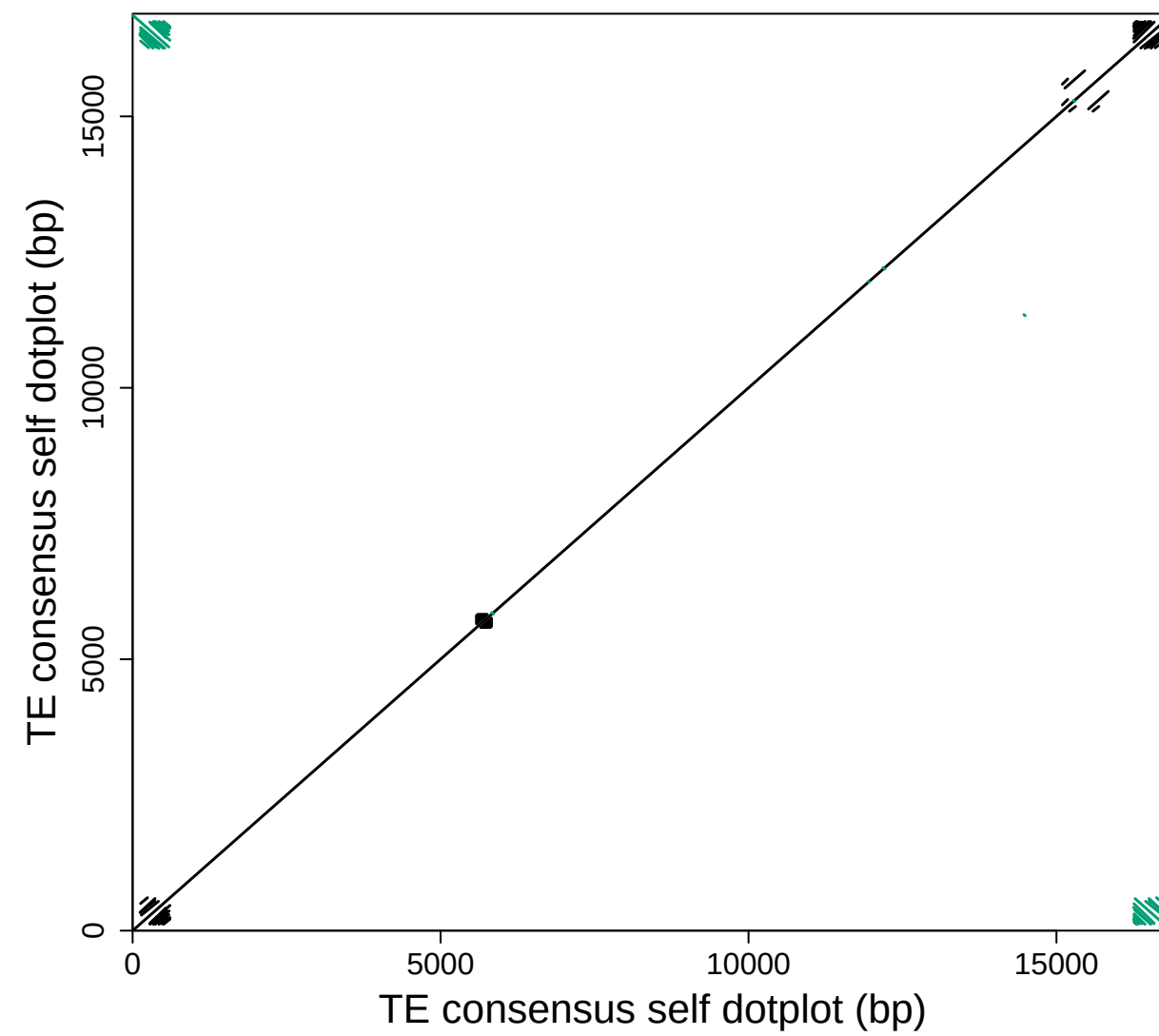
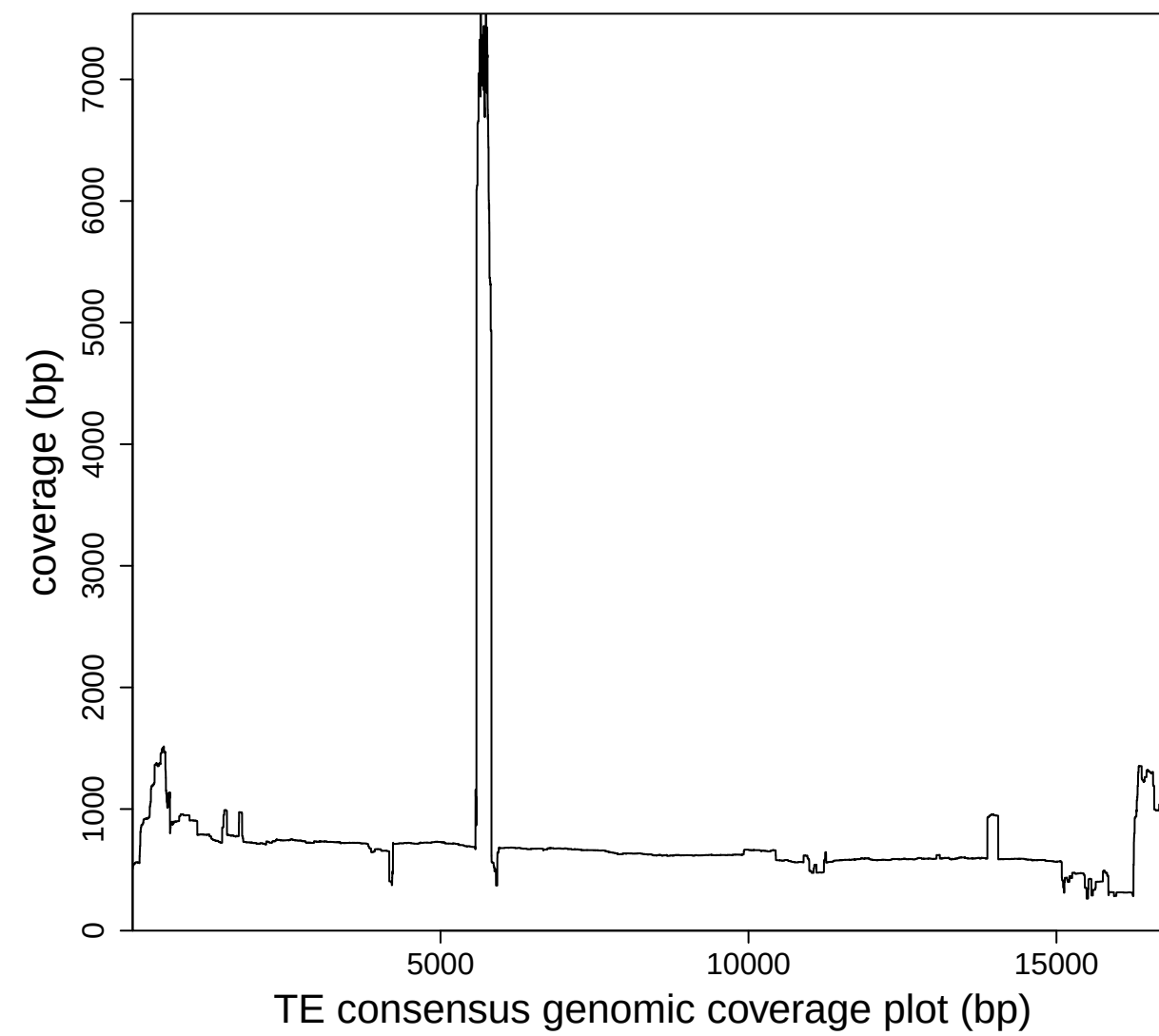
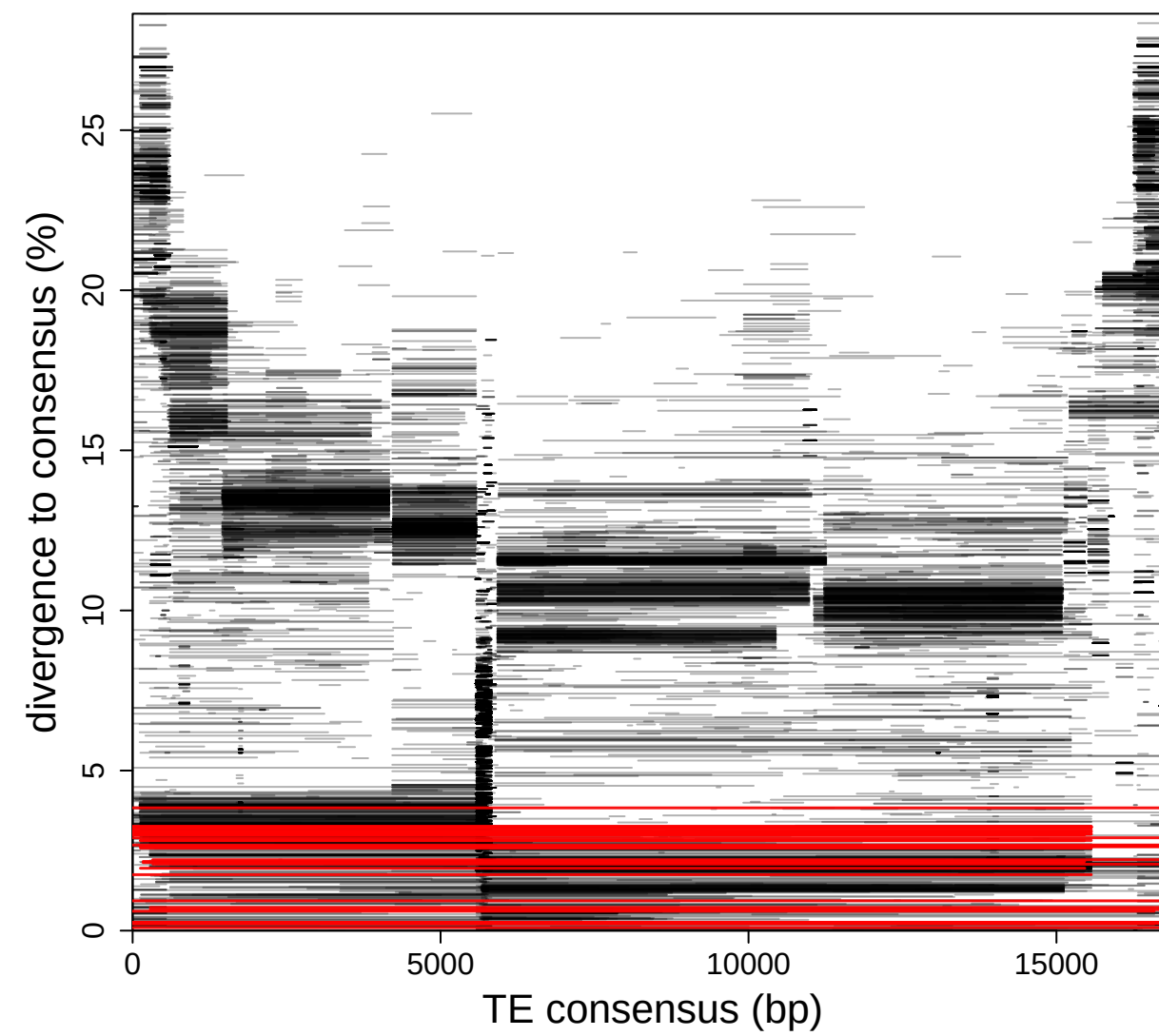
MSA length = 16891





