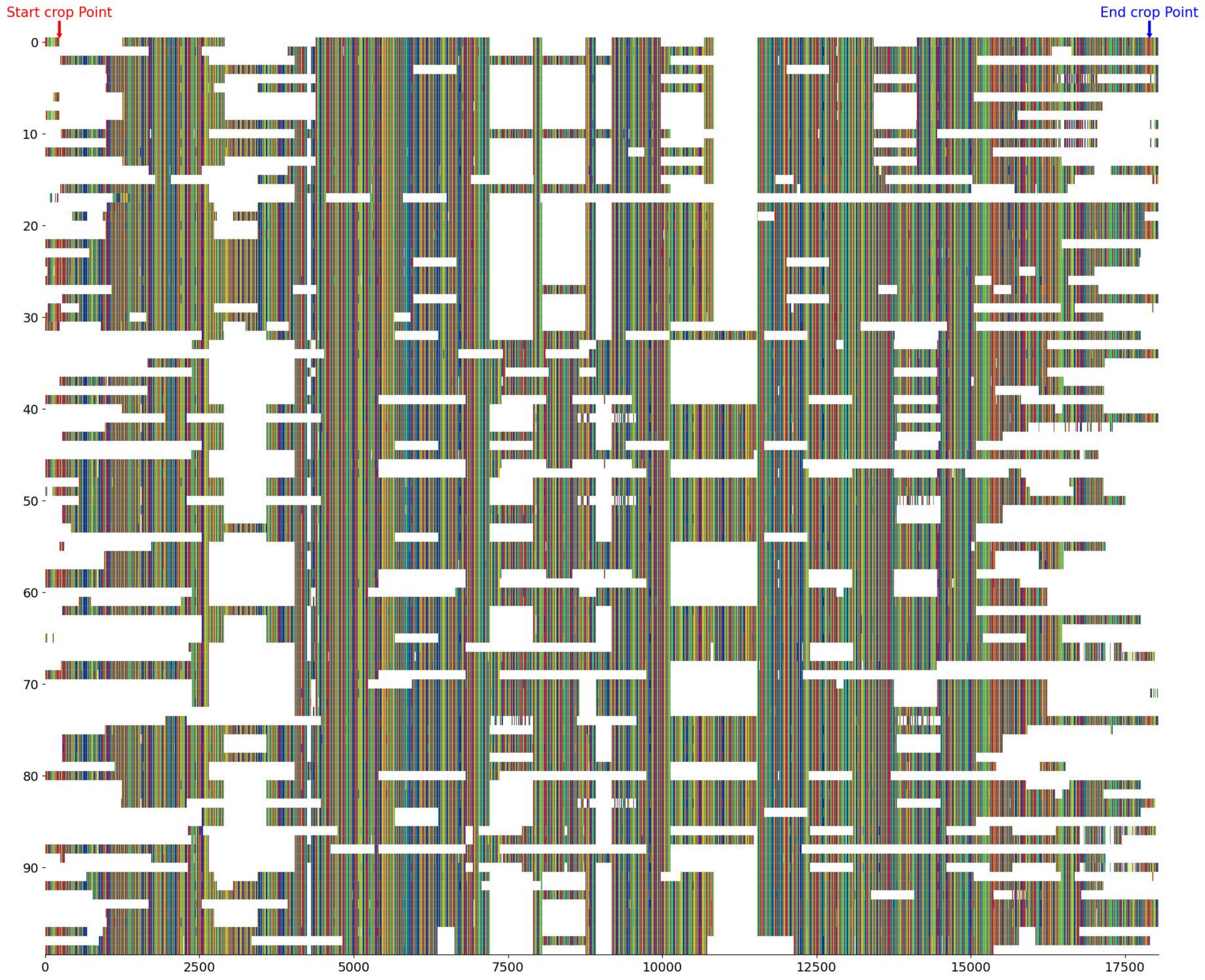
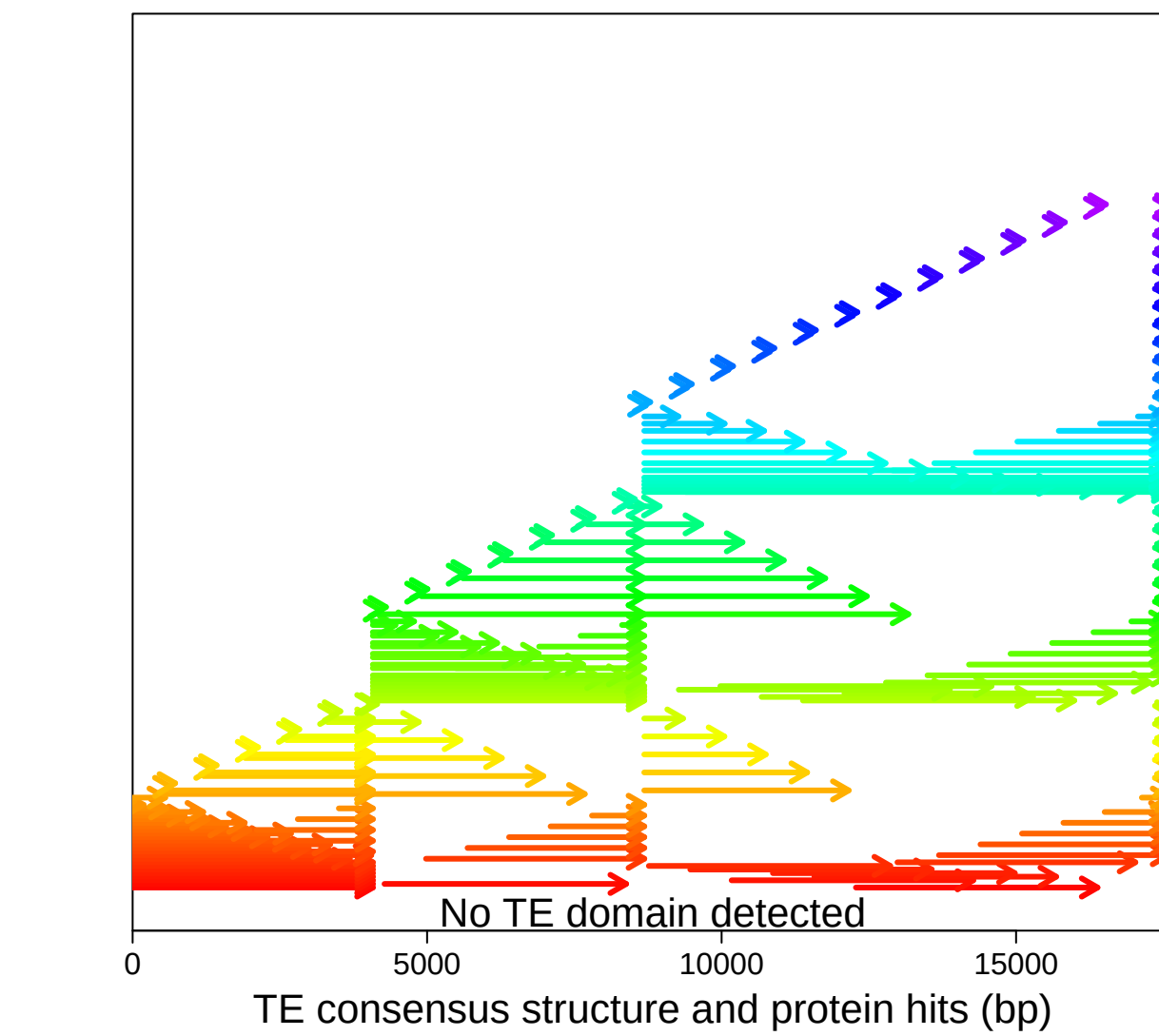
