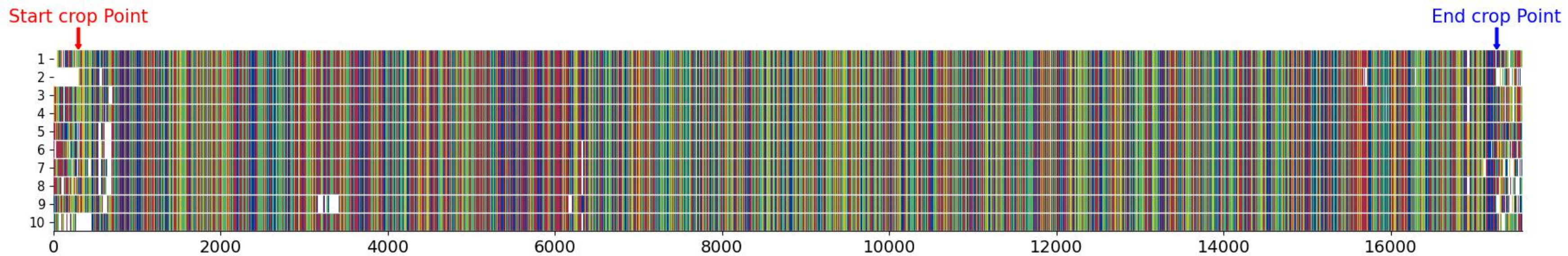


ETIa

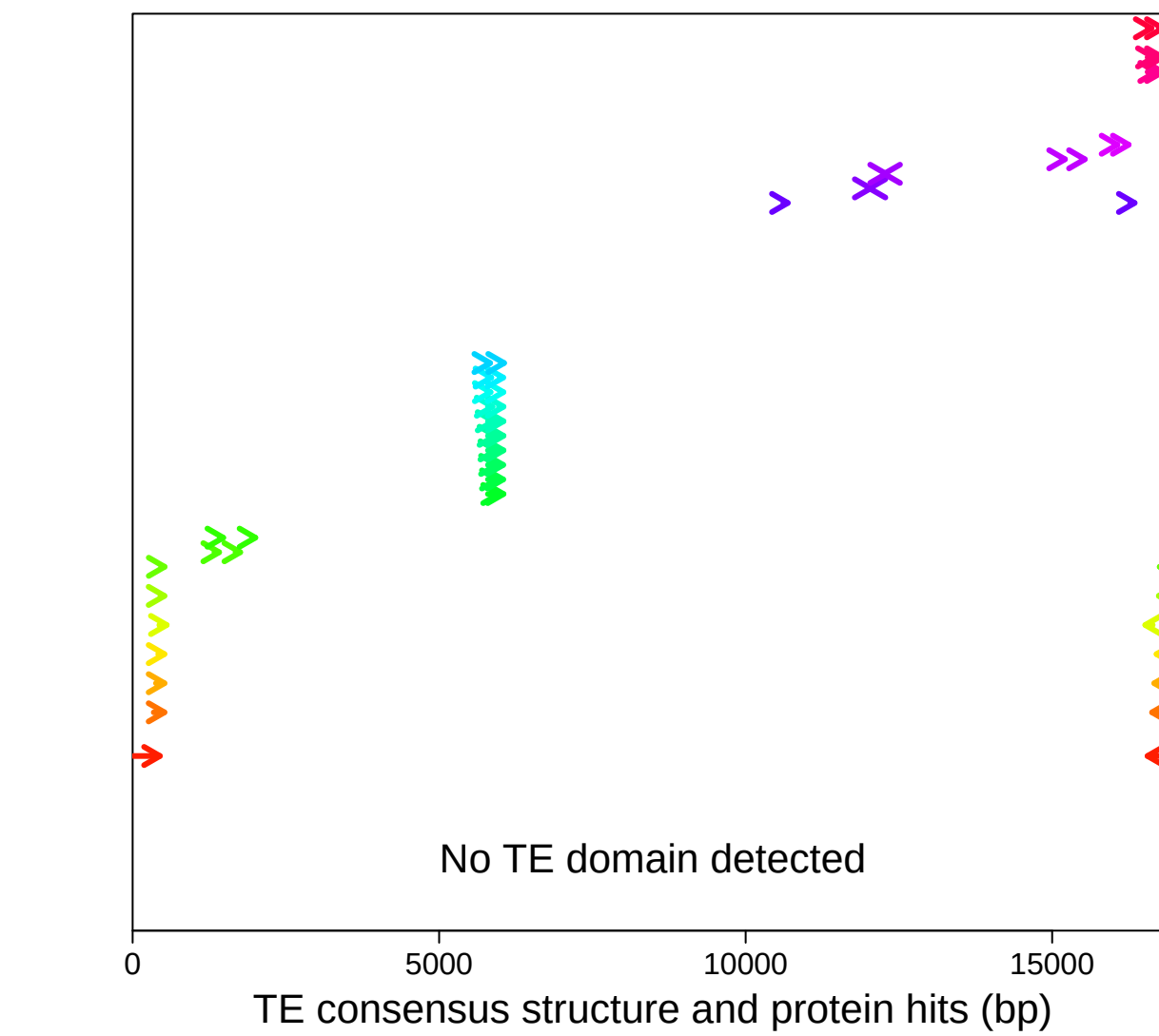
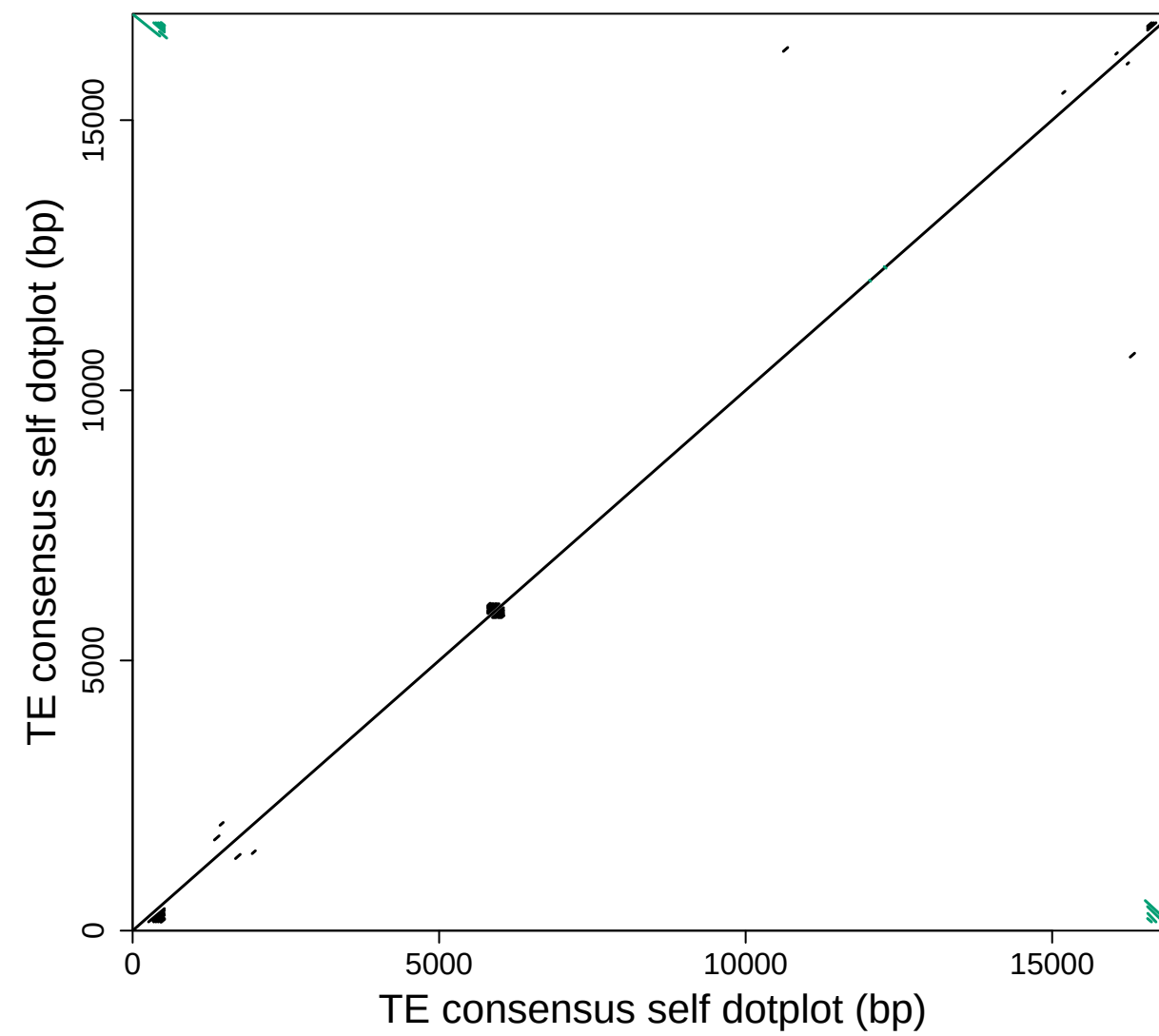
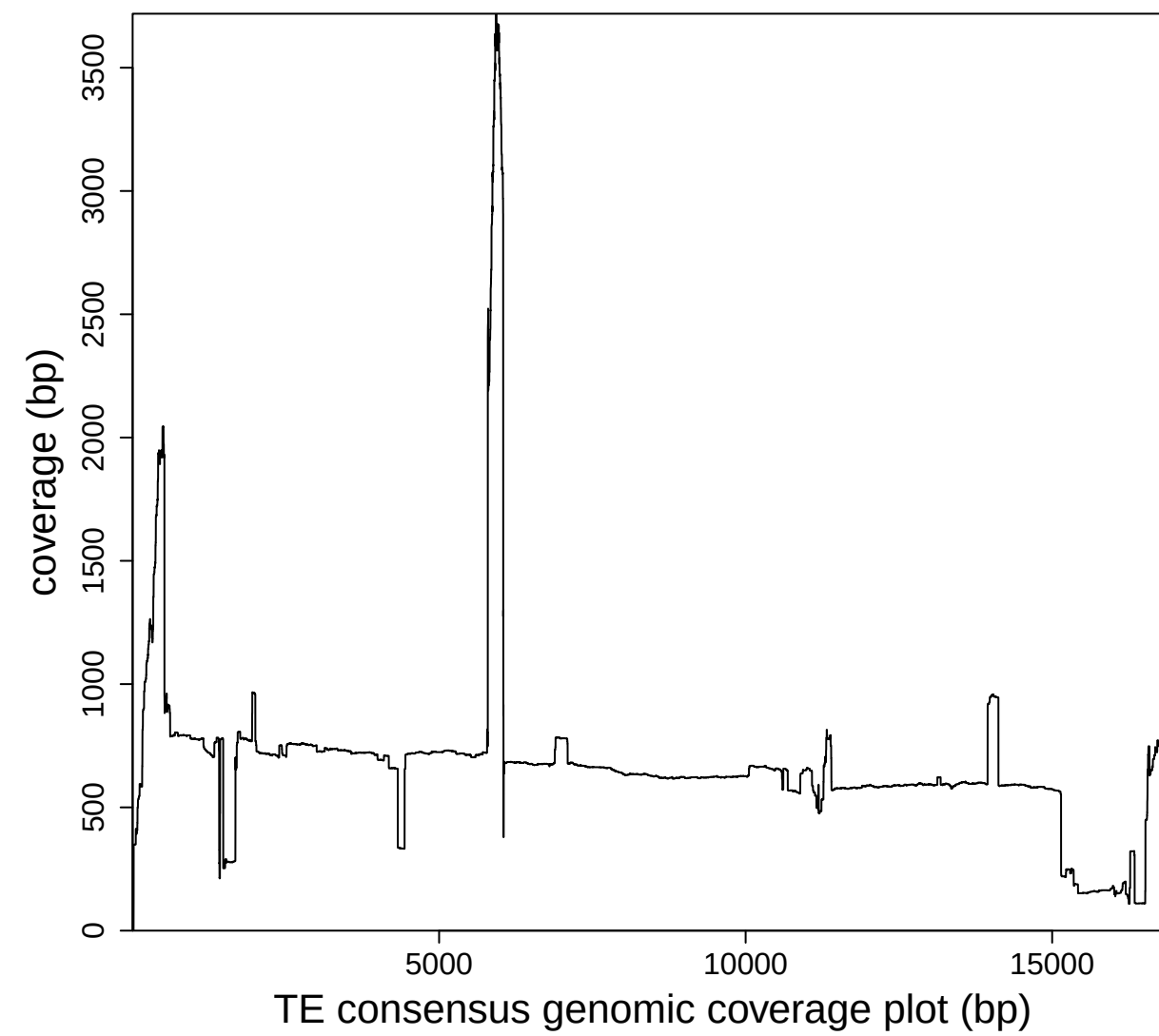