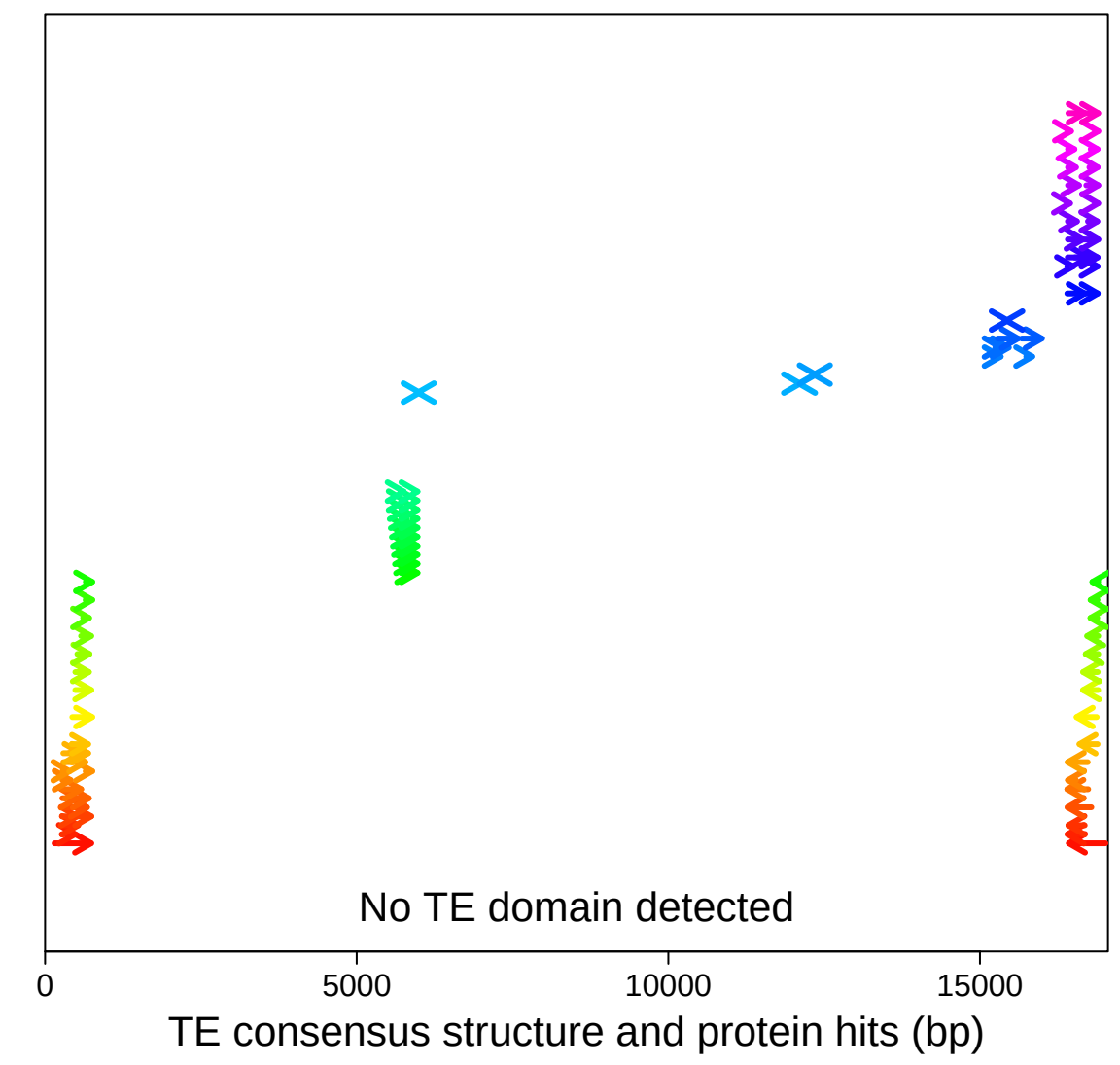
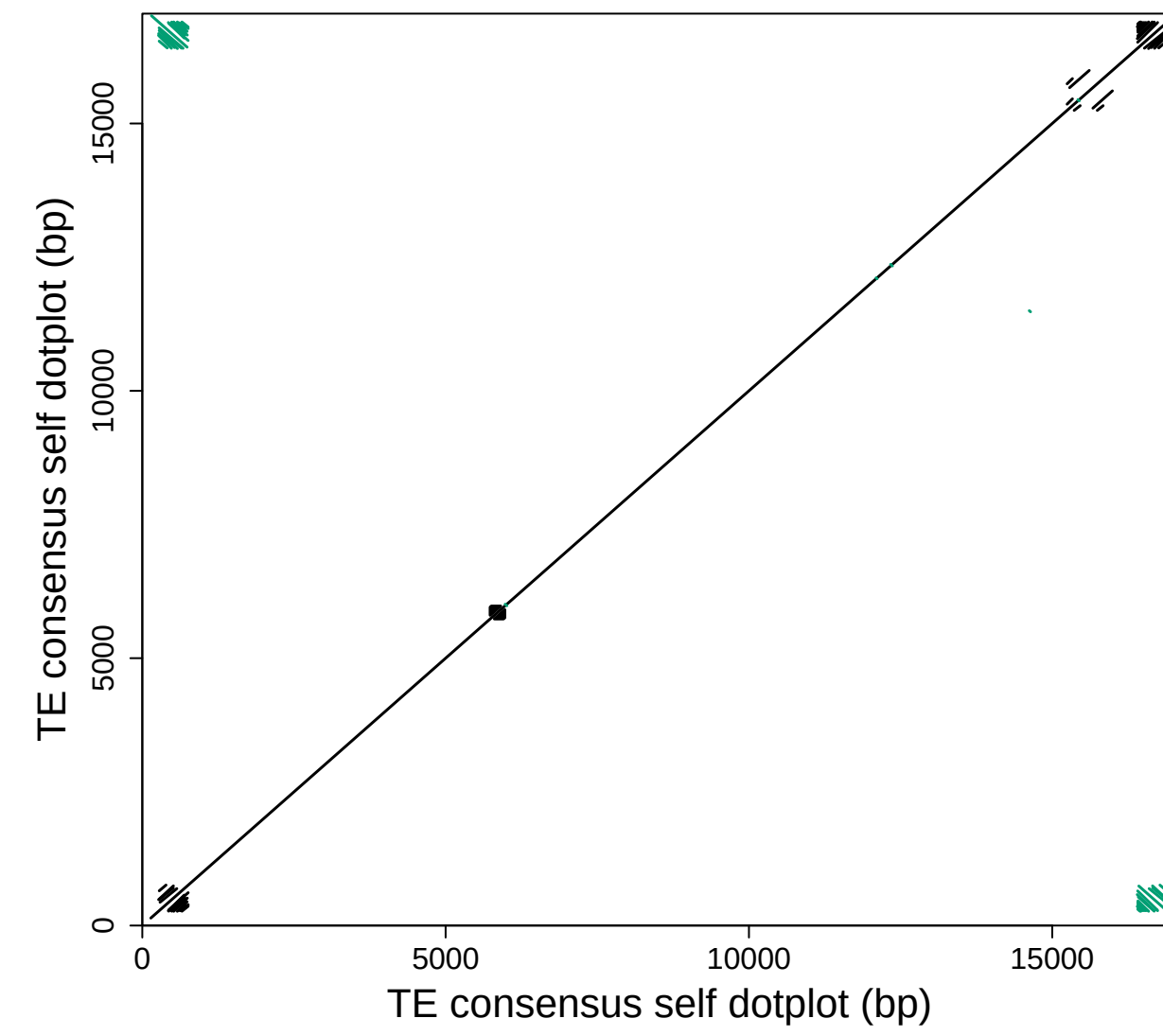
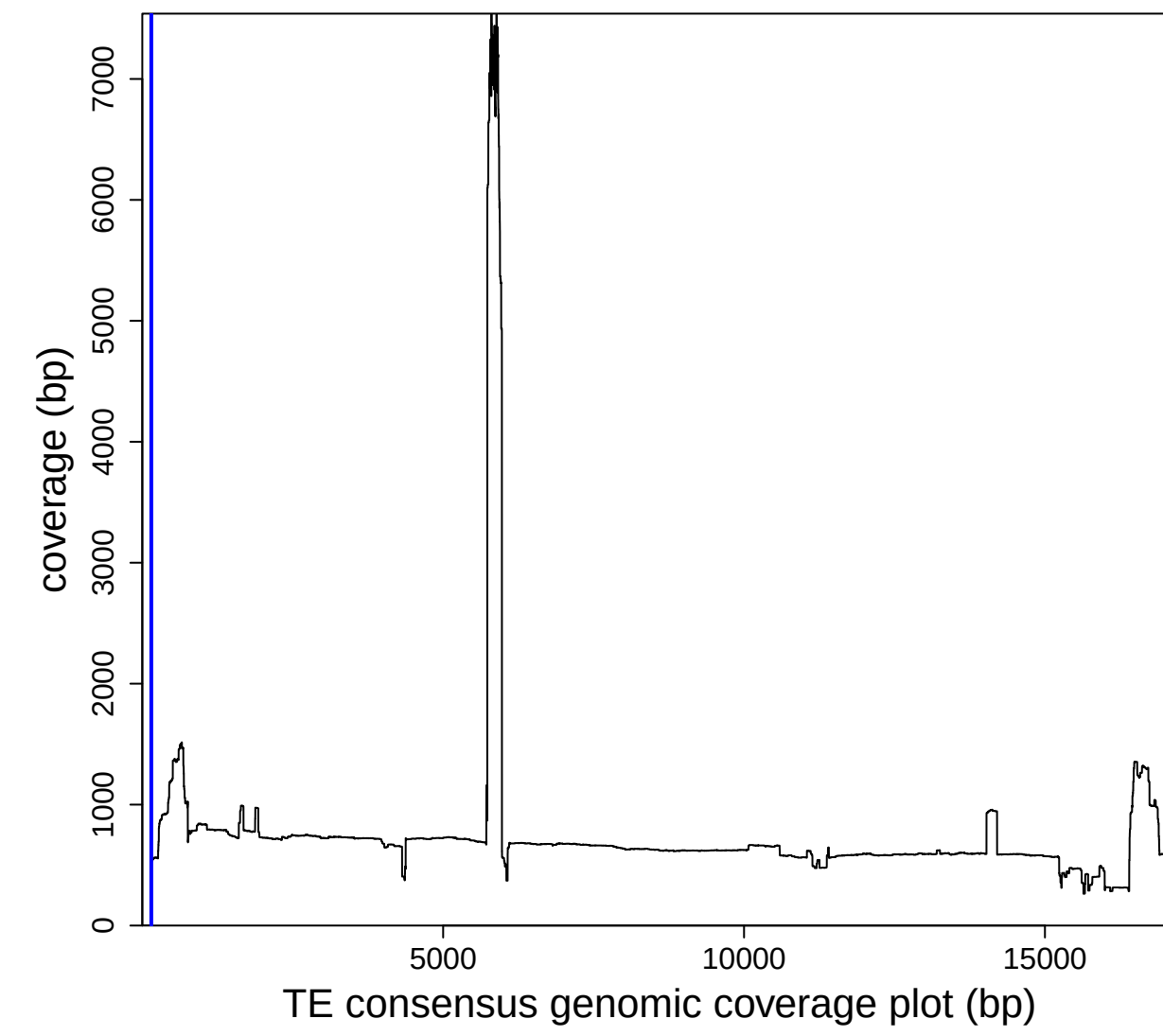