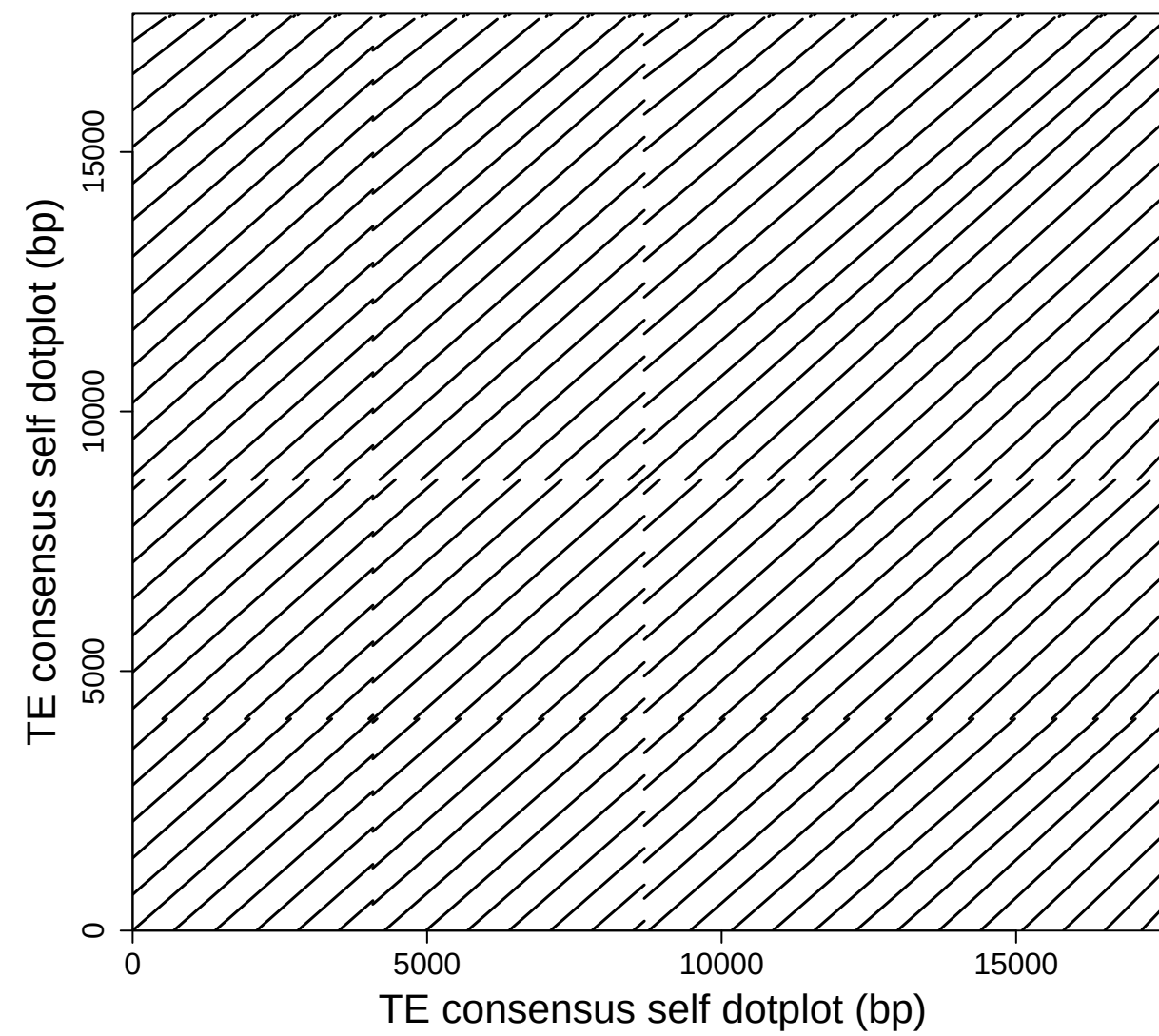
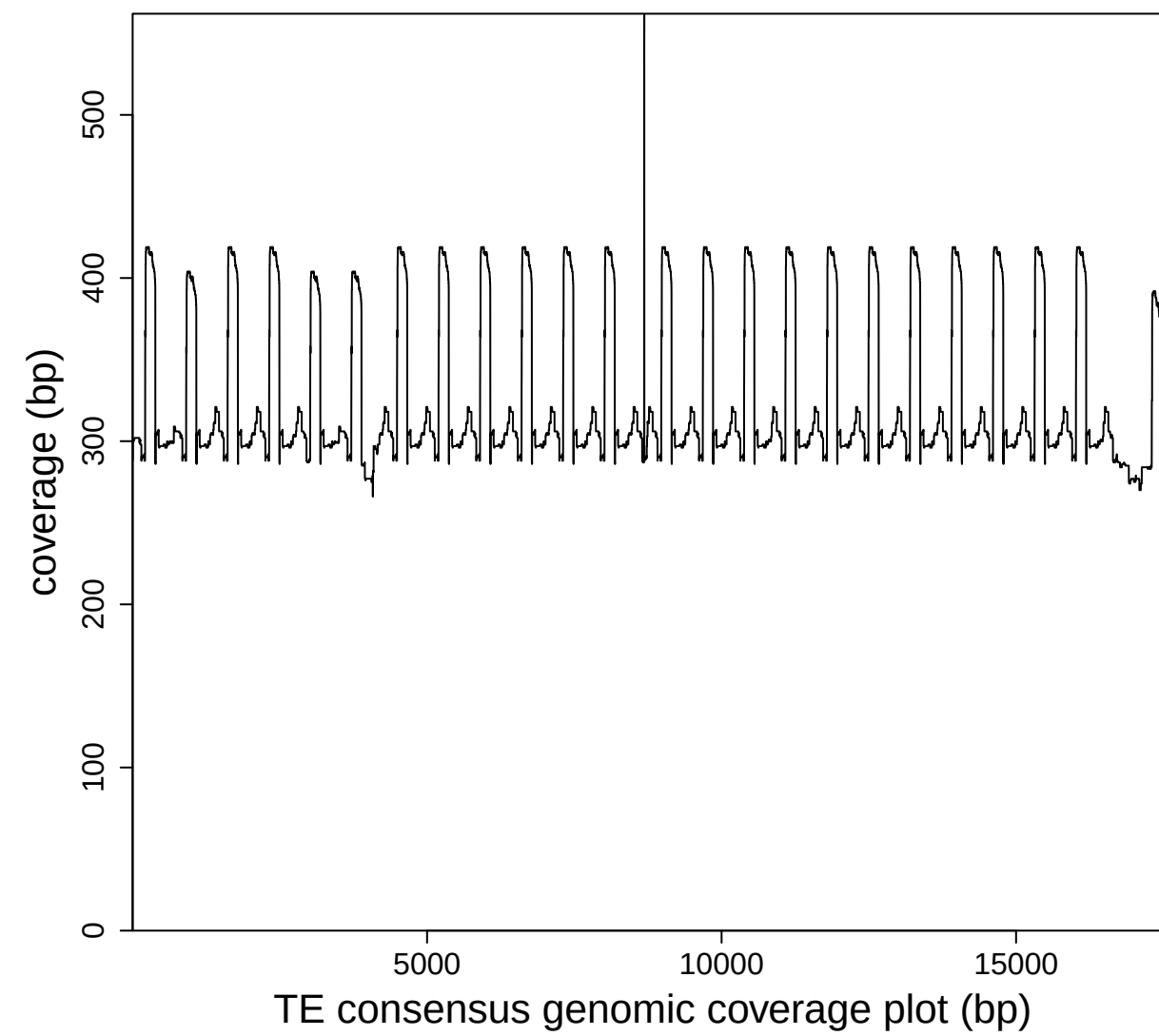