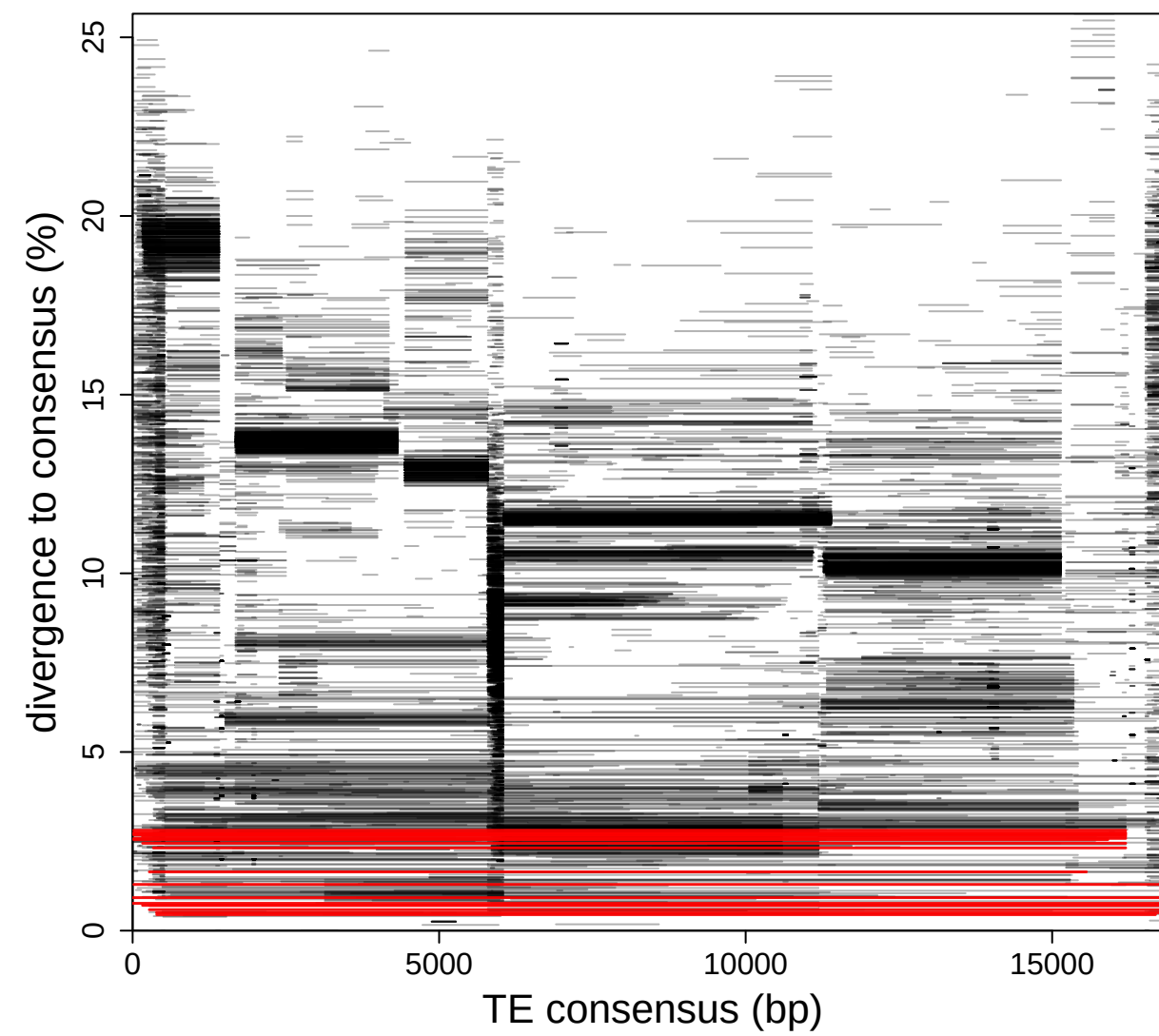
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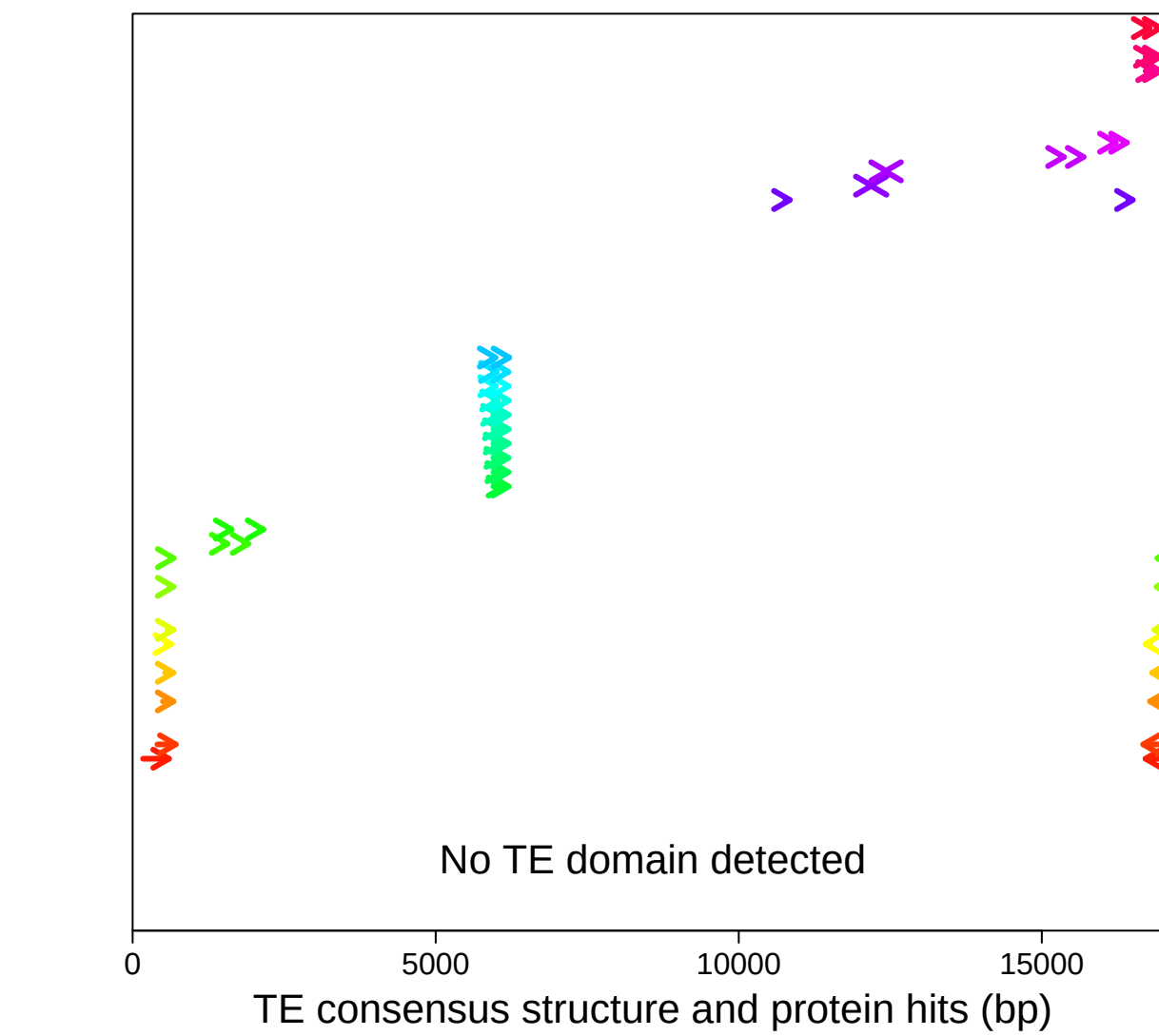