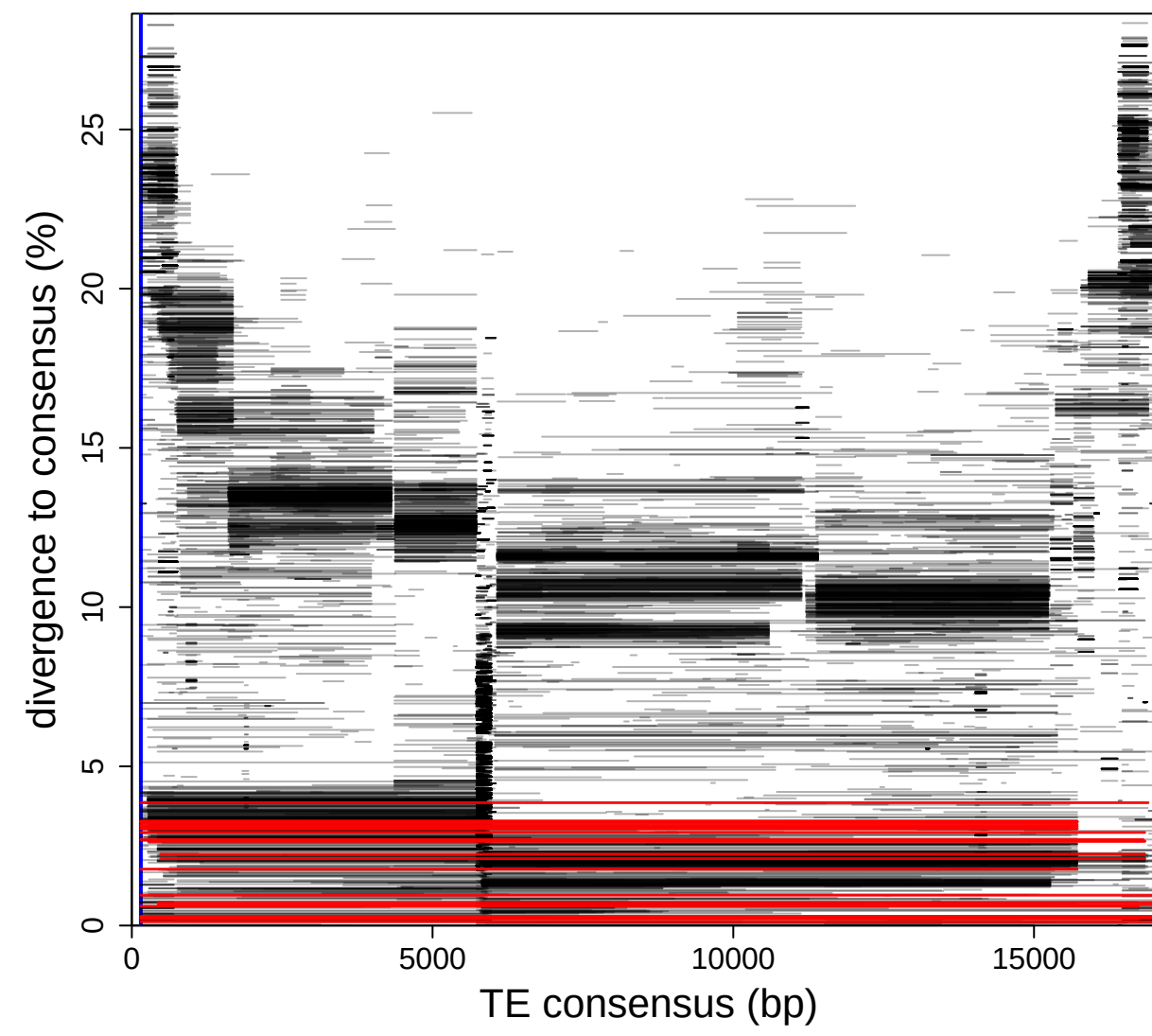
l.b.bed uf.bed g_4.bed fm_1.bed 0 0 bcln.fa aln.fa cl.fa gs.fa
size: 16891bp; fragments: 17784; full length: 