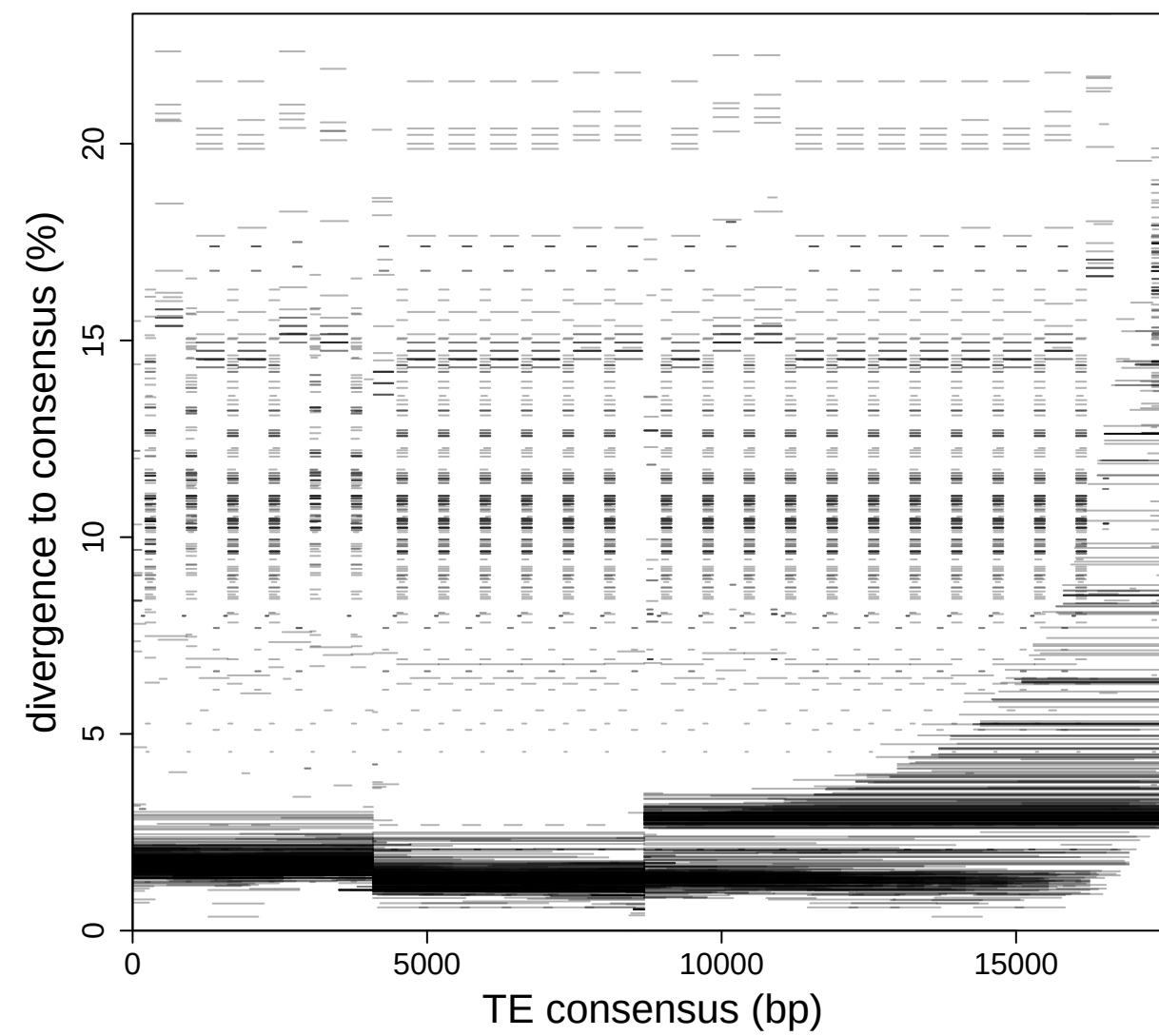






