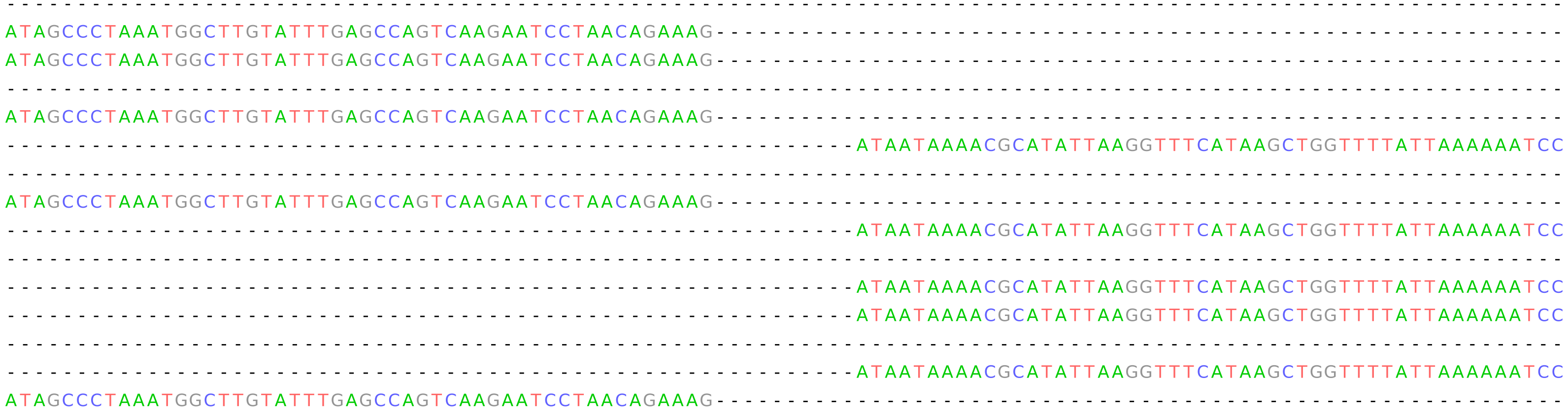
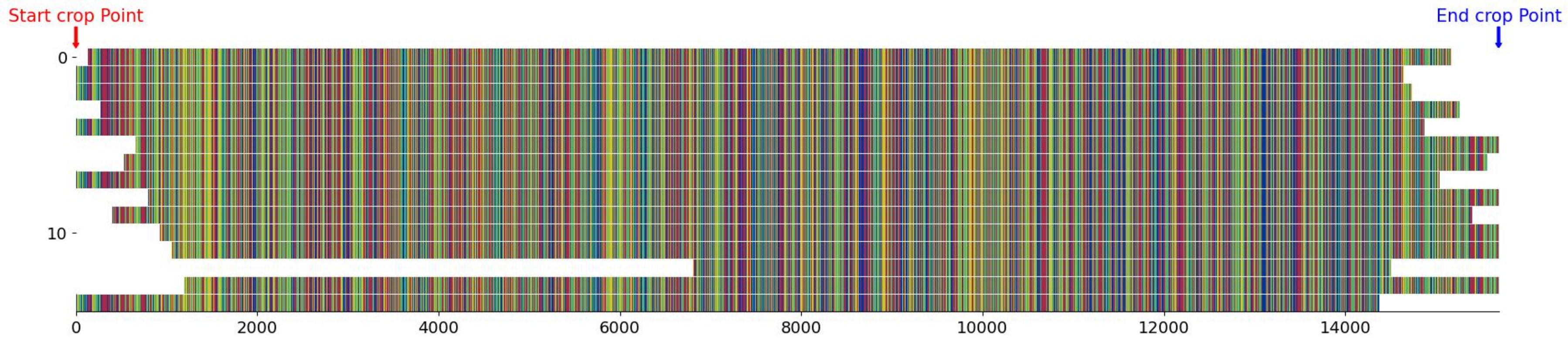