43 (>=15201.9bp)

After TEtrimmer 16891 bp



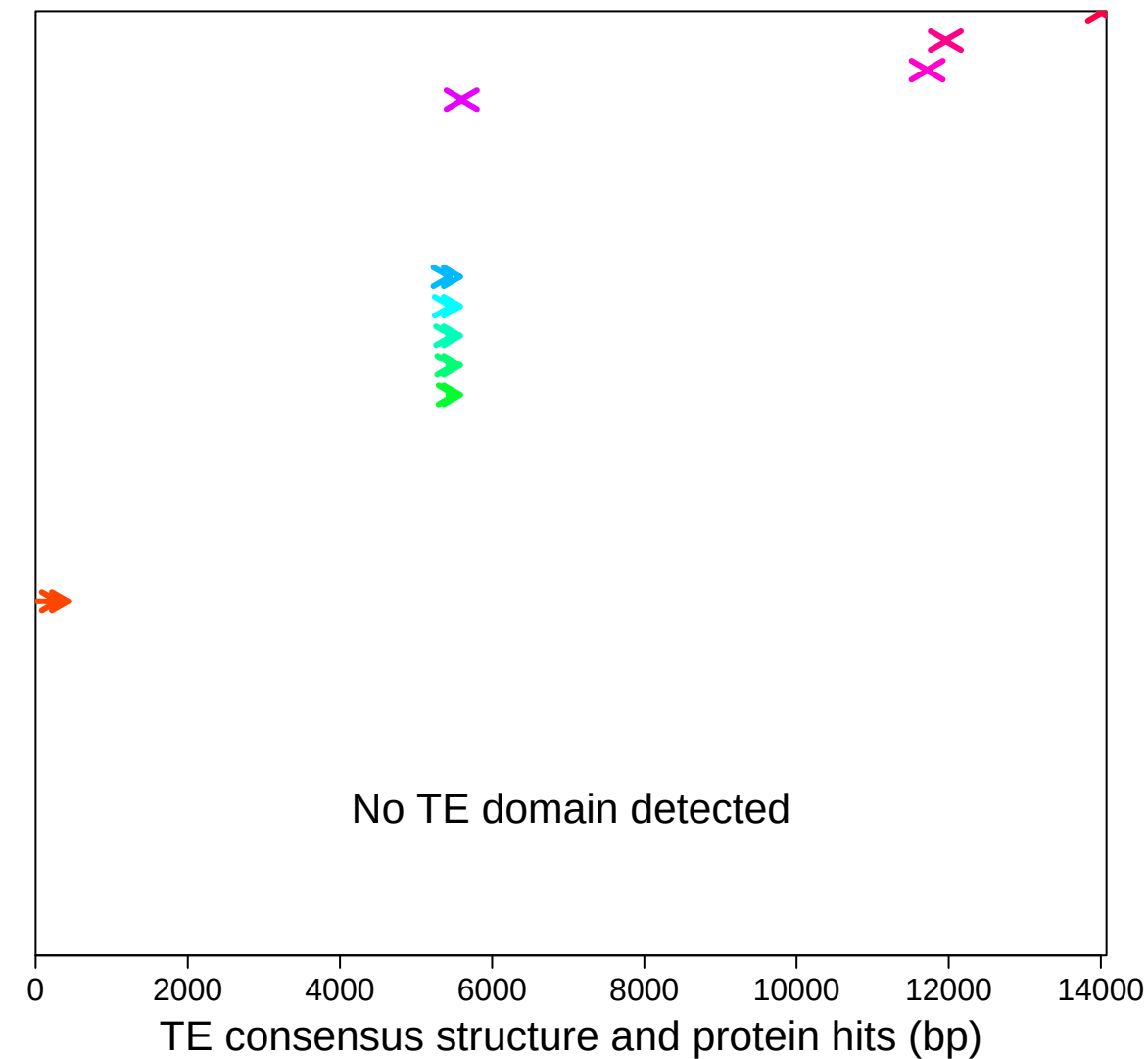
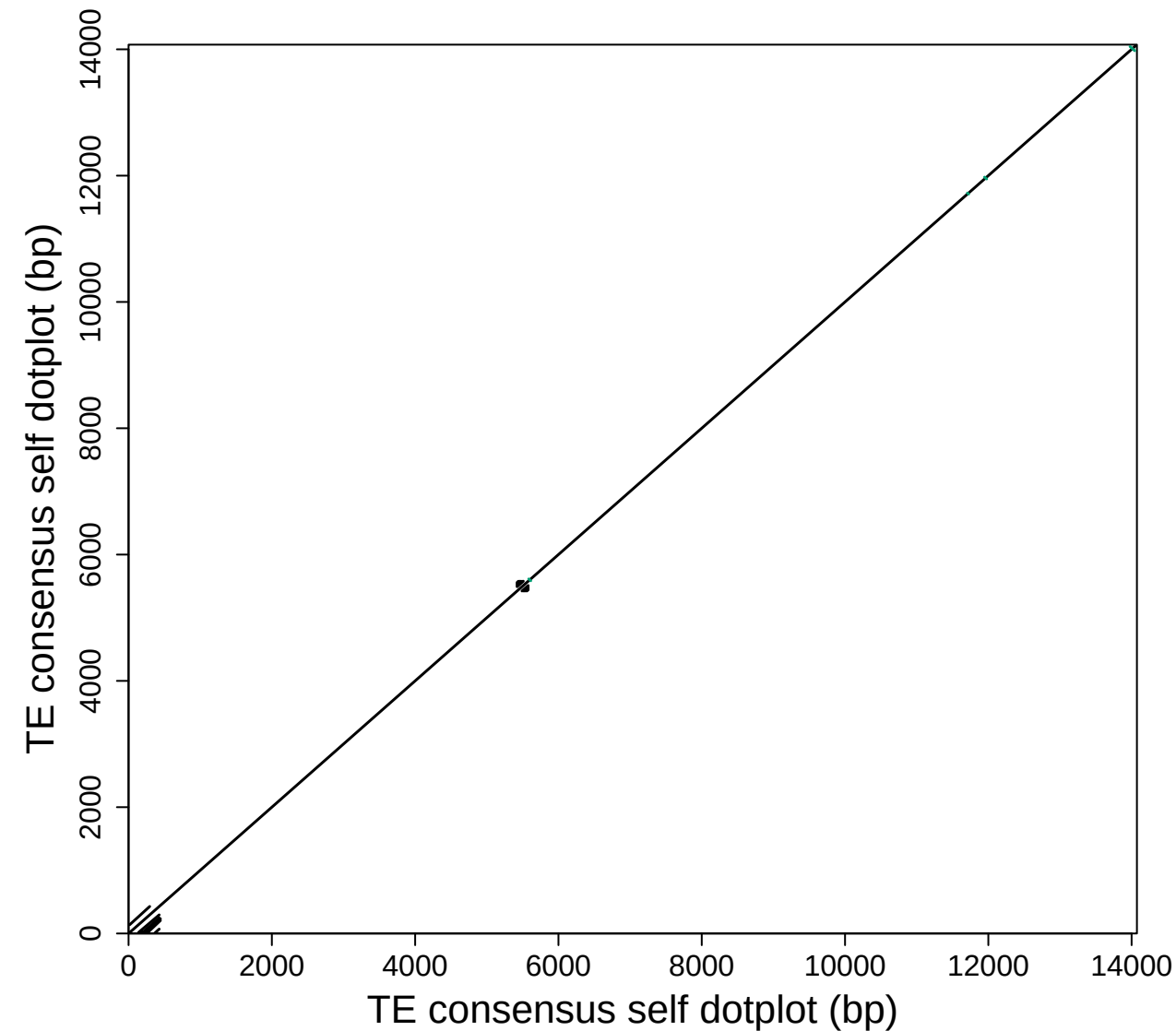