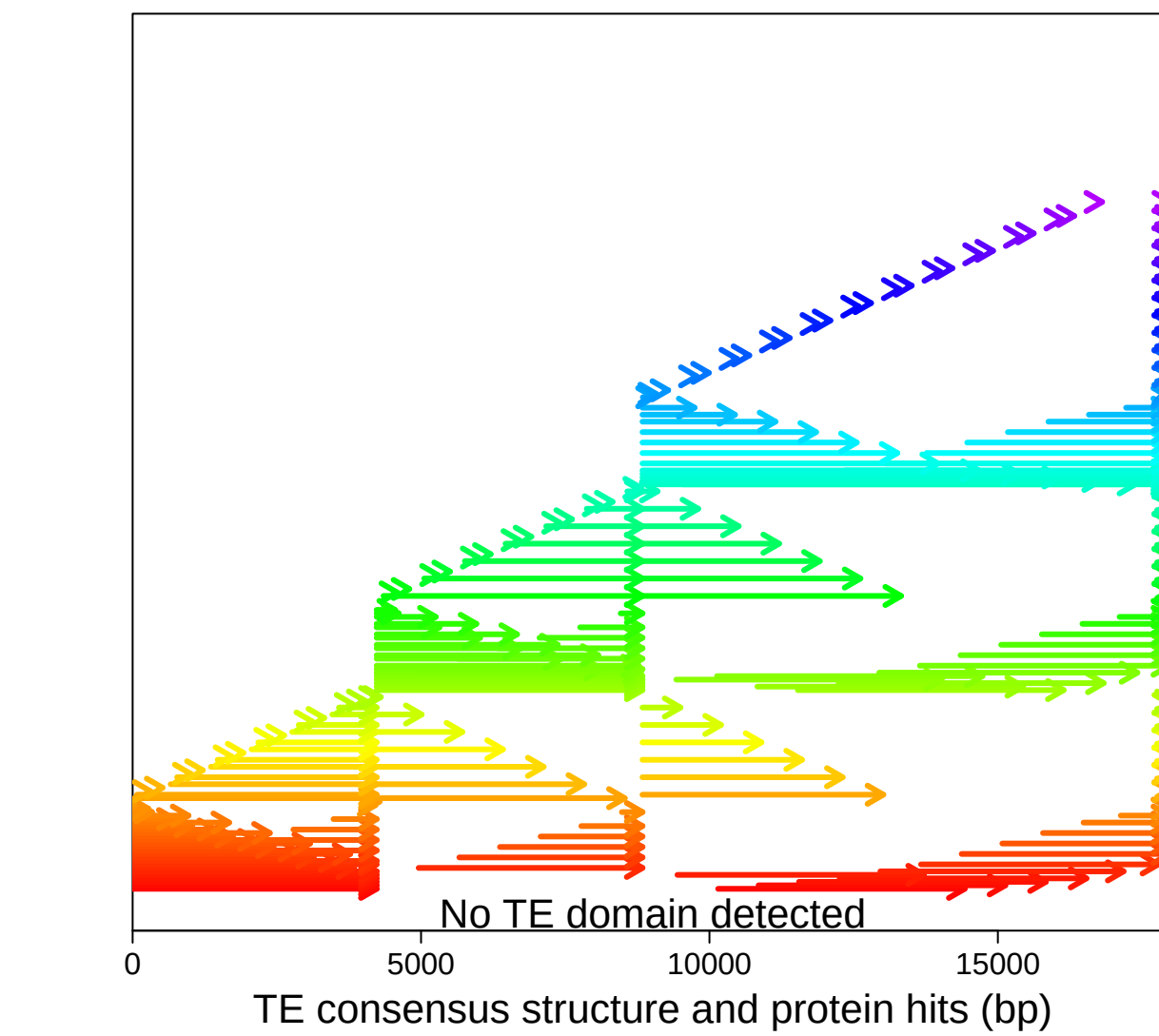
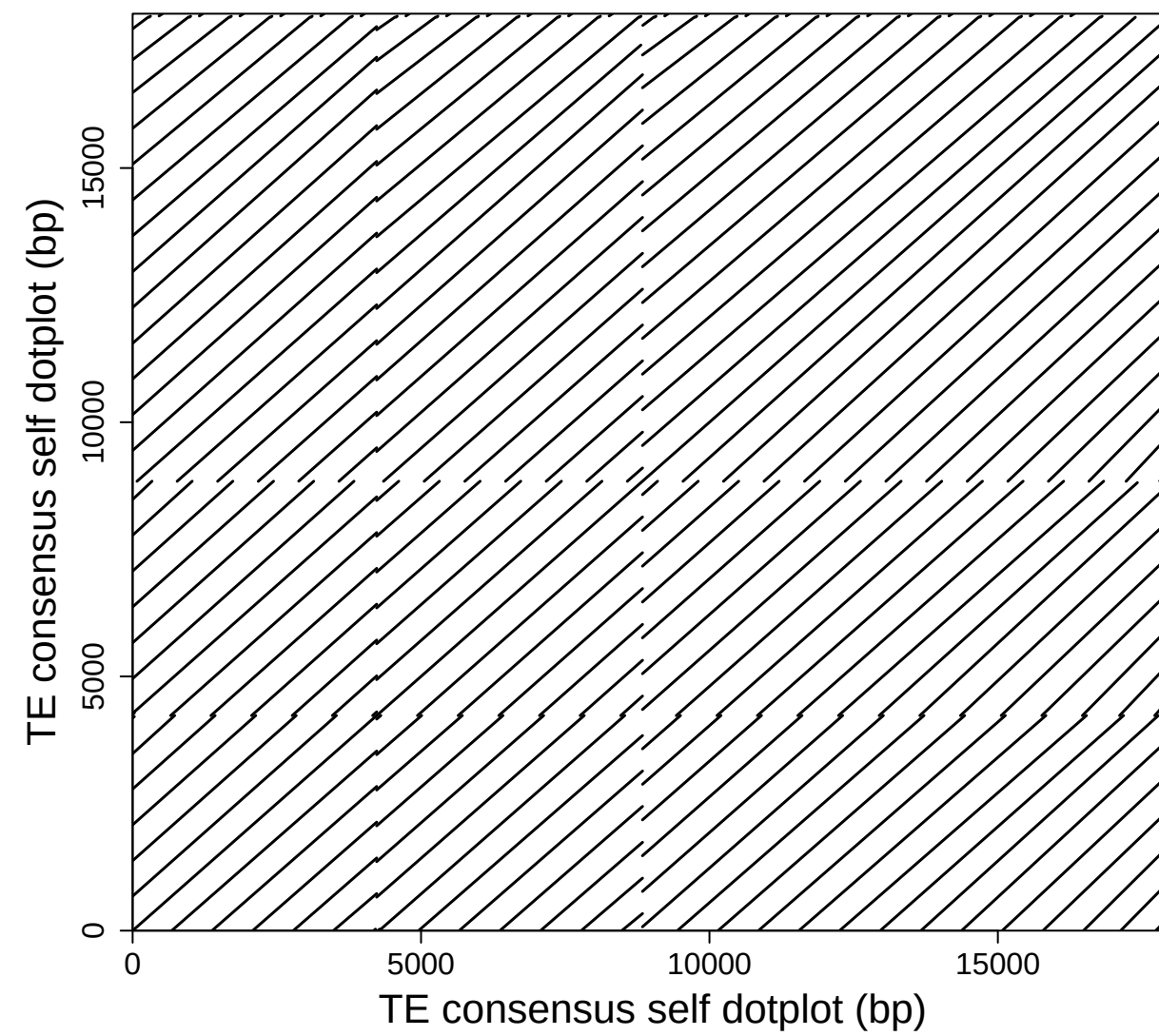
