

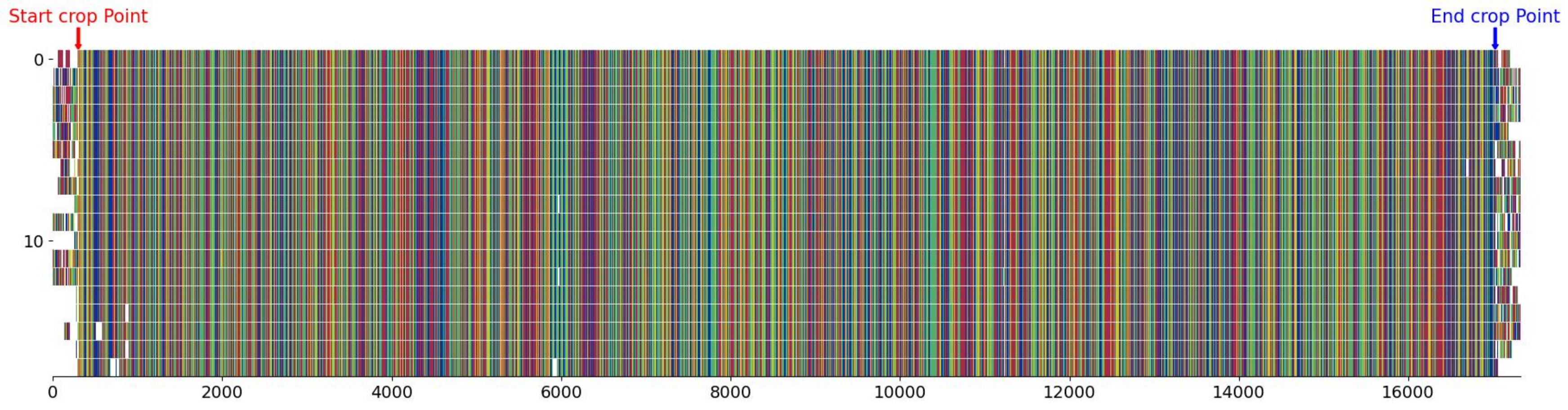
MSA length = 16728

Start crop Point



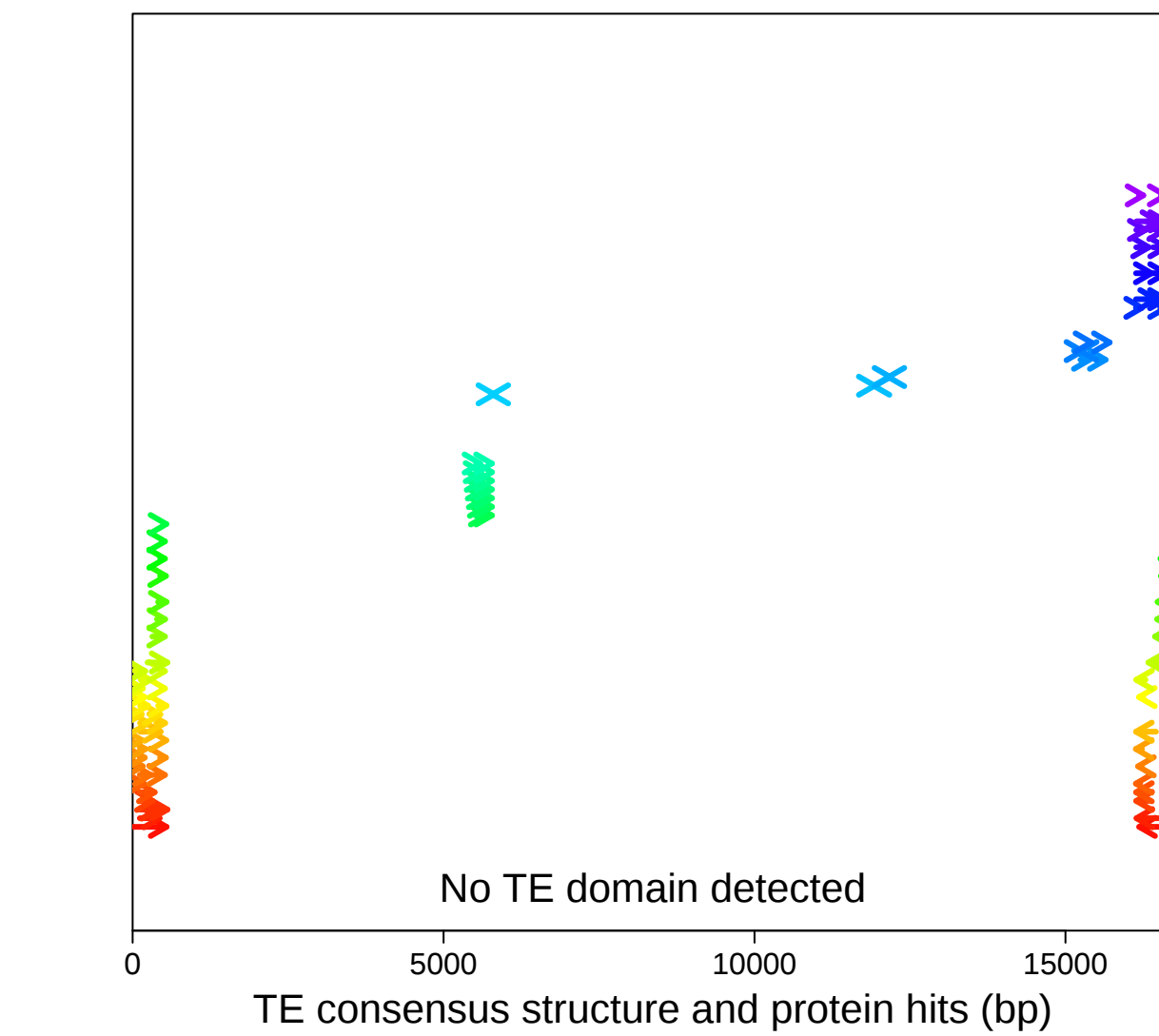
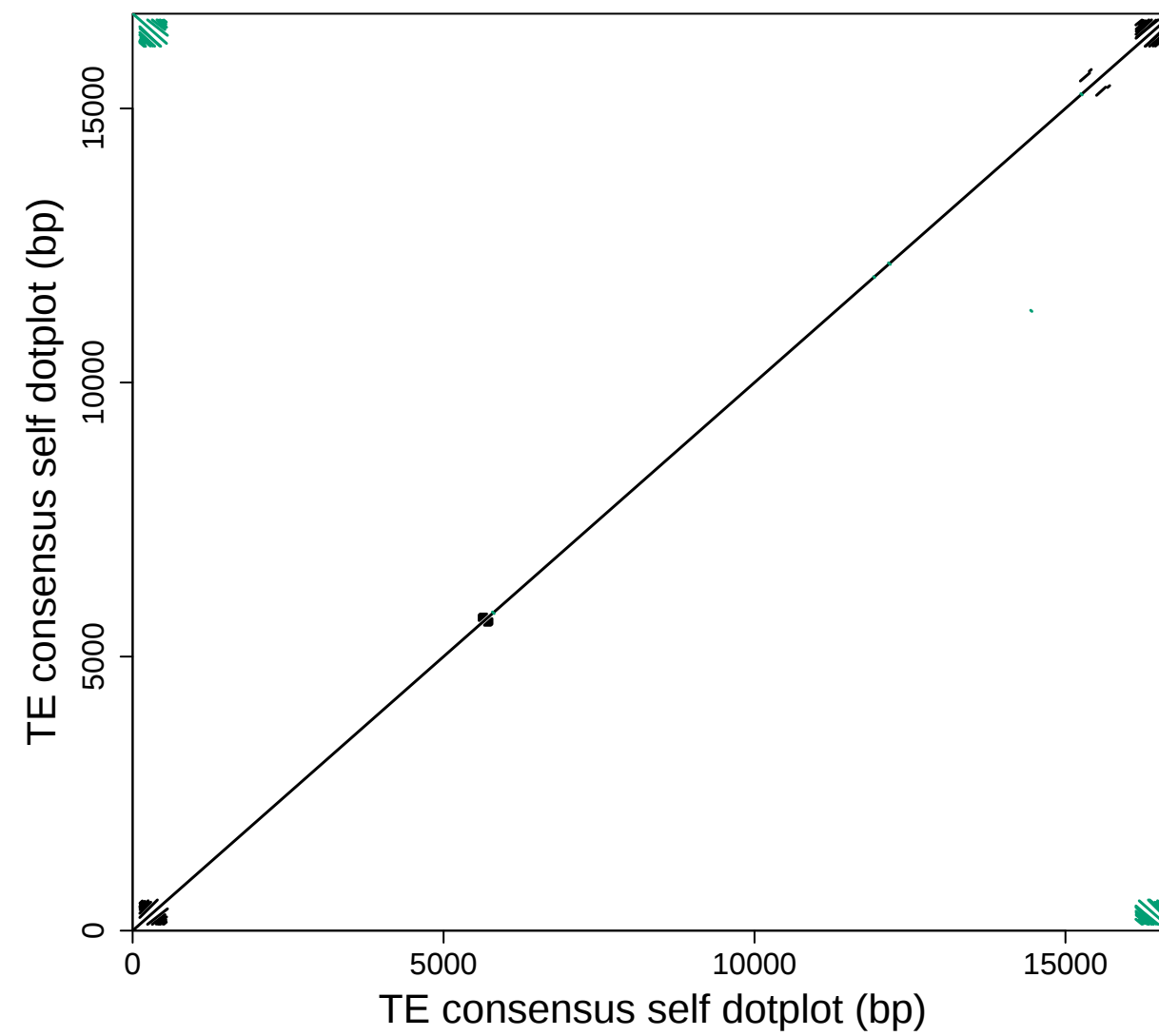