Start crop Point

End crop Point

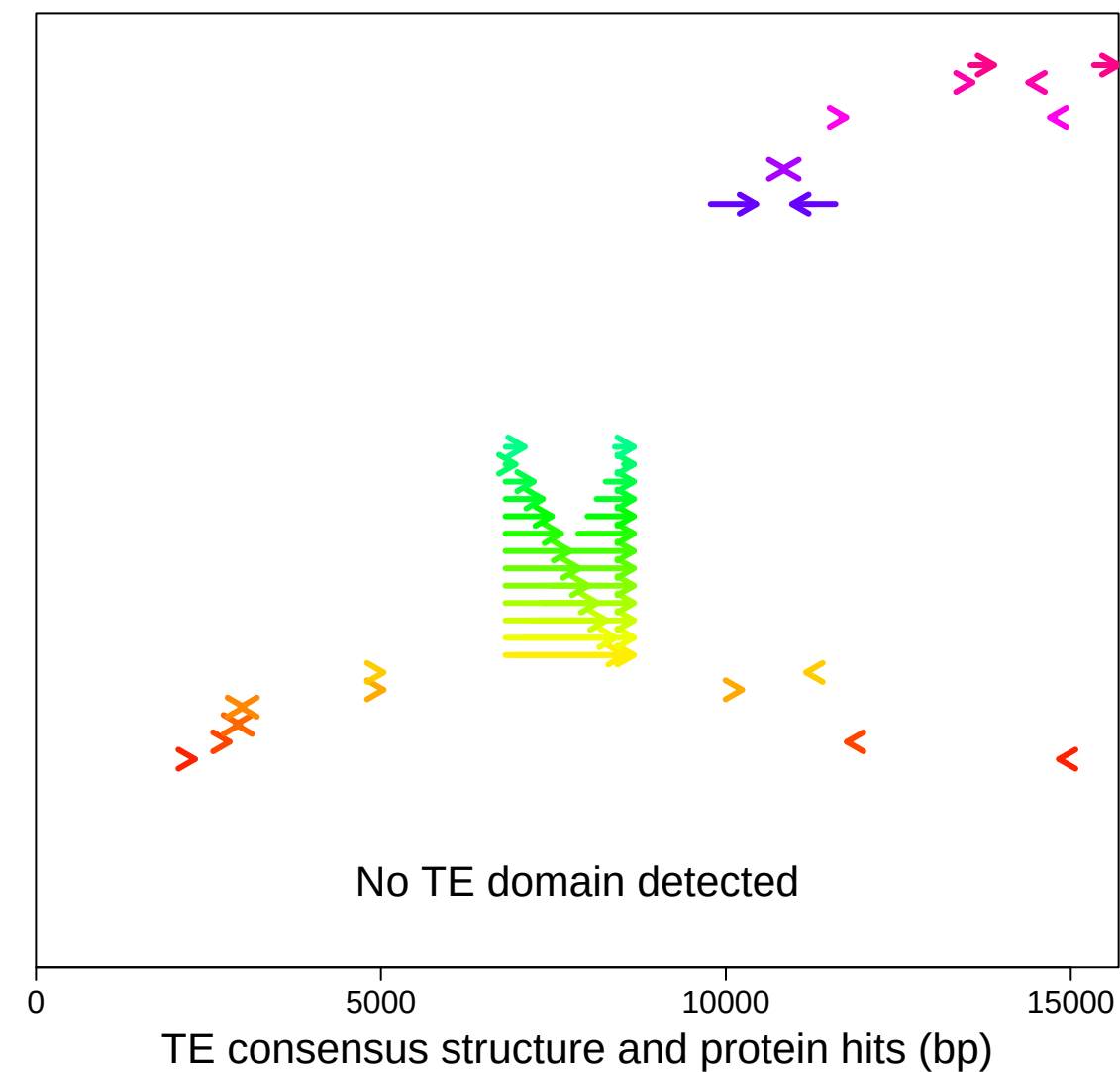
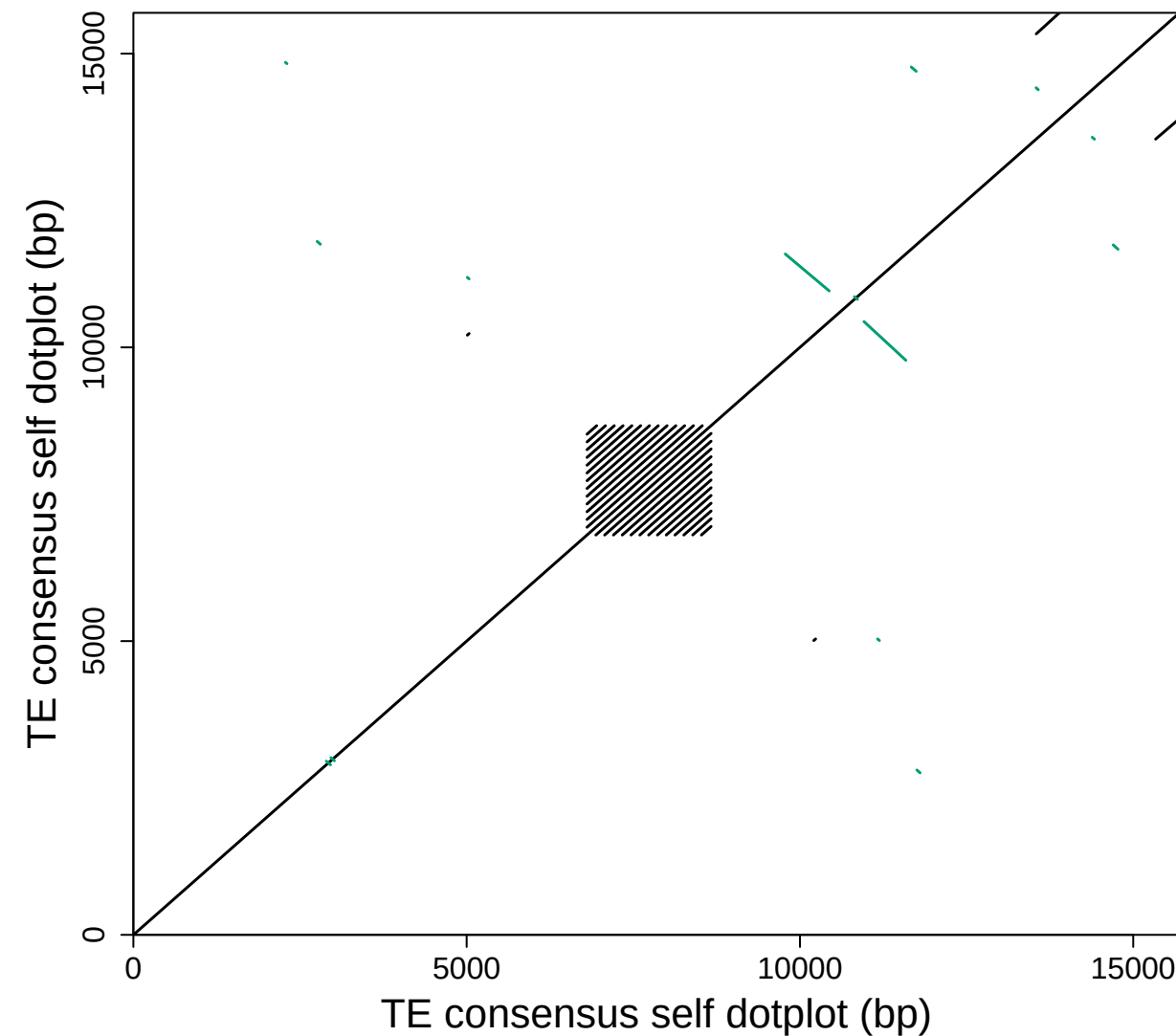