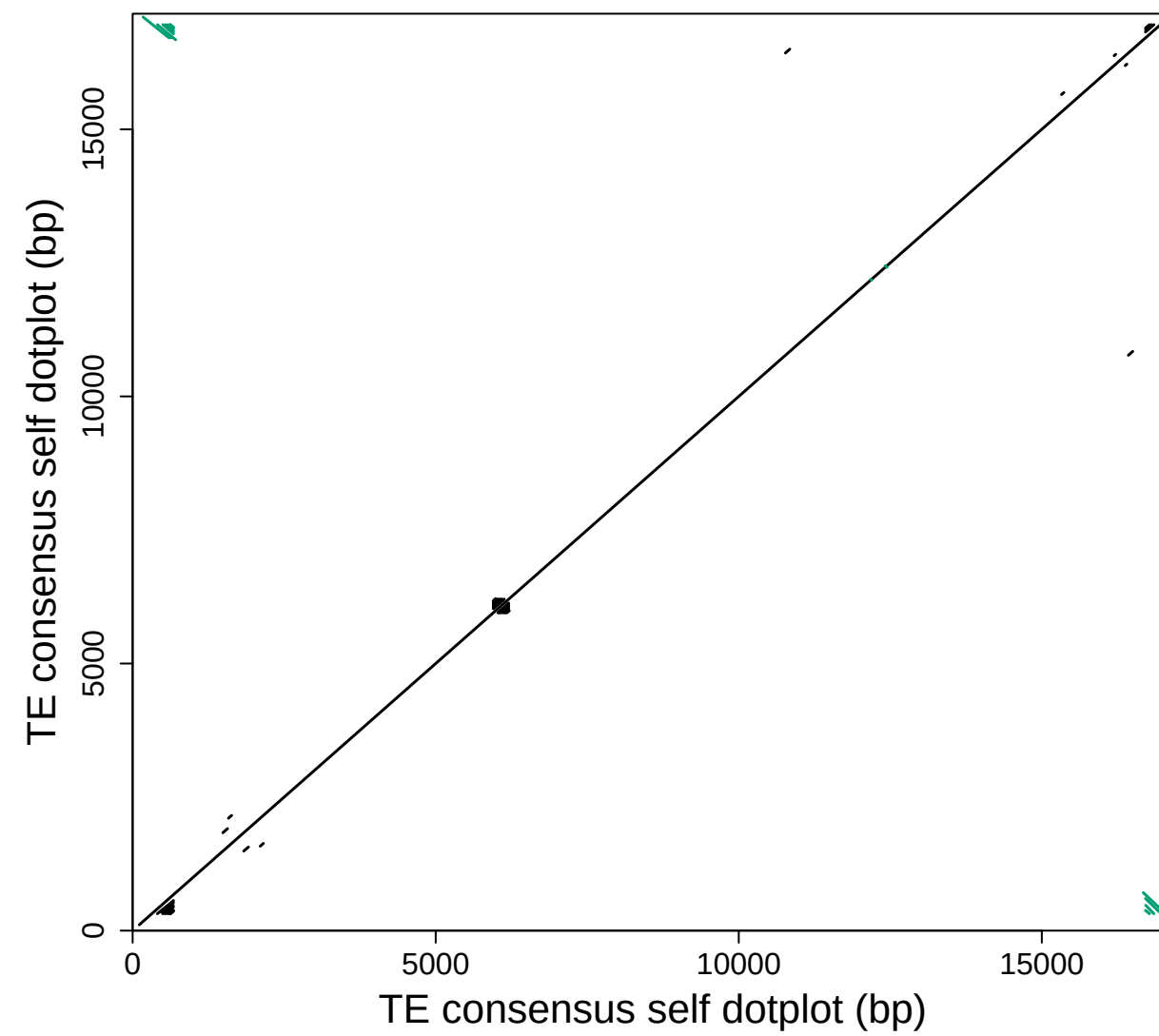
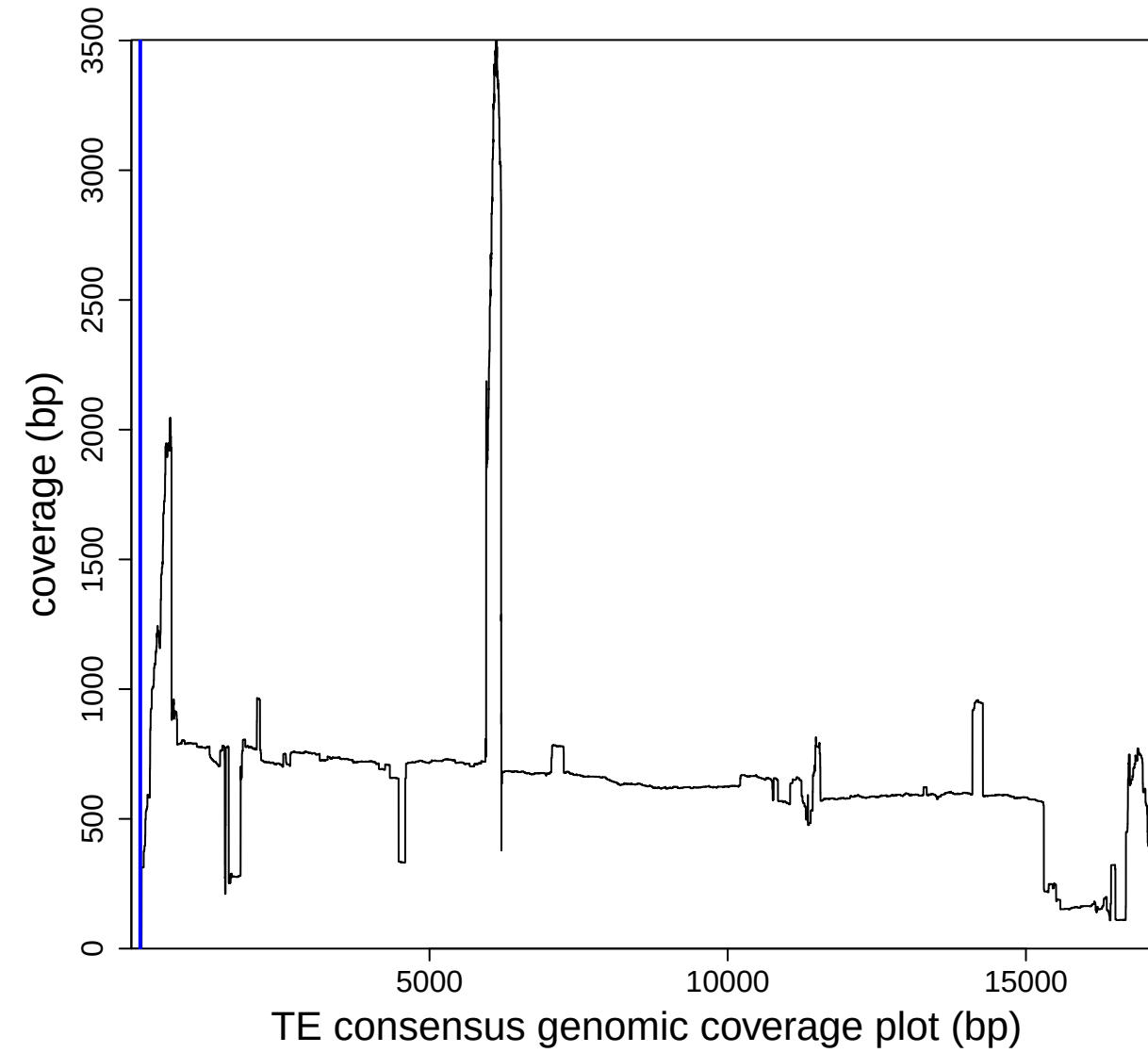
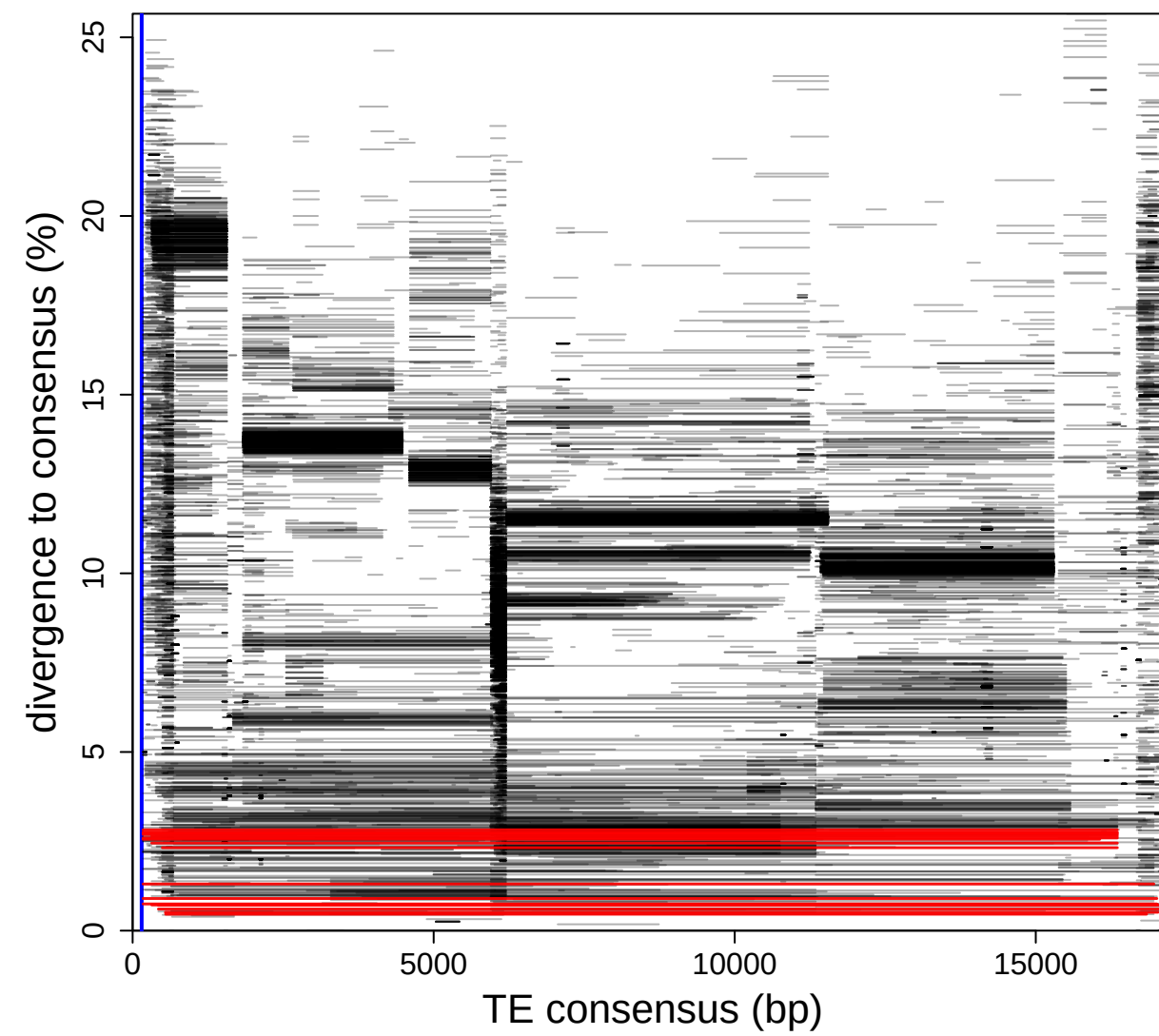
l.b.bed uf.bed g_4.bed fm_1.bed 0 0 bcln.fa aln.fa cl.fa gs.fa