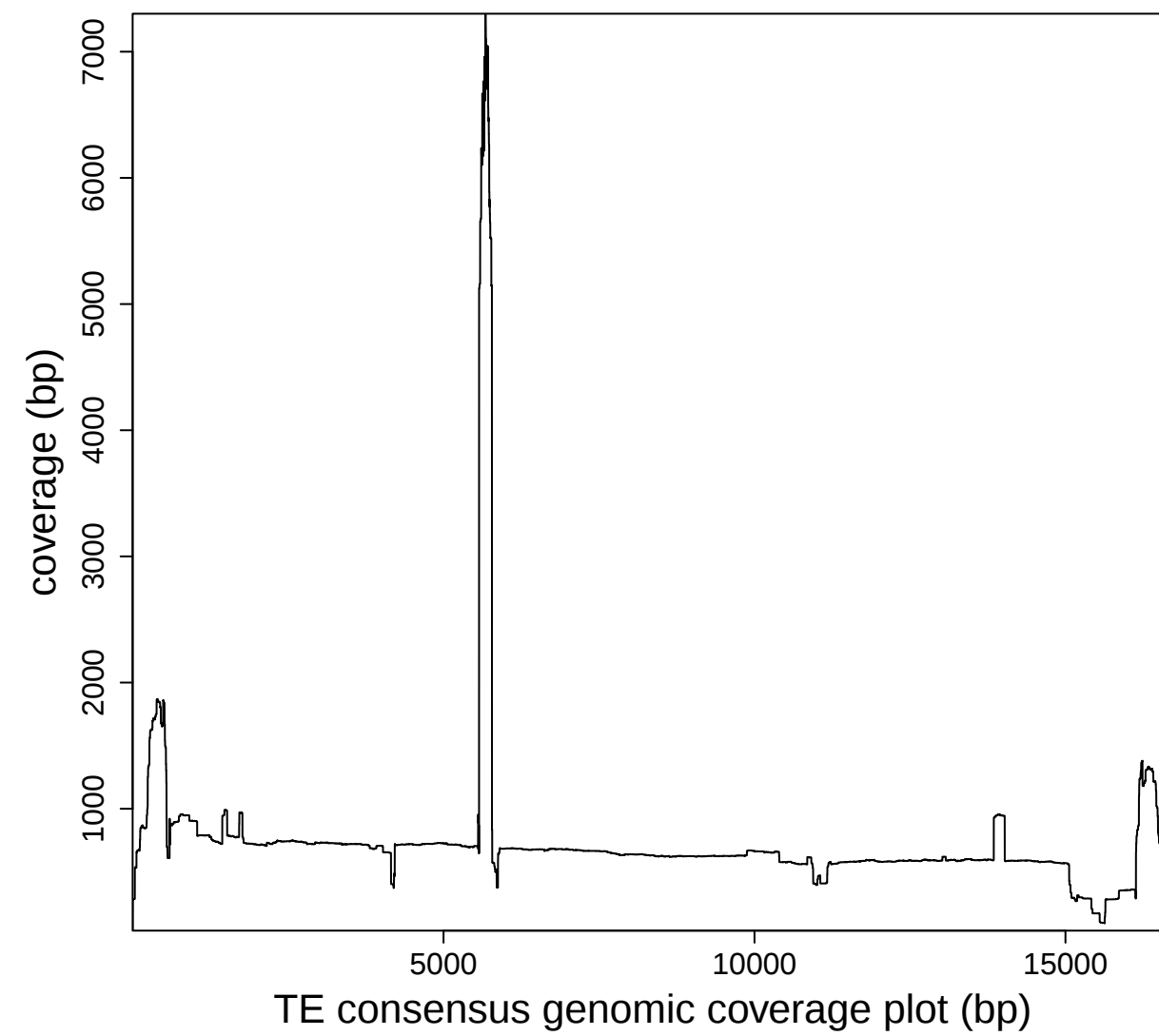
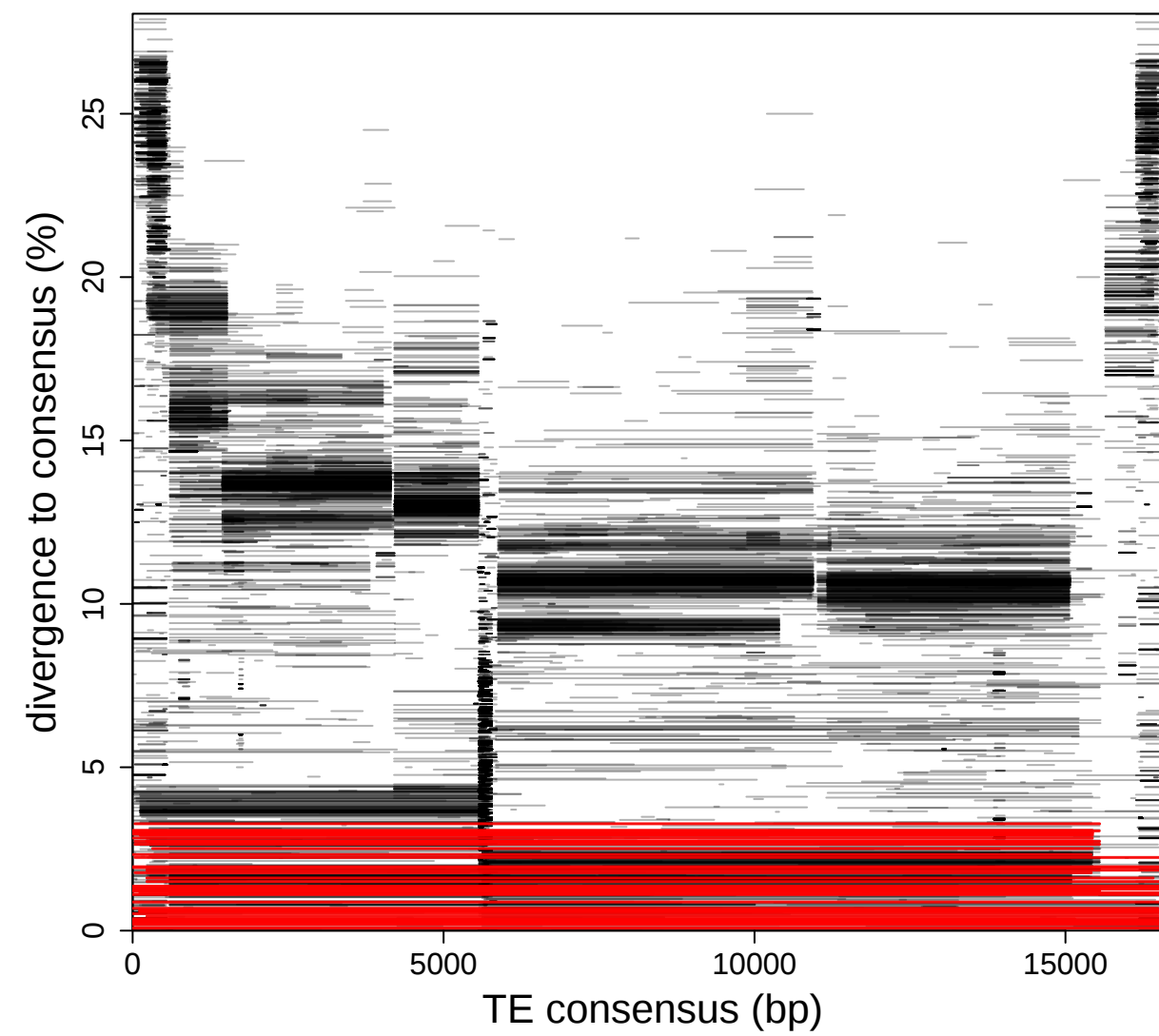
End crop Point





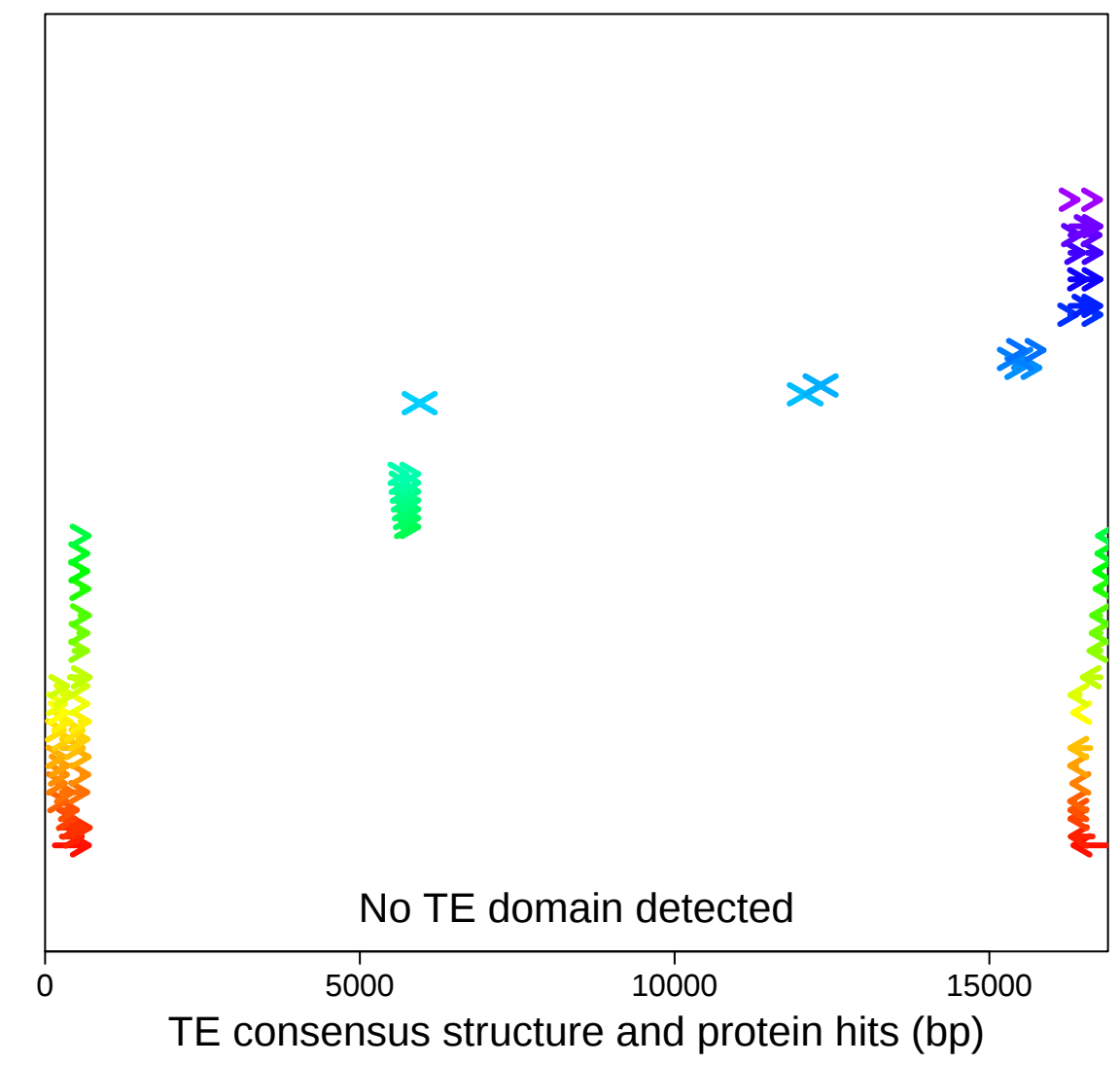