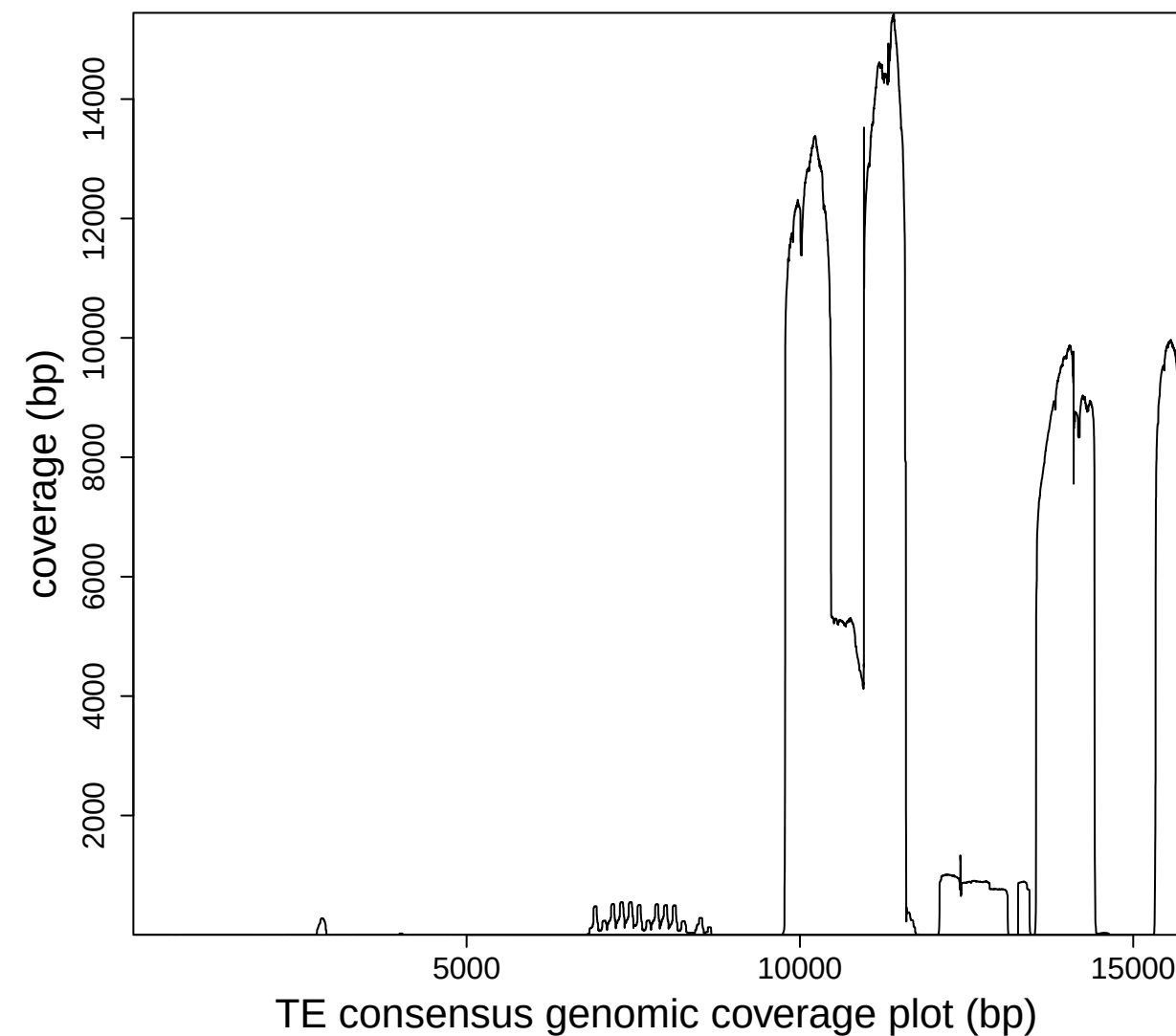
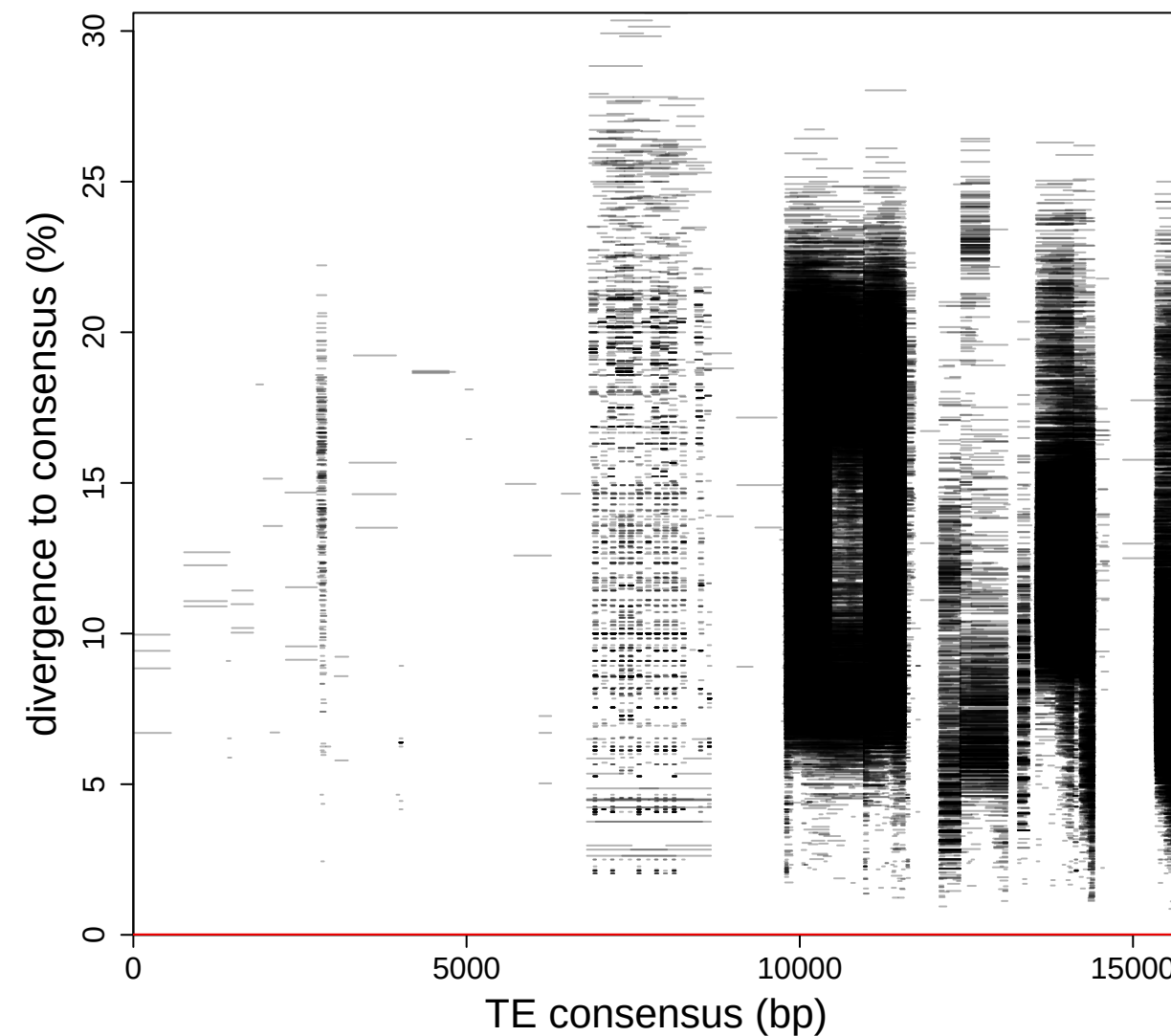
MSA length = 15694