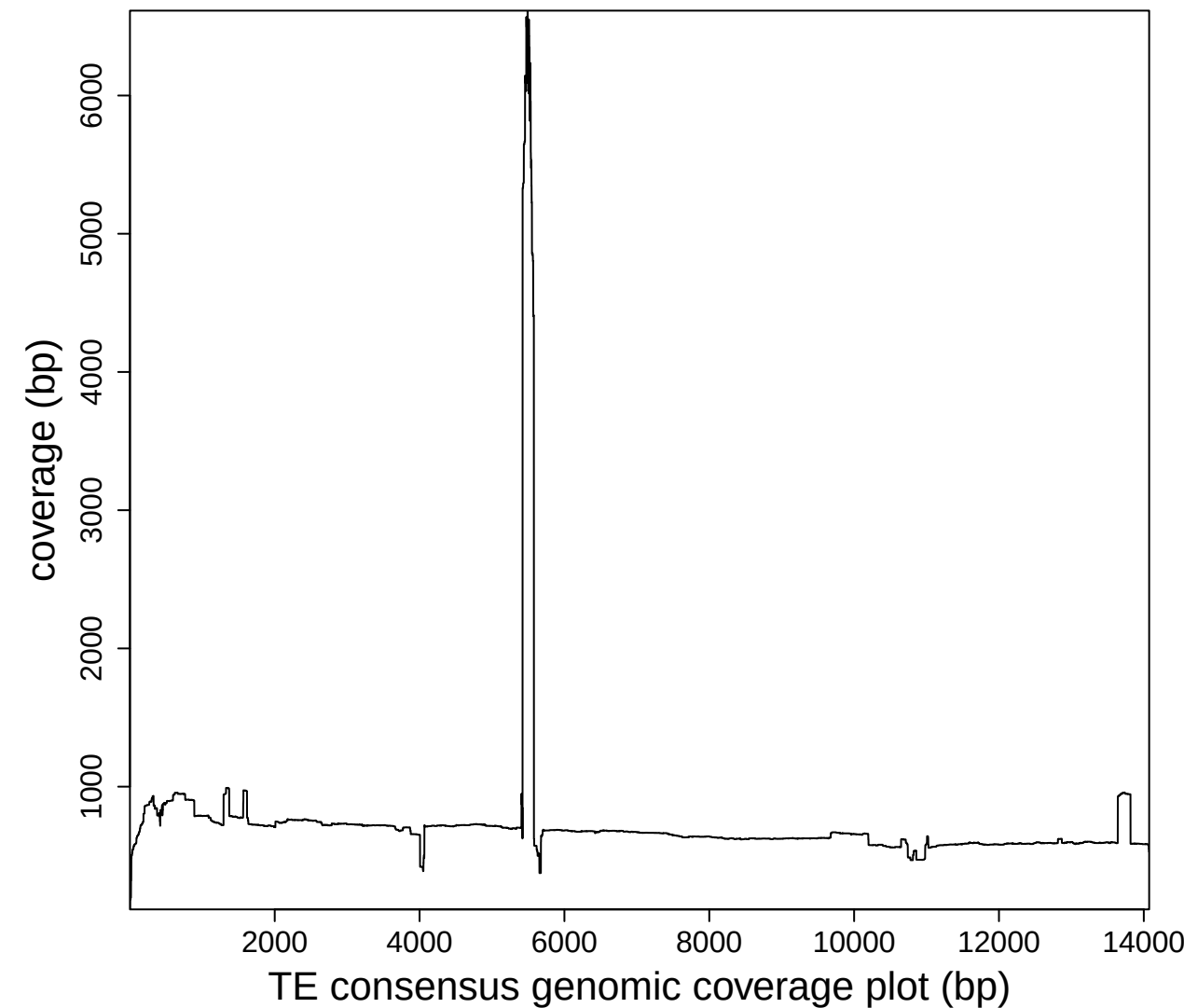
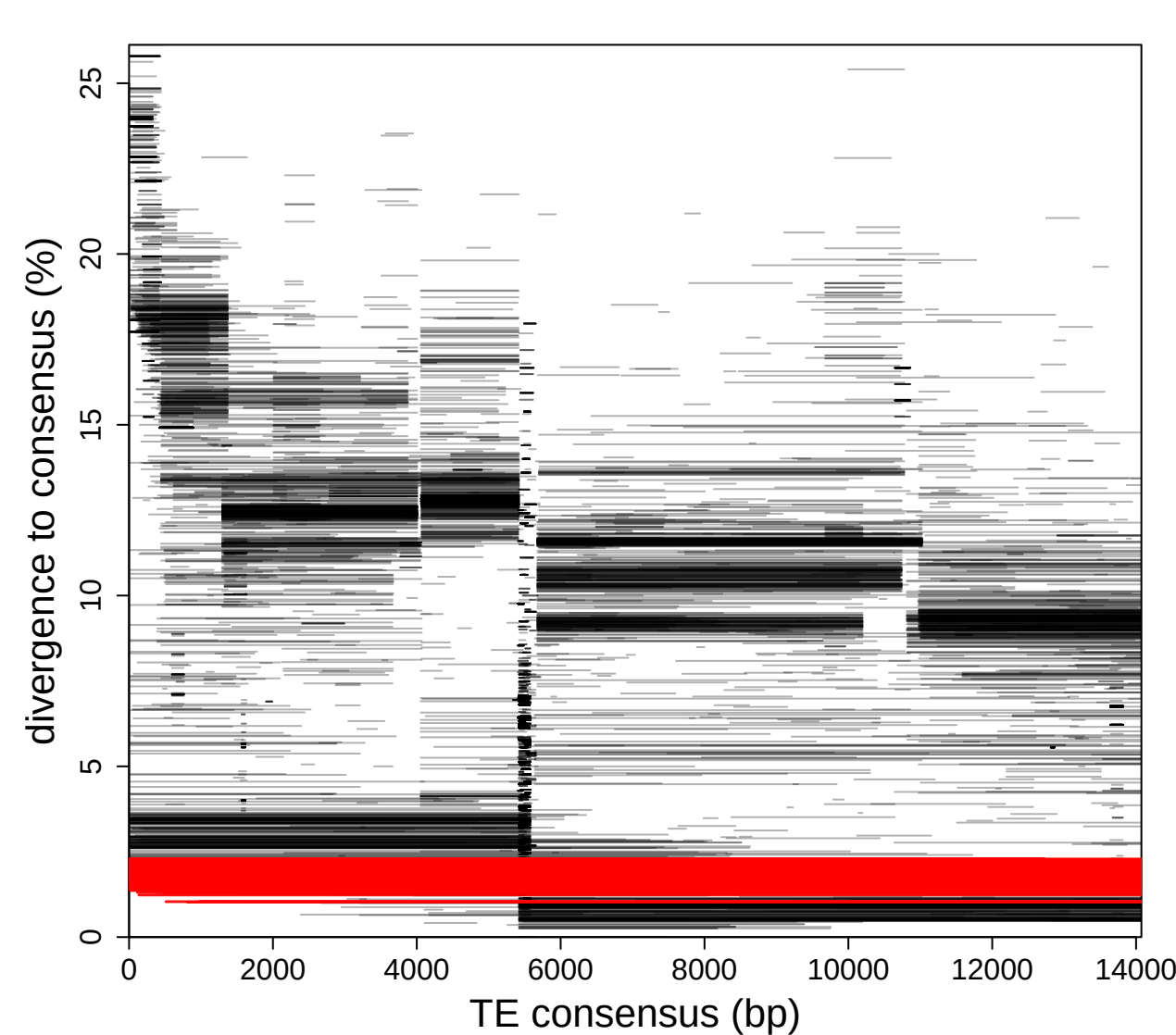
3.fasta.b.bed uf.bed g 4.bed fm 1.bed 0 0 bcln.fa aln.fa cl.fa