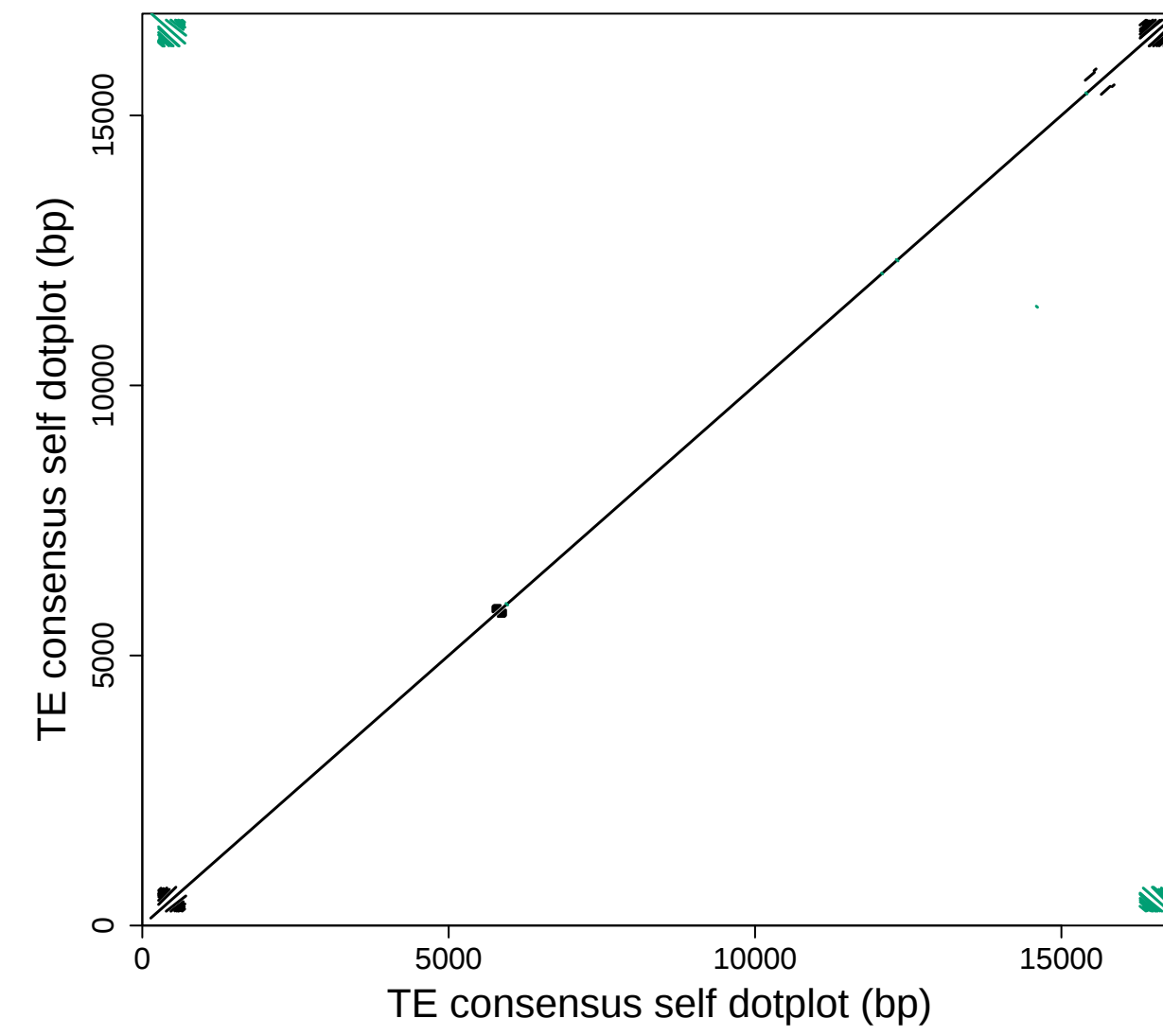
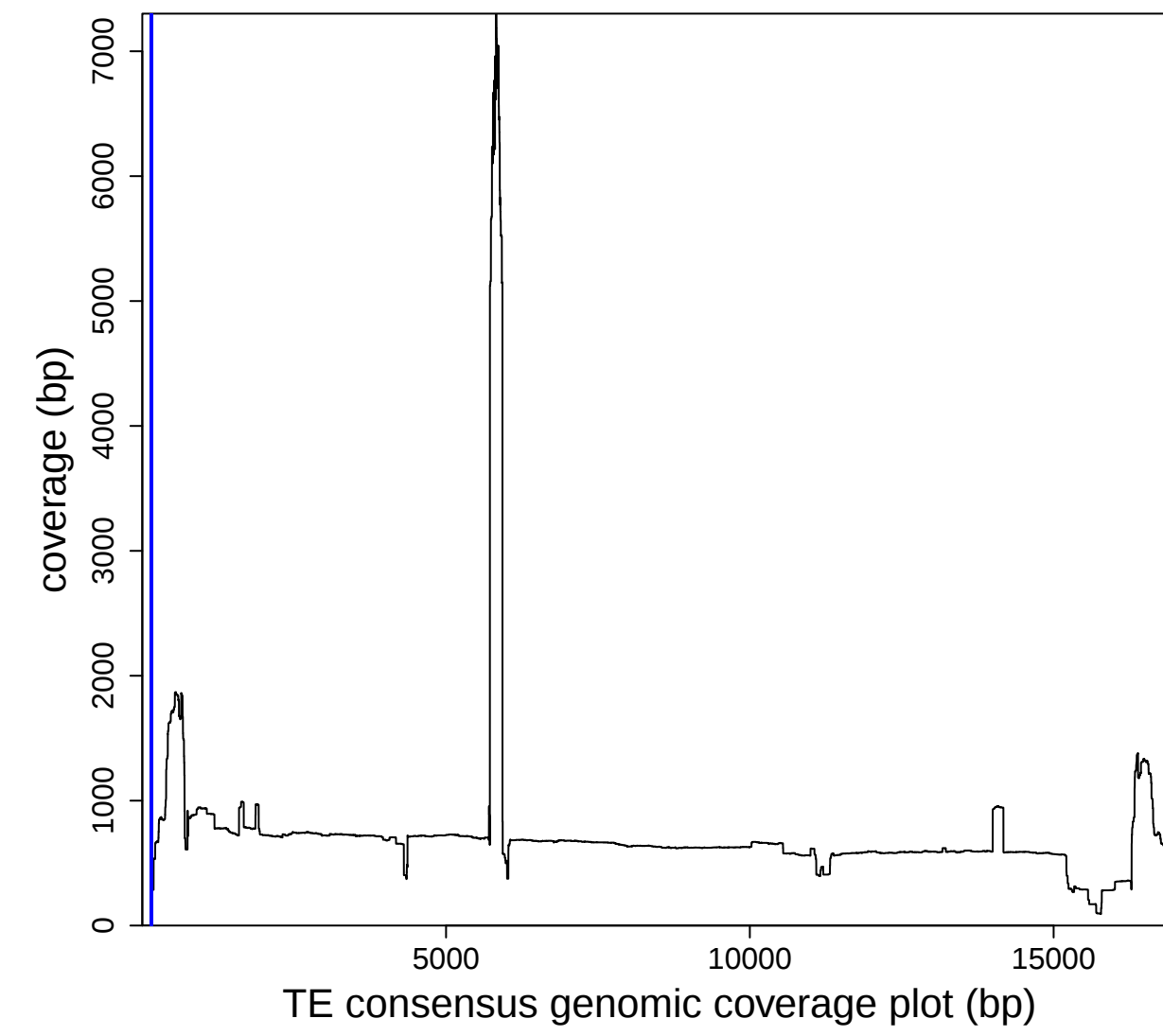
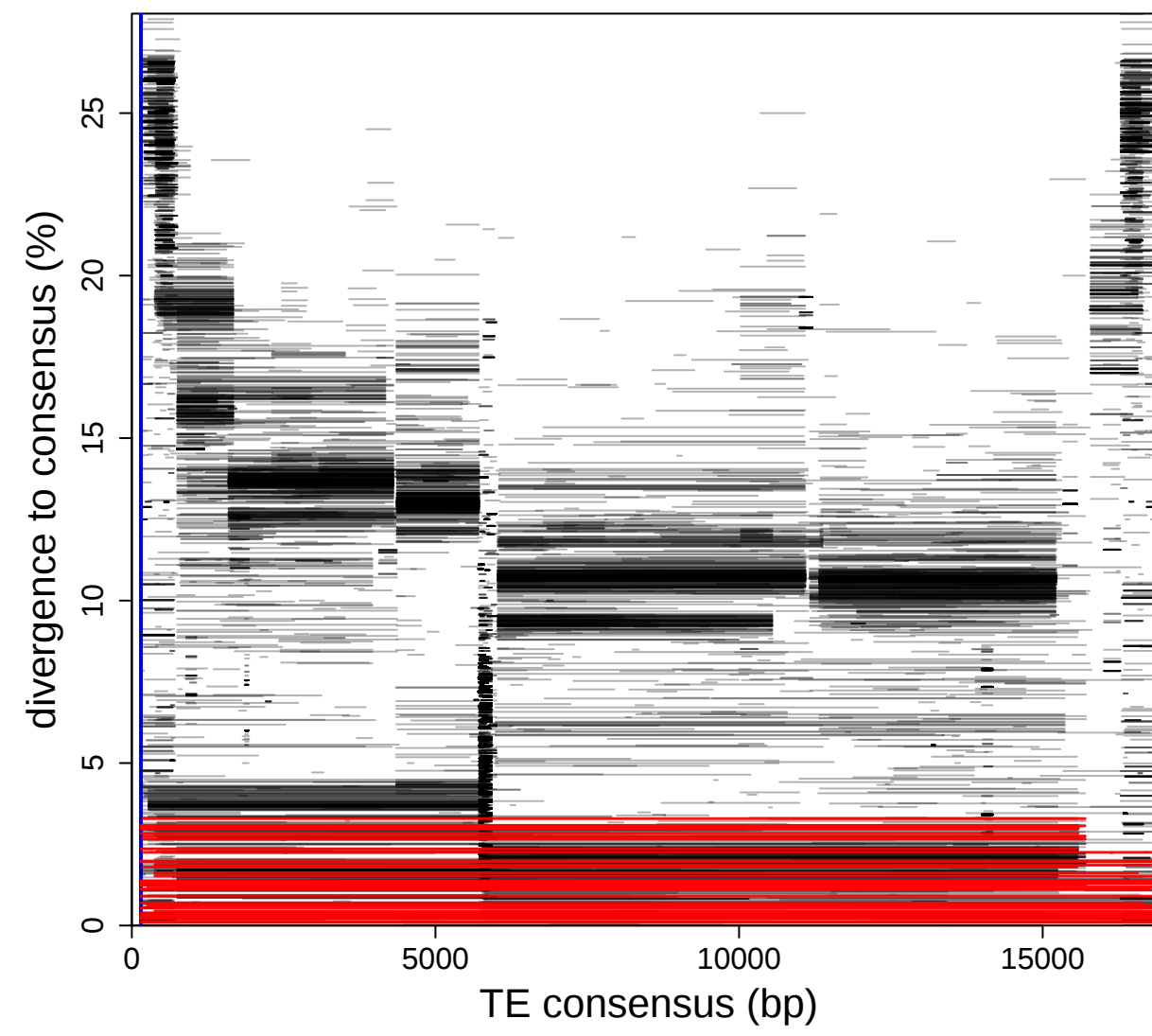
l.b.bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.fa
size: 16728bp; fragments: 16486; full length: 61 (>=15055.2bp)

After TEtrimmer 16728 bp



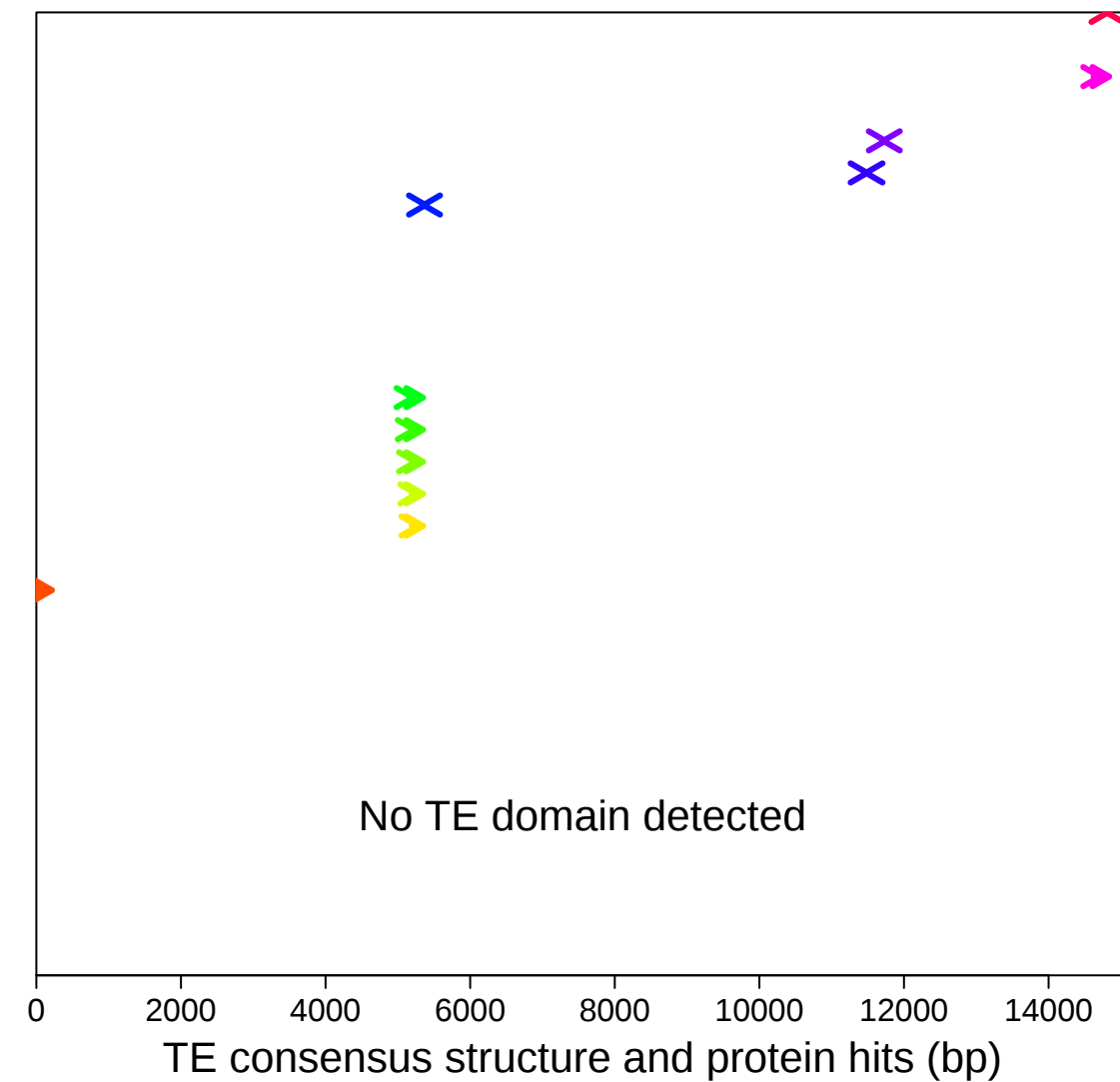