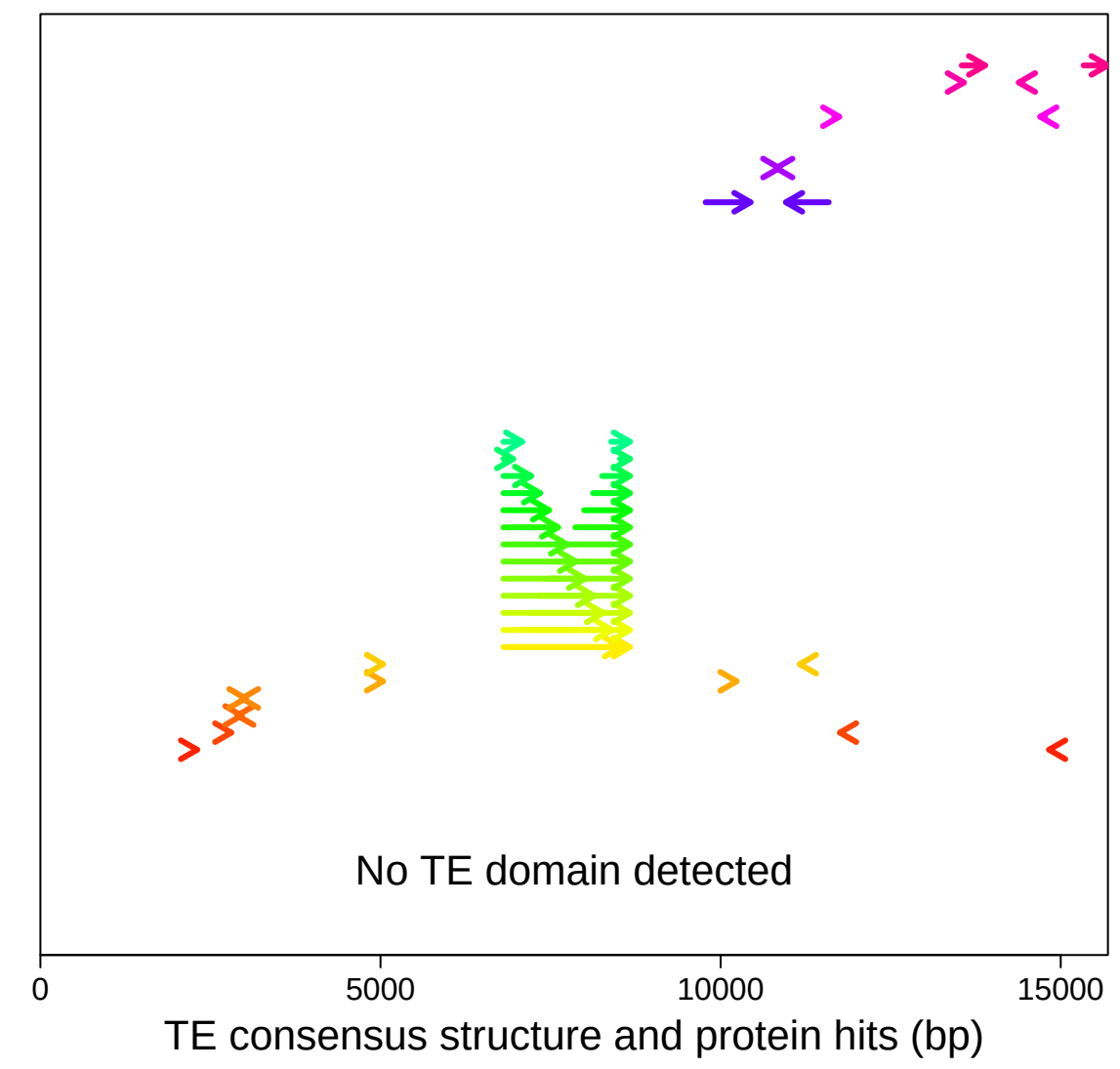
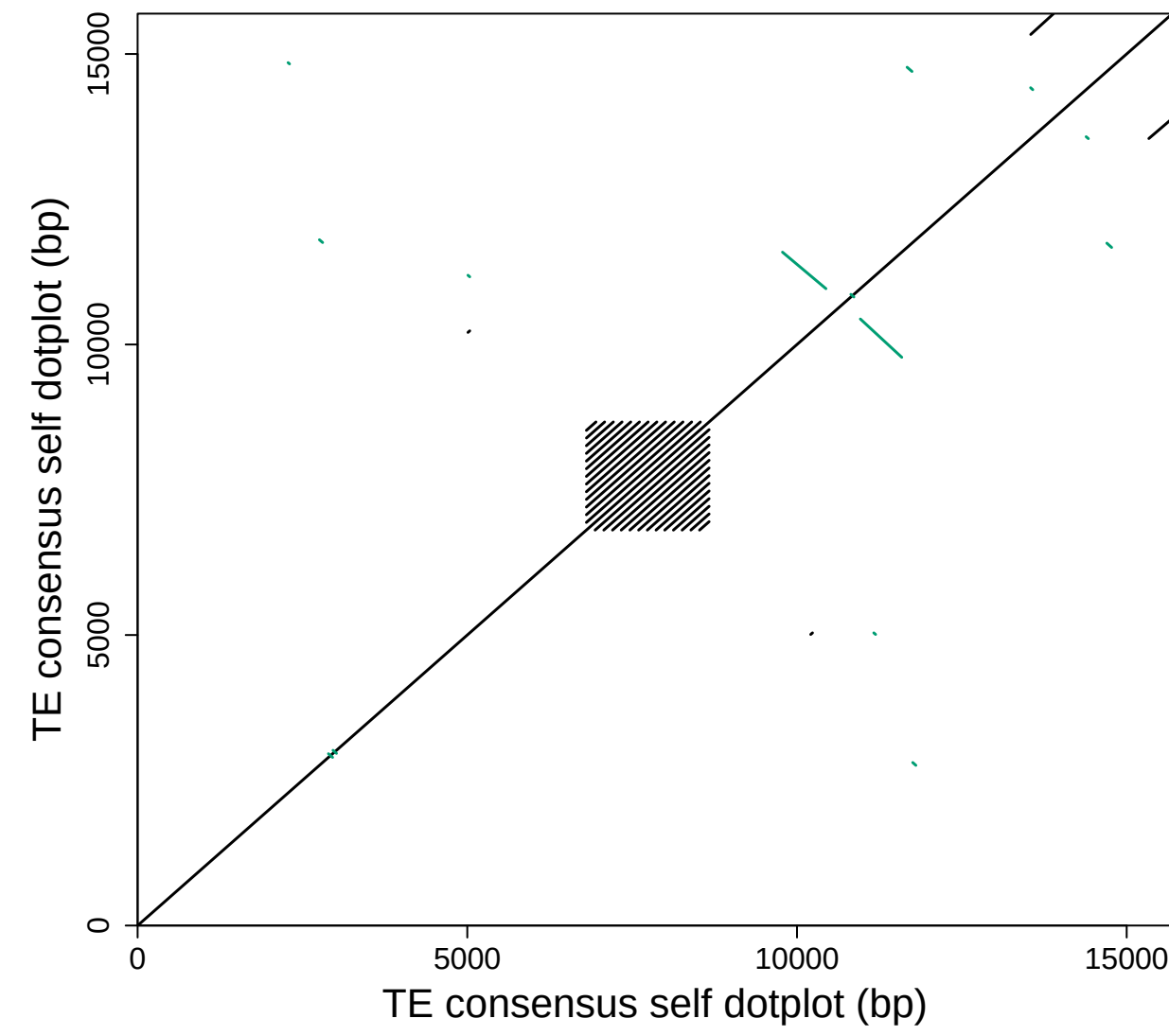
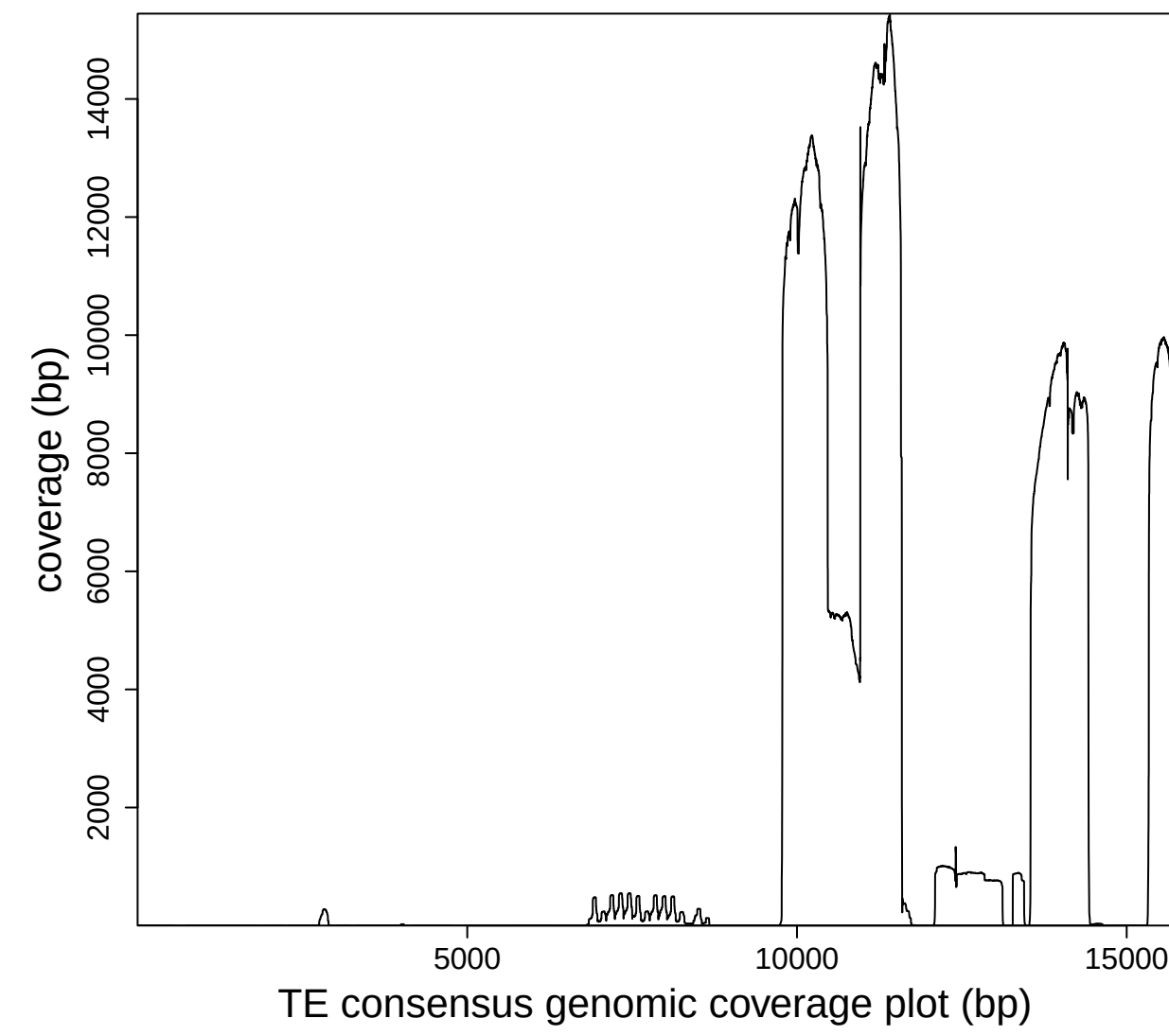
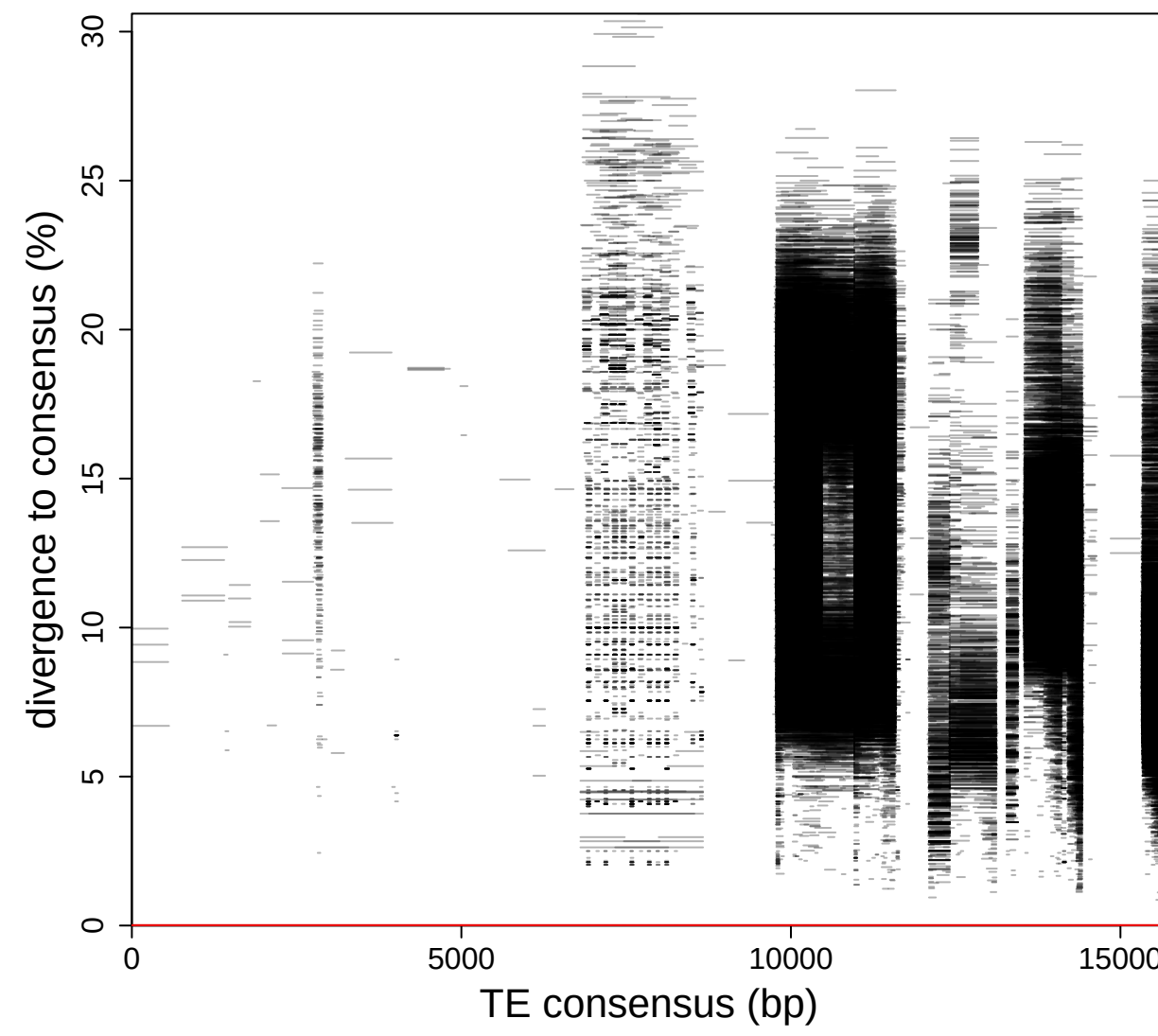
l.b.bed uf.bed g_1.bed fm_1.bed 0 0 bcIn.fa aln.fa cl.fa gs.fa
size: 15694bp; fragments: 72495; full length: 1 (>=14124.6bp)