size: 16970bp; fragments: 12765; full length: 18 (>=15273bp)

After TEtrimmer 16970 bp



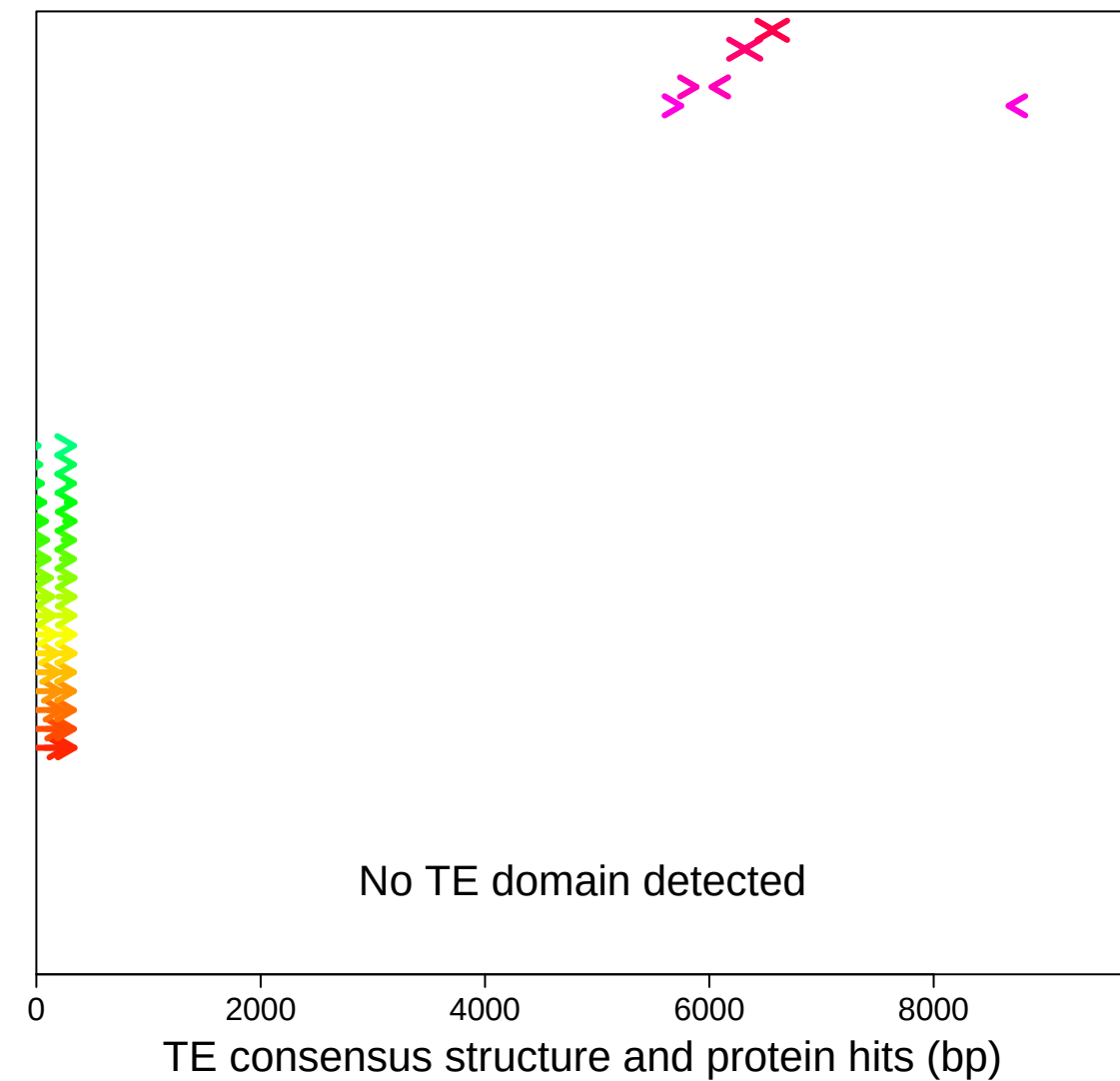