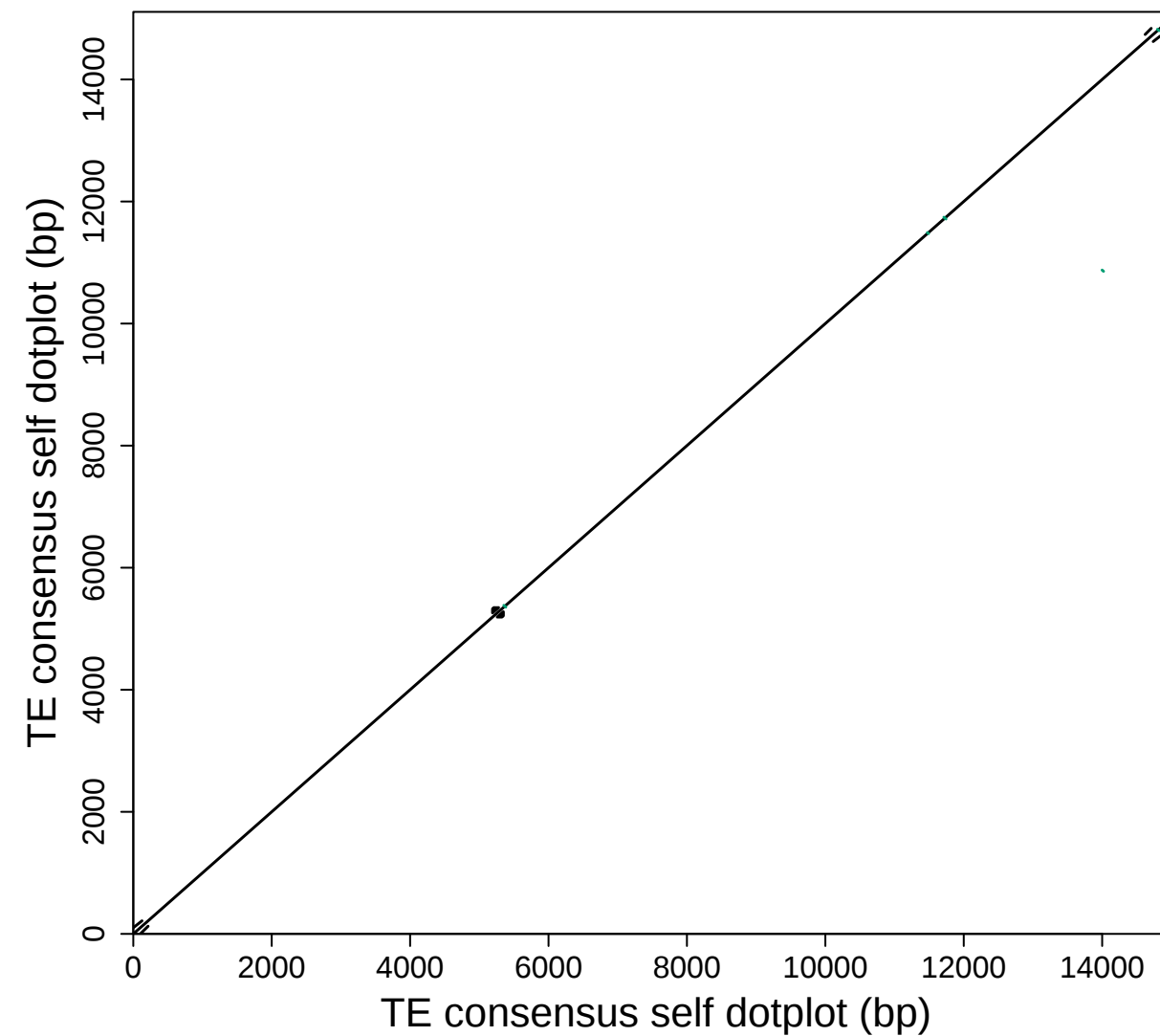
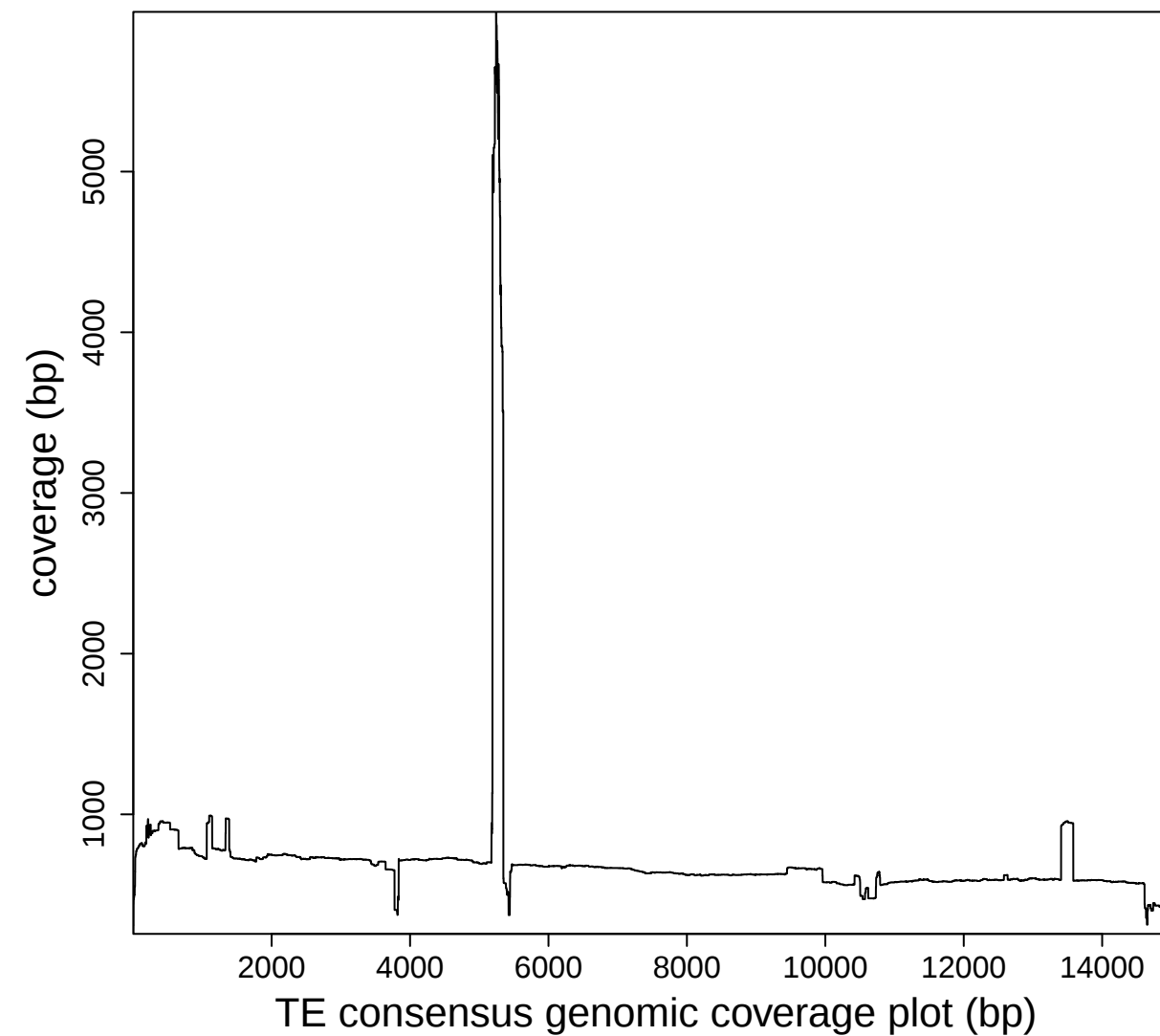
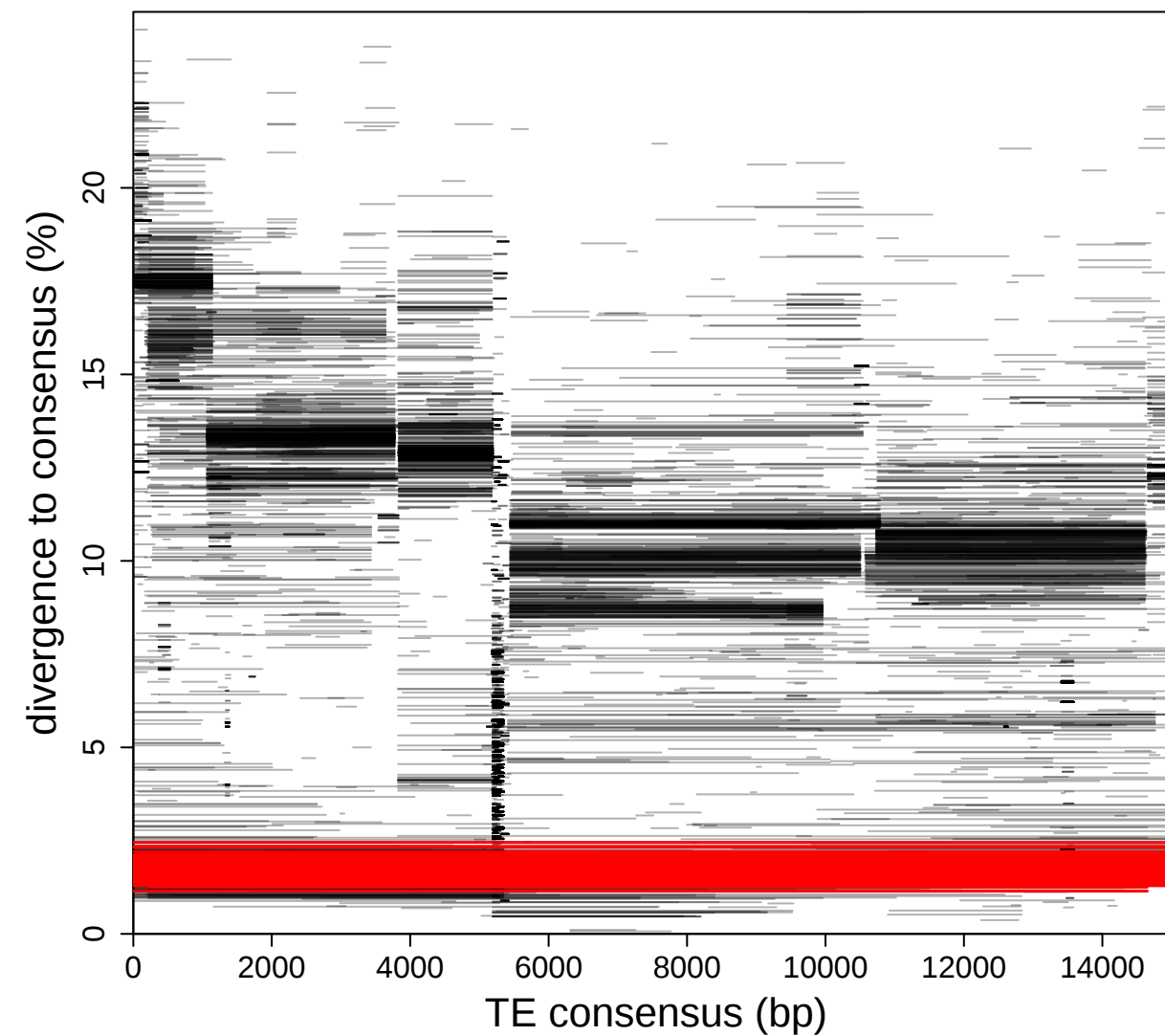