After TEtrimmer 15694 bp

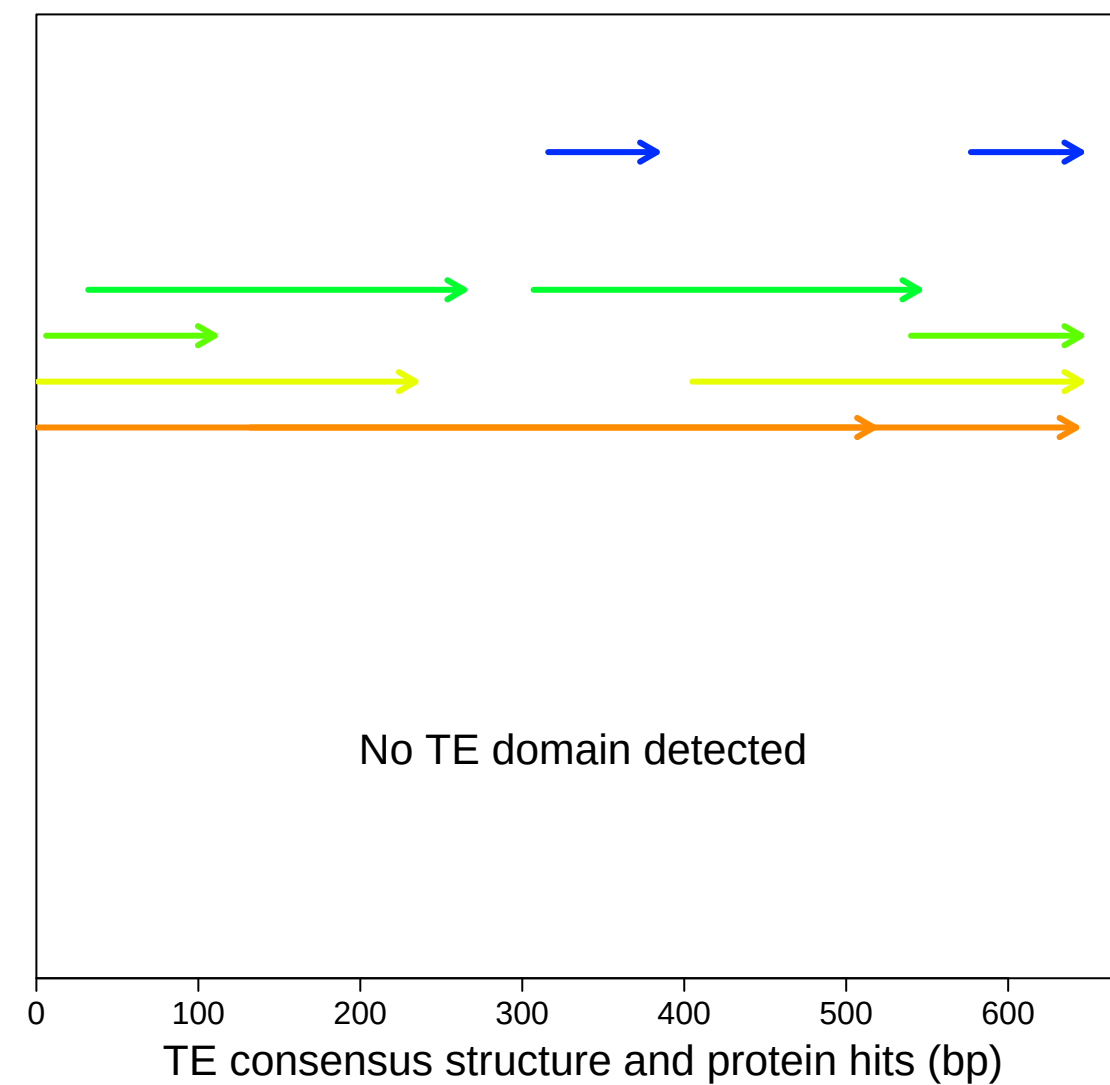
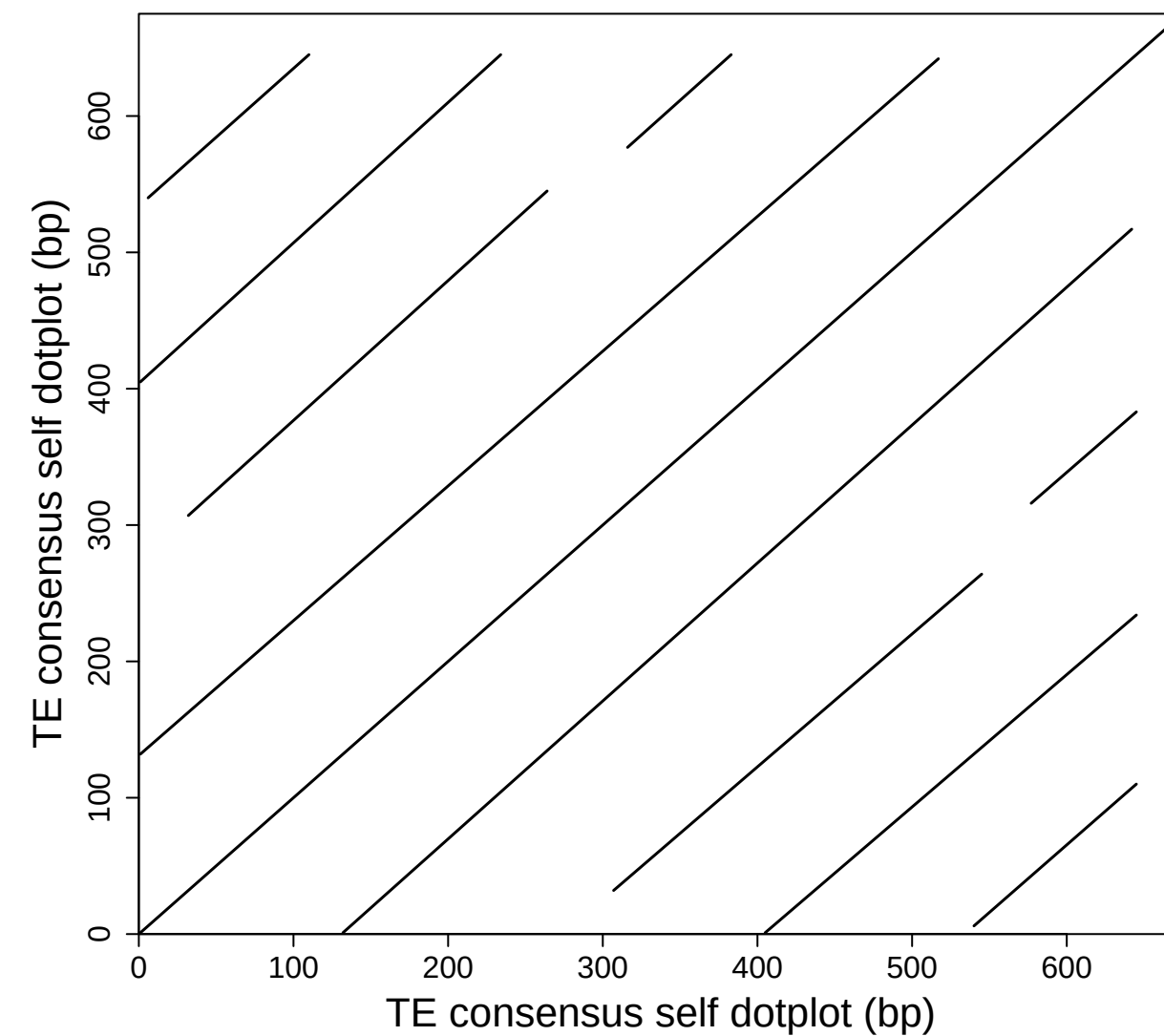