size: 17192bp; fragments: 17669; full length: 29 (>=15472.8bp)

After TEtrimmer Extended plot Blue lines are boundaries

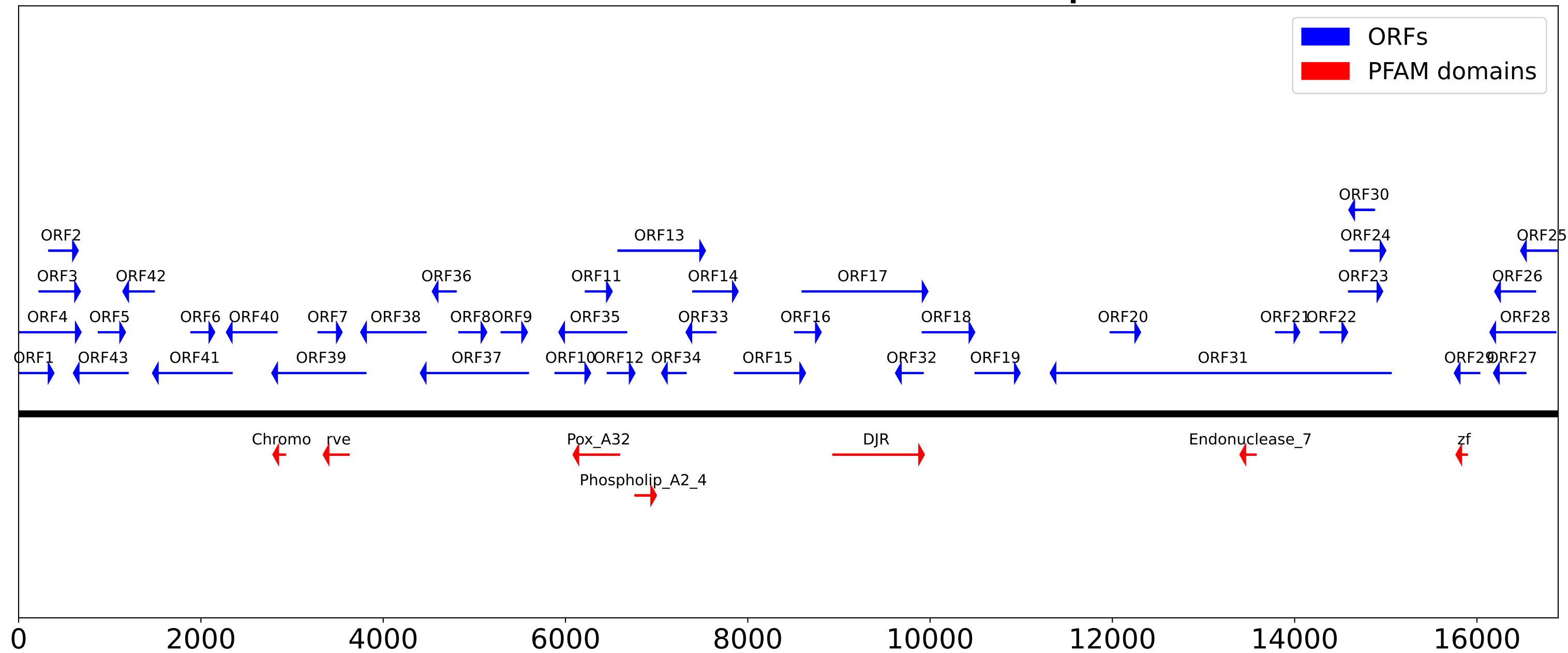


TE: rnd_4_family_733
size: 14074bp; fragments: 12196; full length: 133 (≥ 12666.6 b

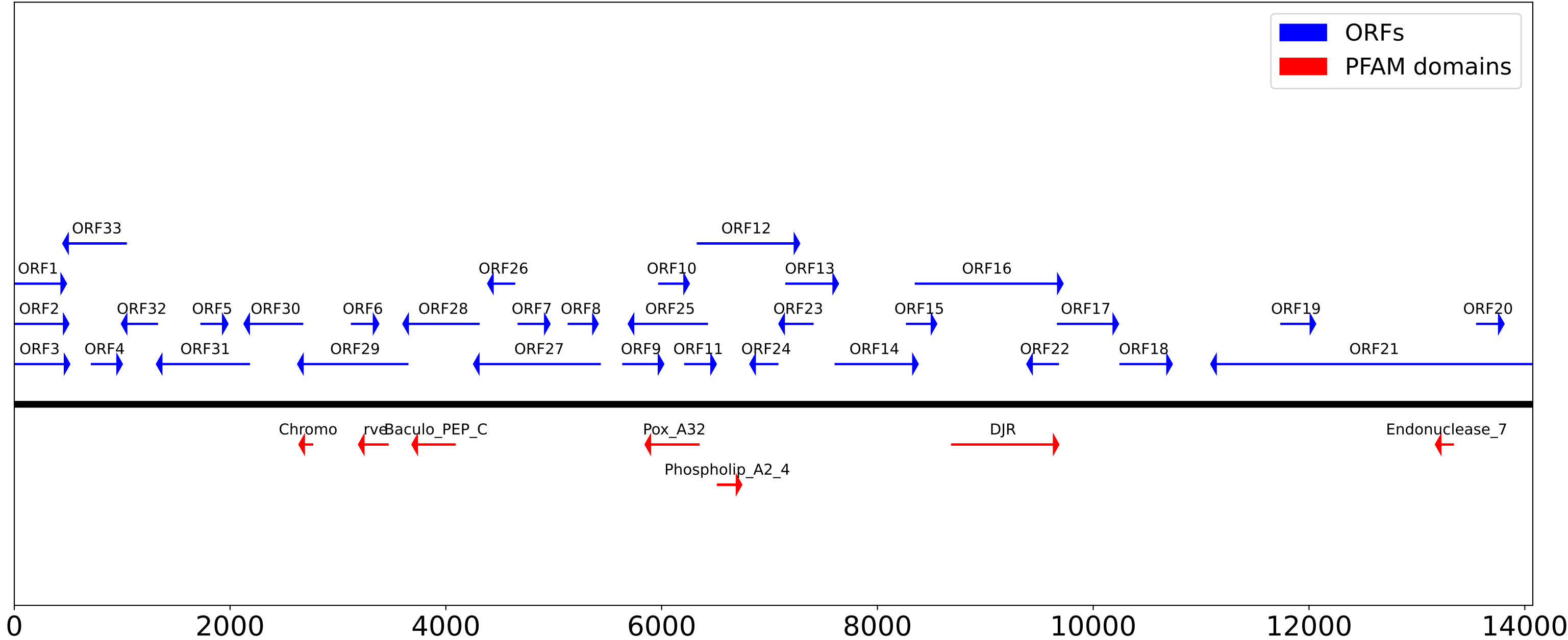
Before TEtrimmer 14074 bp



After TETrimmer ORF and PFAM domain plot



Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