5.fasta.b.bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa
size: 17028bp; fragments: 16474; full length: 54 (>=15325.2bp)

After TEtrimmer Extended plot Blue lines are boundaries

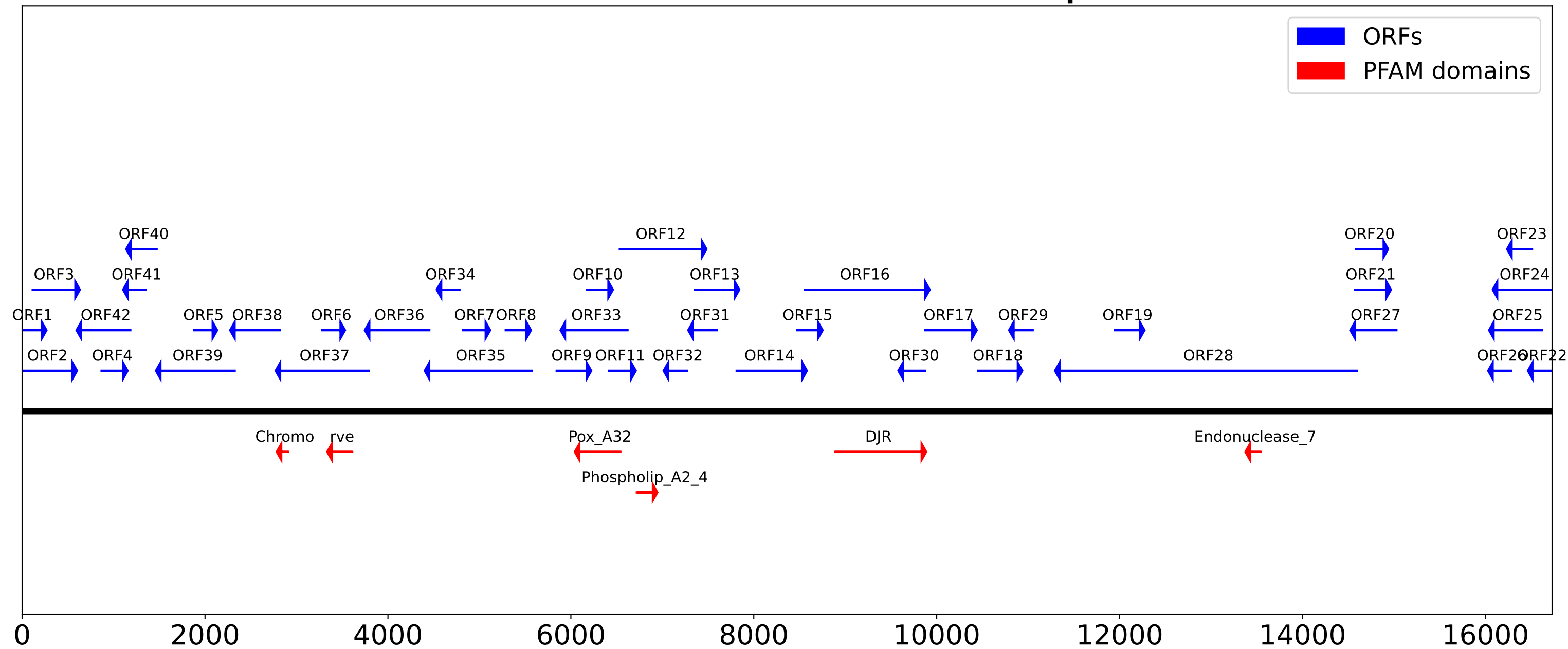


TE: rnd_4_family_735
size: 15107bp; fragments: 11047; full length: 128 (≥ 13596.3 b