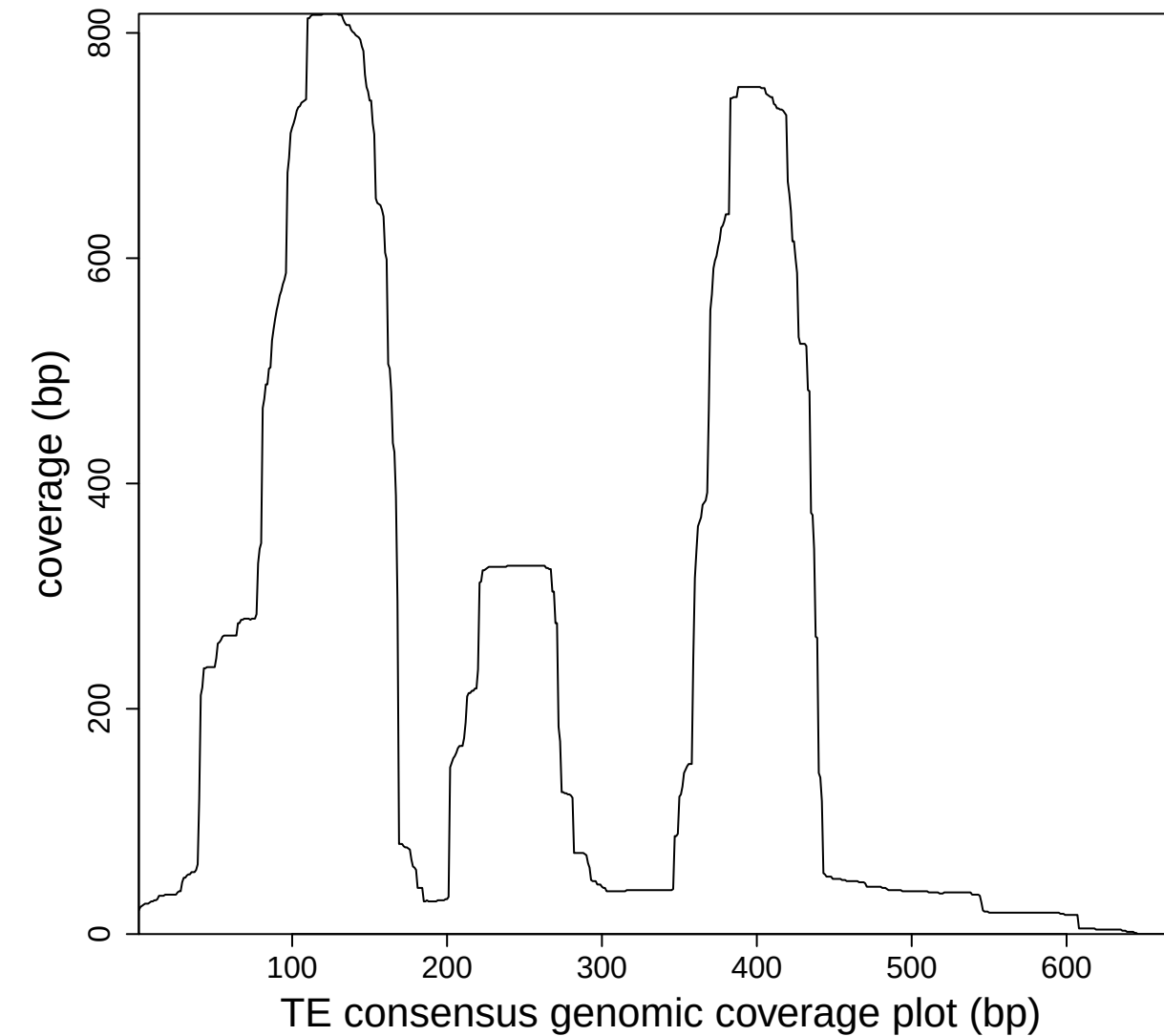
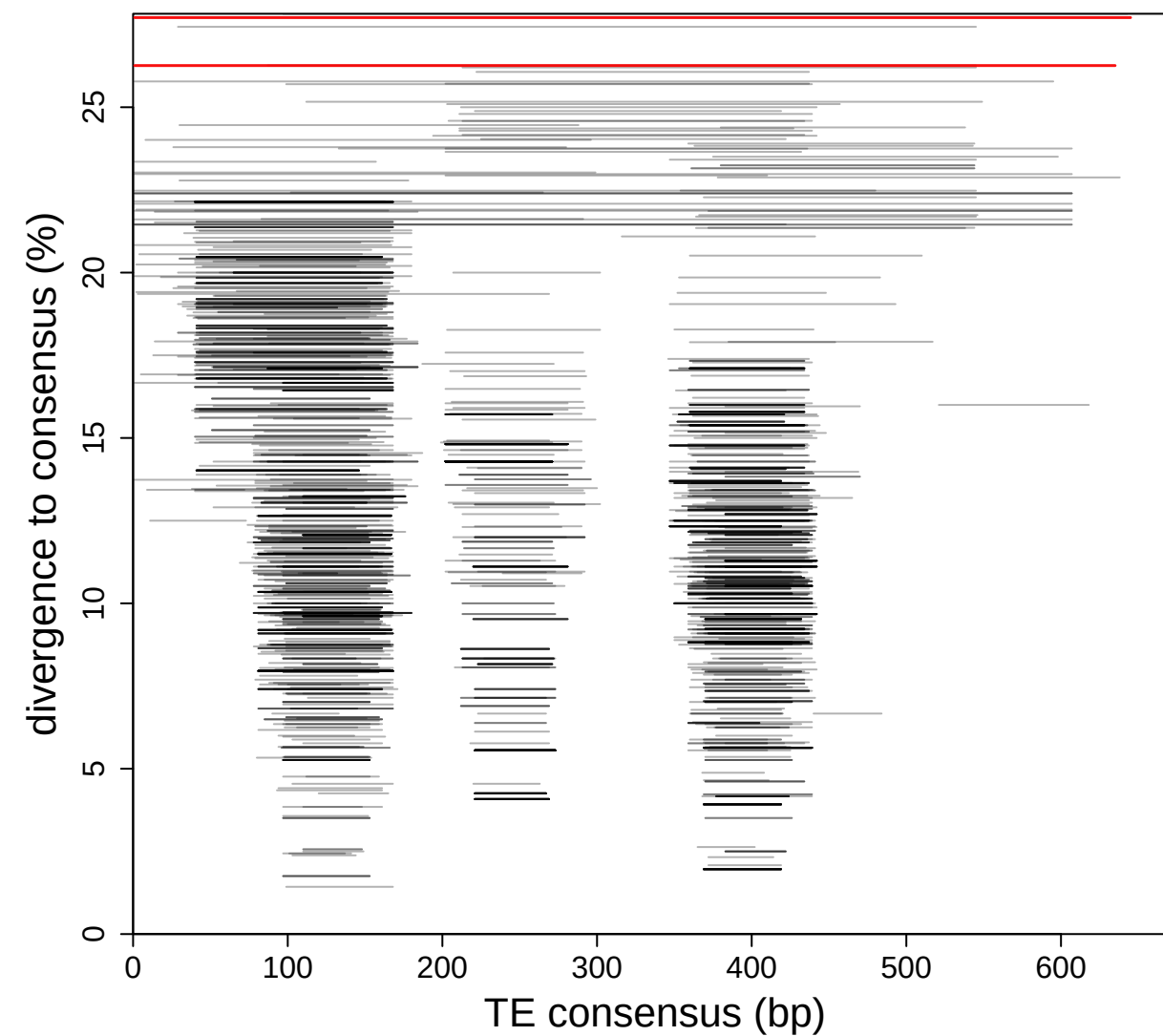