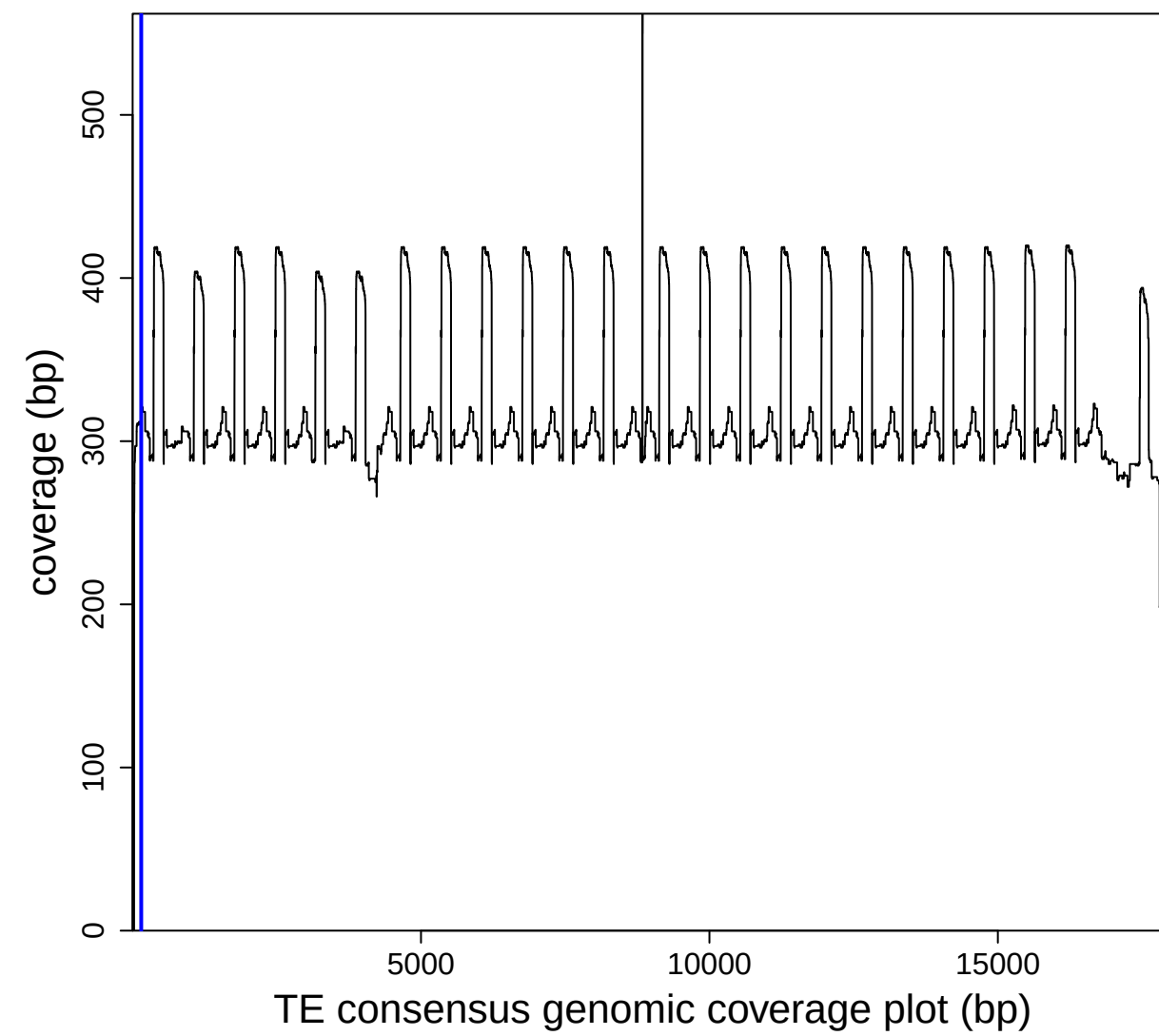
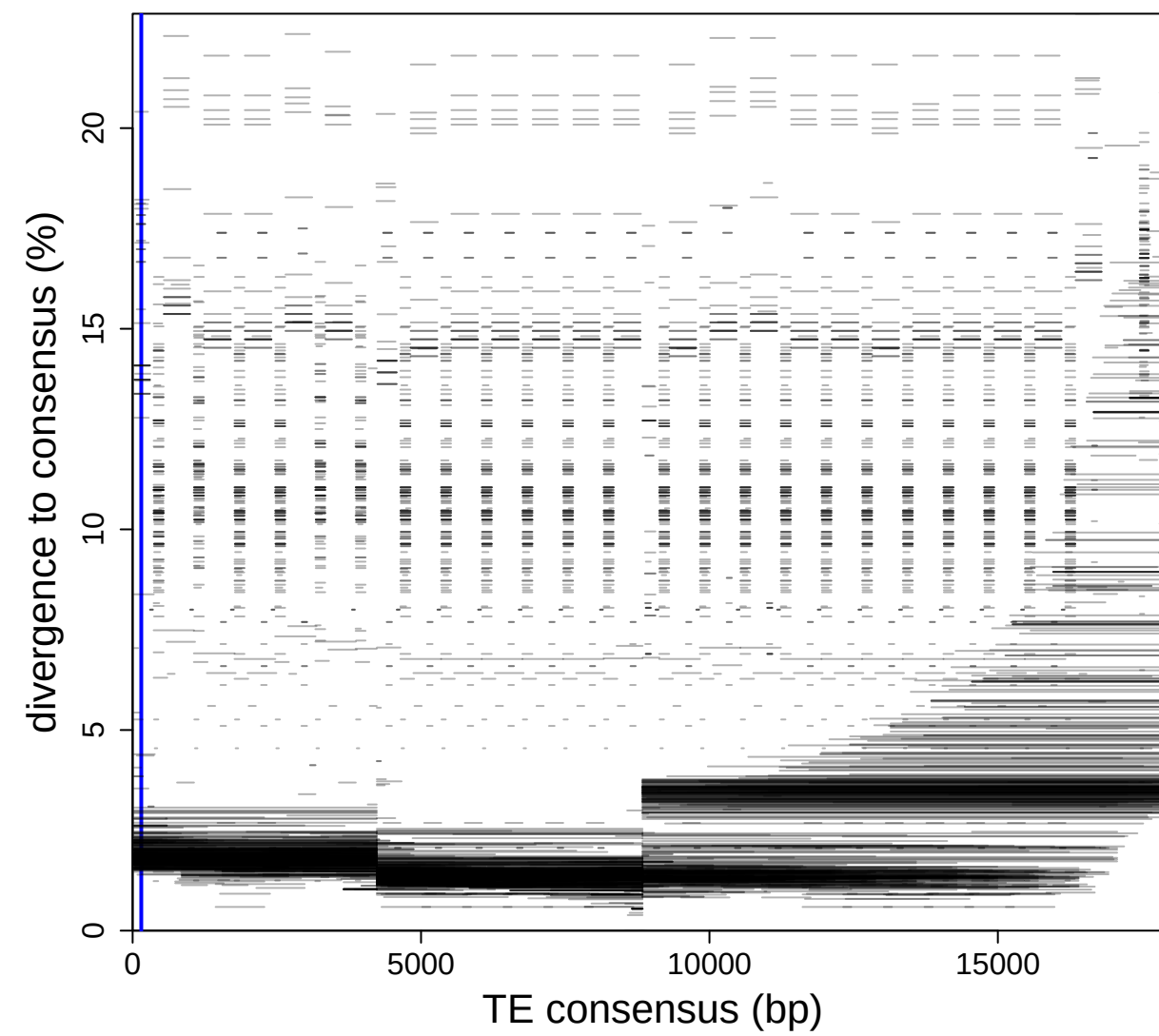