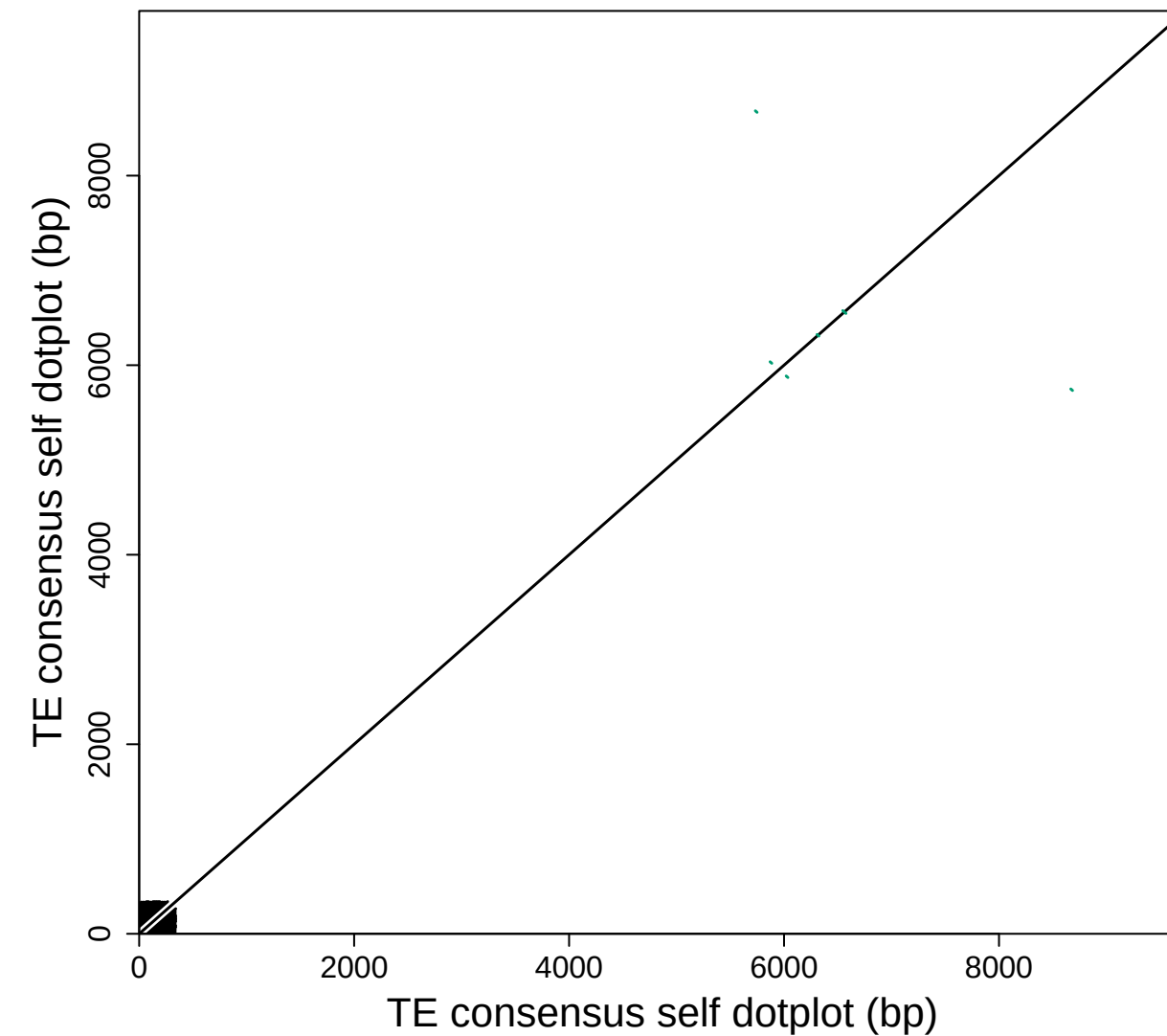
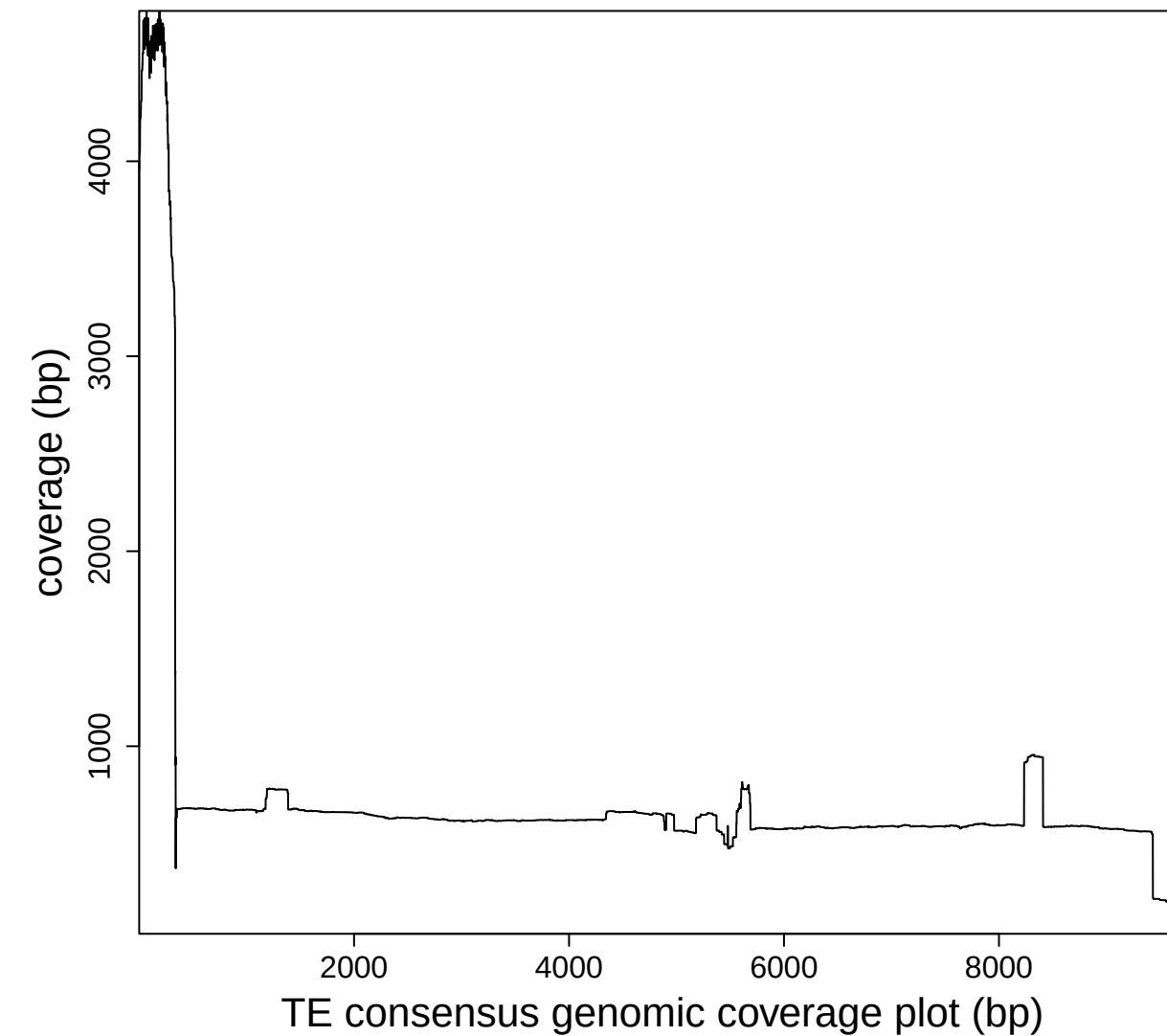
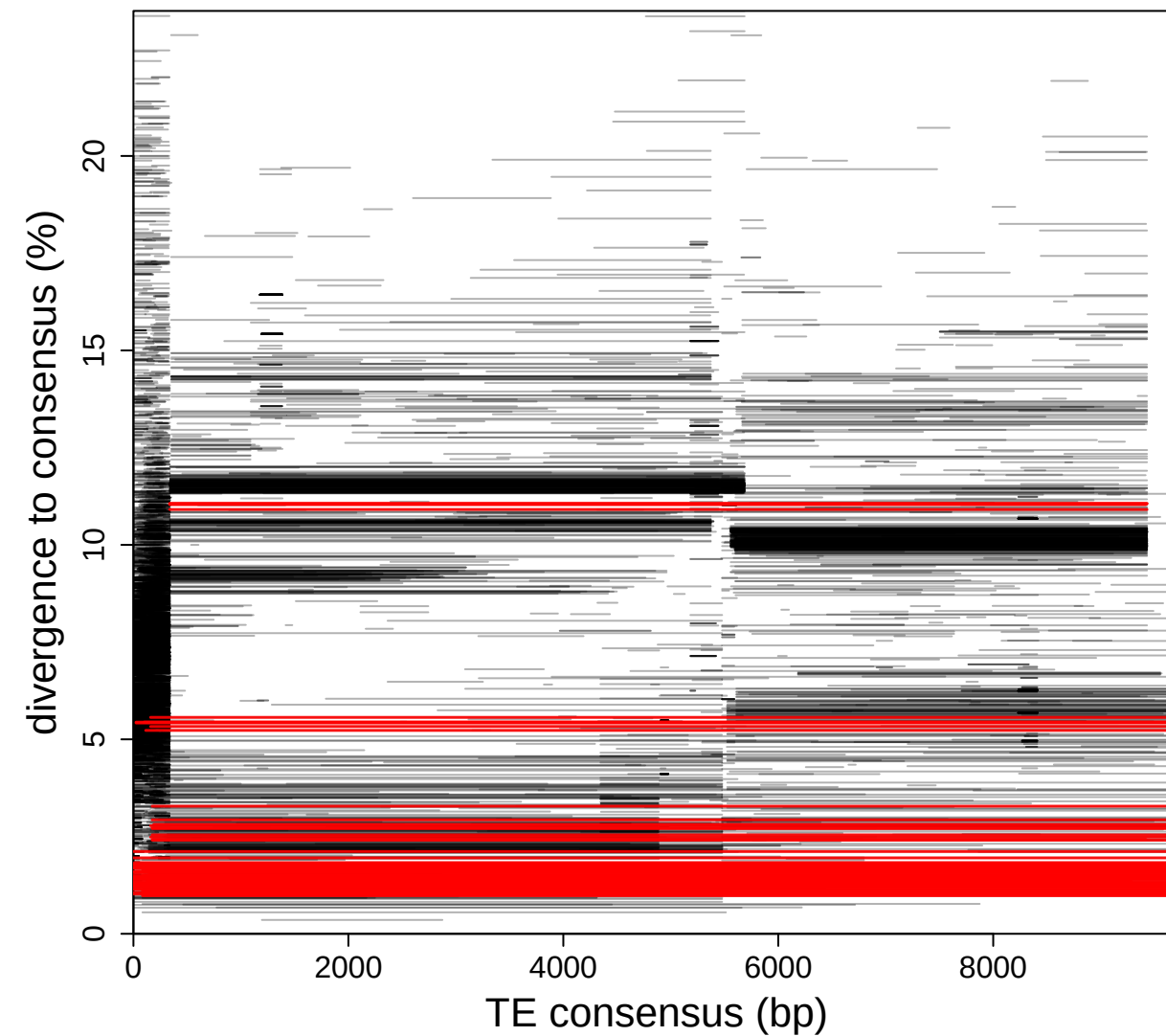
7.fasta.b.bed uf.bed g 4.bed fm 1.bed 0 0 bcln.fa aln.fa cl.fa