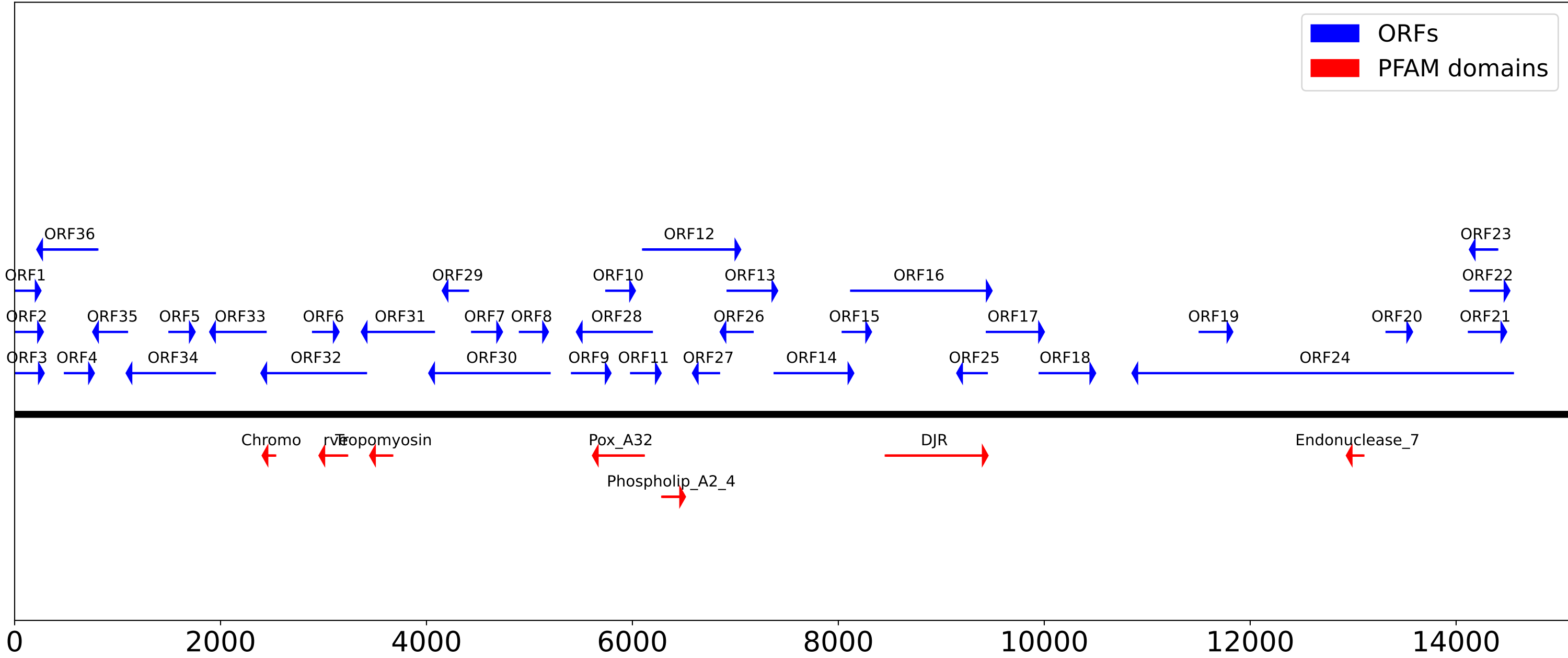
Before TEtrimmer 15107 bp



After TETrimmer ORF and PFAM domain plot



Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