8.fasta.b.bed uf.bed g 1.bed fm 1.bed 0 0 bcln.fa aln.fa cl.fa
size: 15694bp; fragments: 72495; full length: 1 (>=14124.6bp)

After TEtrimmer Extended plot Blue lines are boundaries

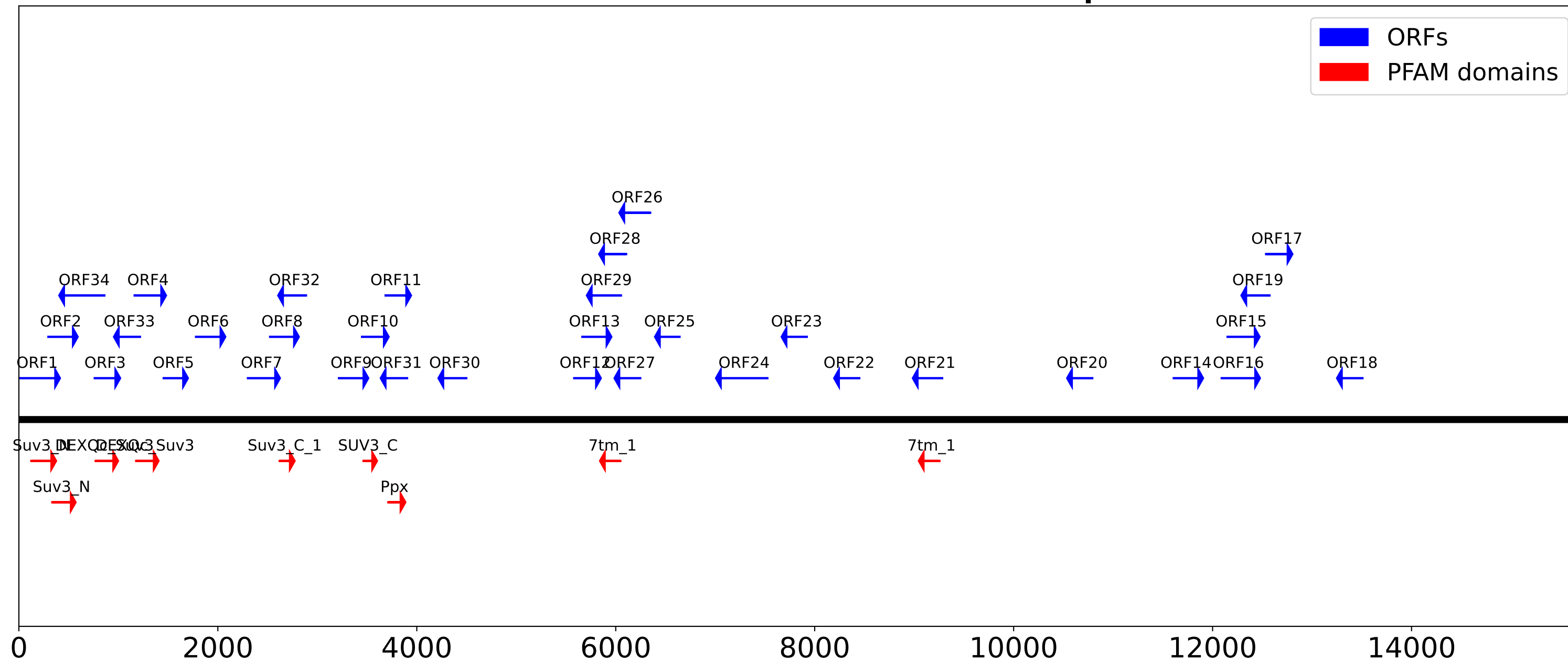


TE: rnd_4_family_318
size: 675bp; fragments: 1835; full length: 2 (≥ 607.5 bp)

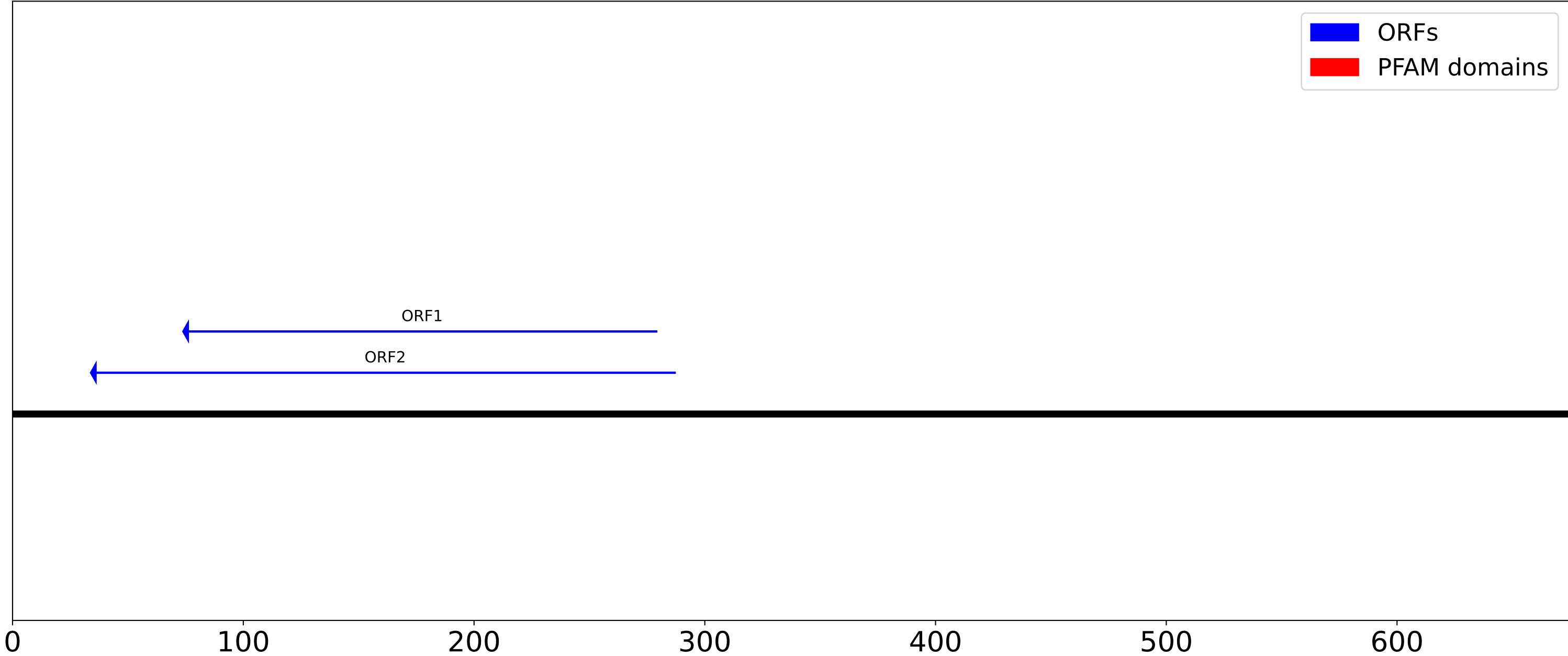


Before TEtrimmer 675 bp

After TETrimmer ORF and PFAM domain plot



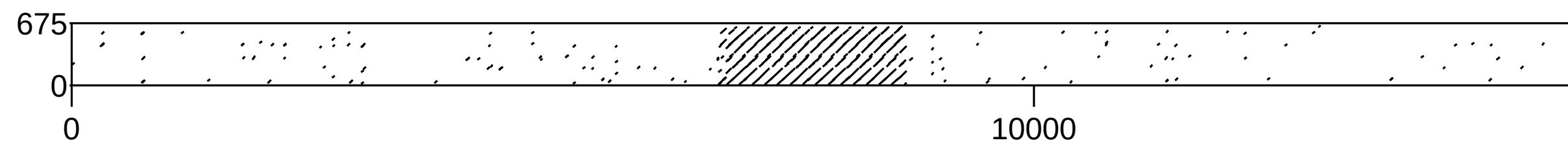
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

nsus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)