size: 17278bp; fragments: 12308; full length: 17 (>=15550.2bp)

After TEtrimmer Extended plot Blue lines are boundaries

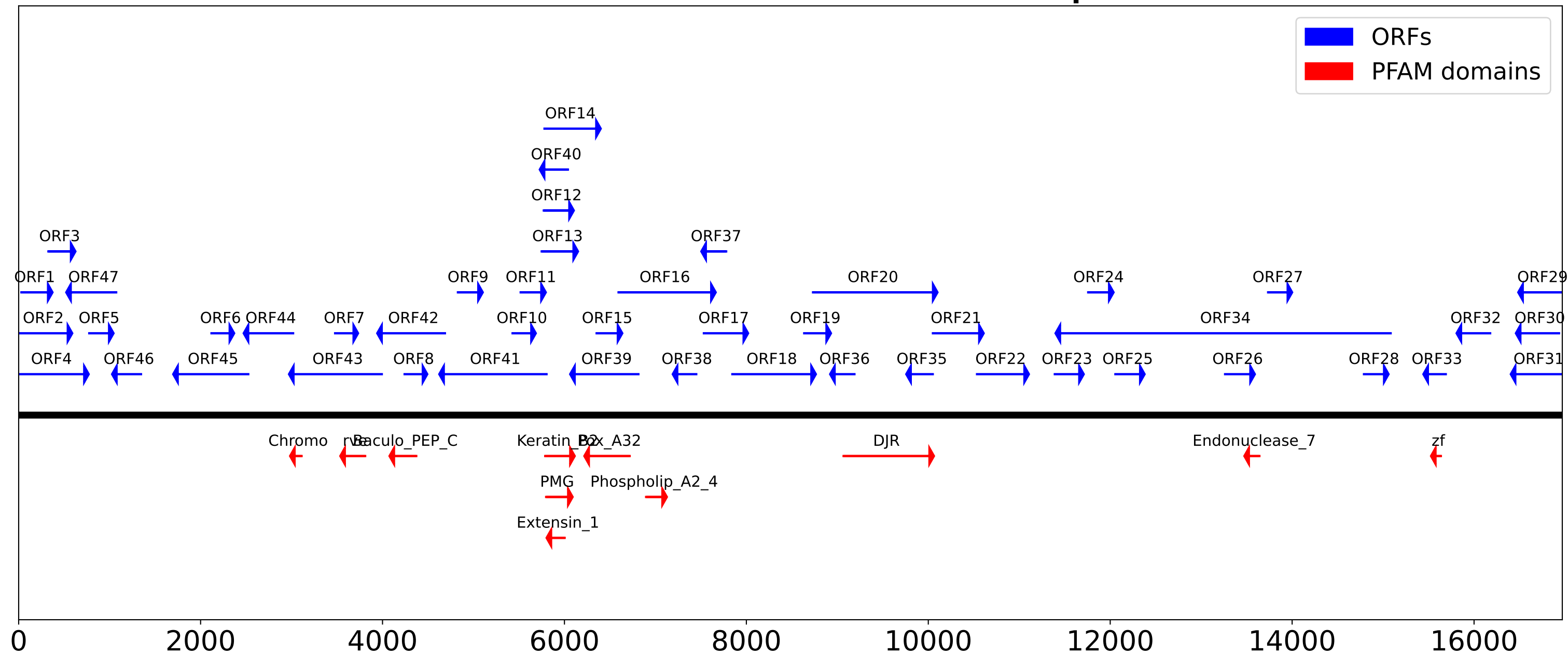


TE: rnd_5_family_757
size: 9738bp; fragments: 10307; full length: 84 (≥ 8764.2 bp)

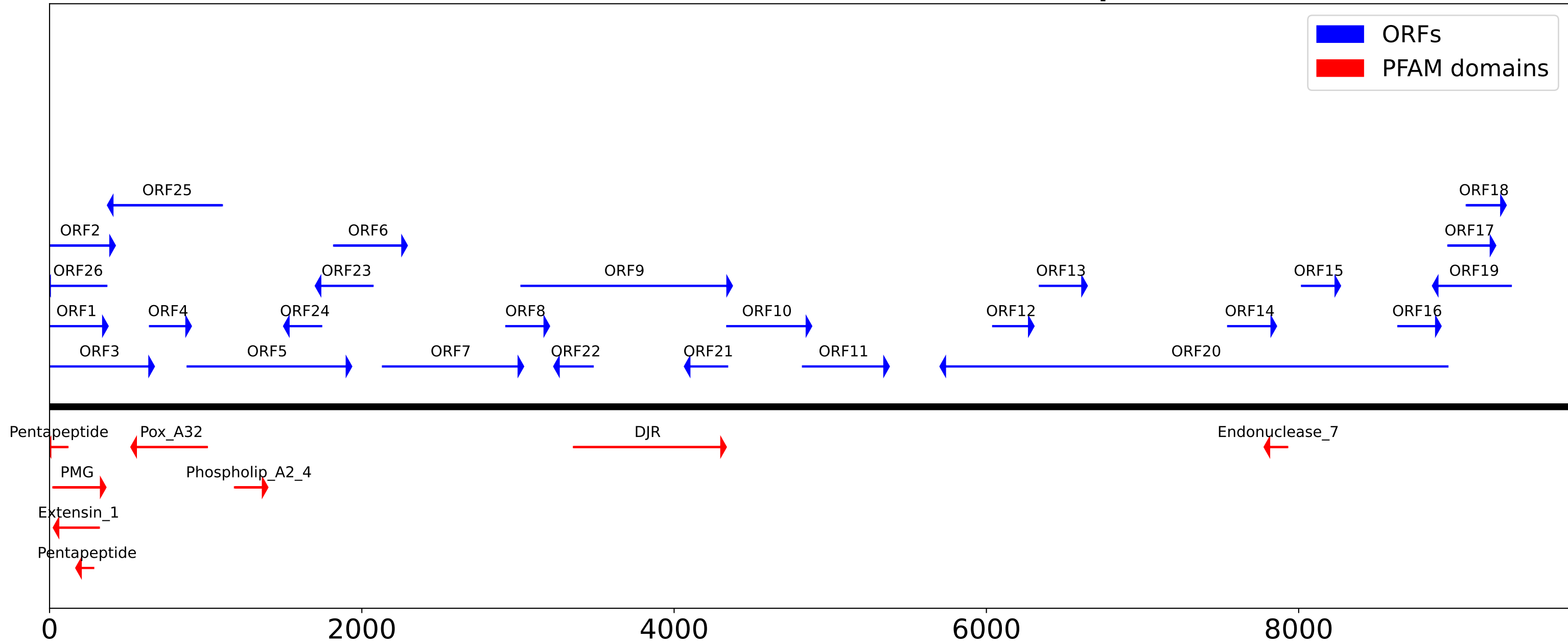
Before TEtrimmer 9738 bp



After TETrimmer ORF and PFAM domain plot



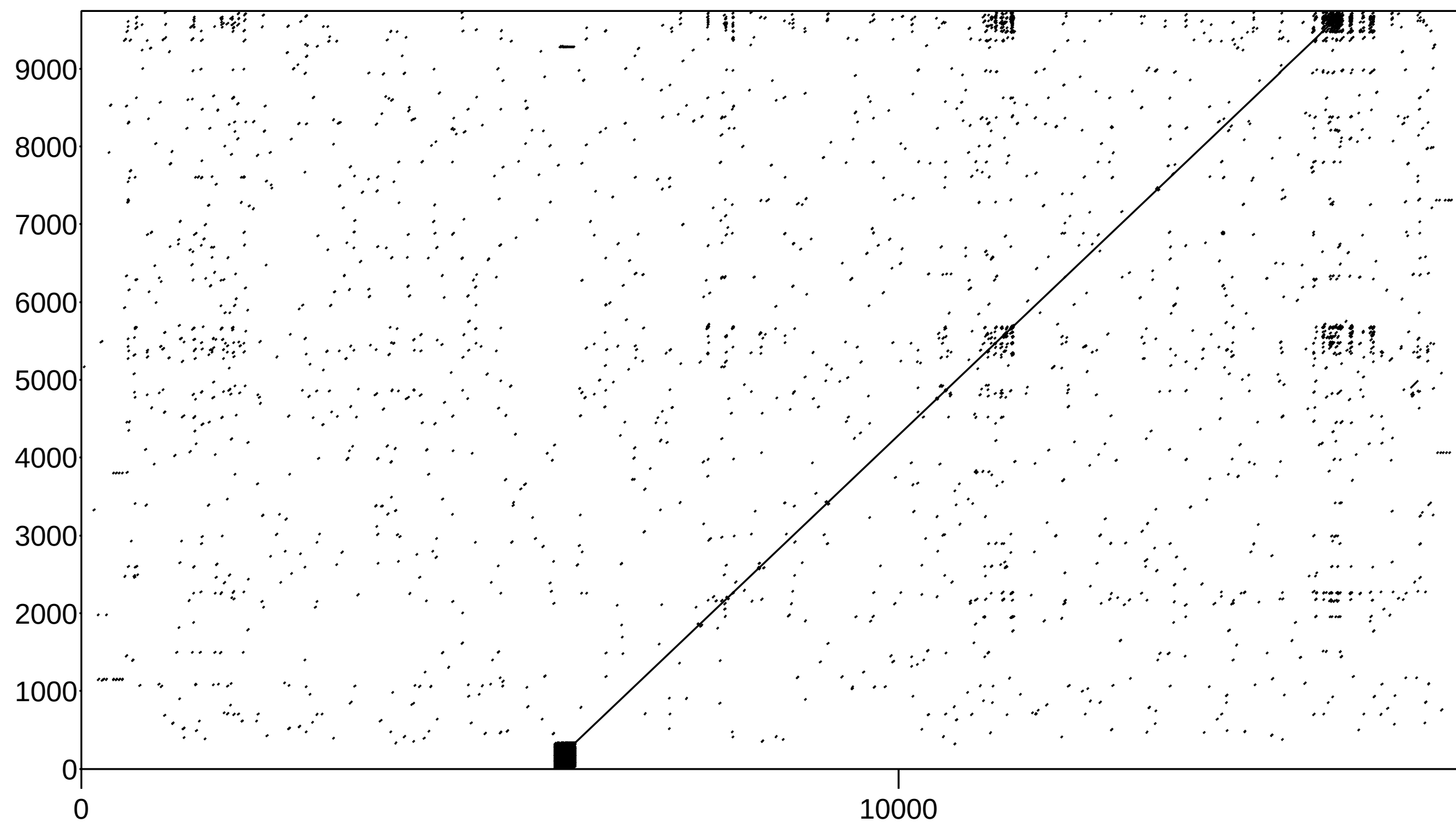
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)