bed_uf.bed g_1.bed fm_1.bed 0 0 bcln.fa aln.fa cl.fa gs.fa c
size: 17663bp; fragments: 5675; full length: 0 (>=15896.7bp)

After TEtrimmer 17663 bp

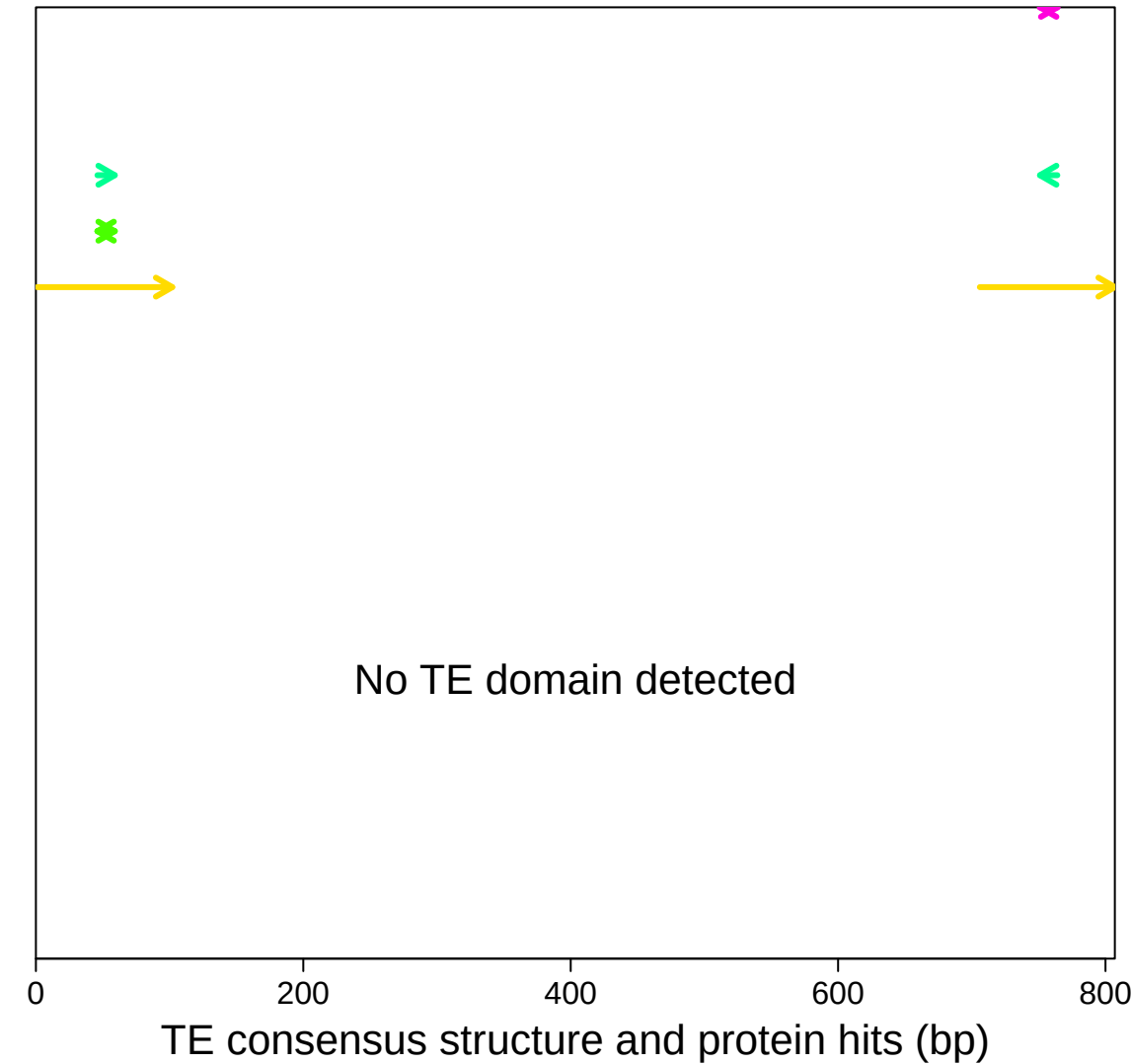
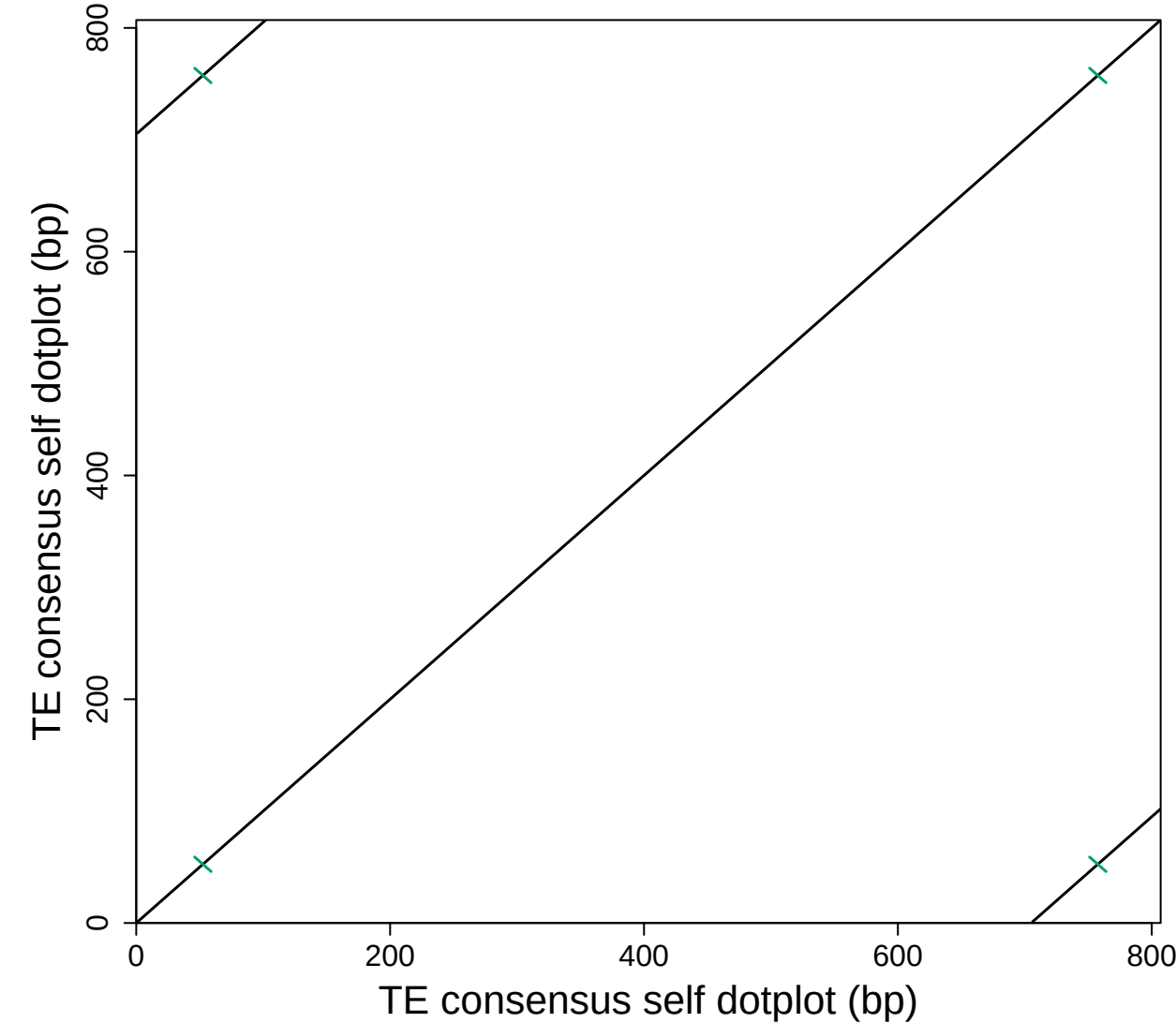
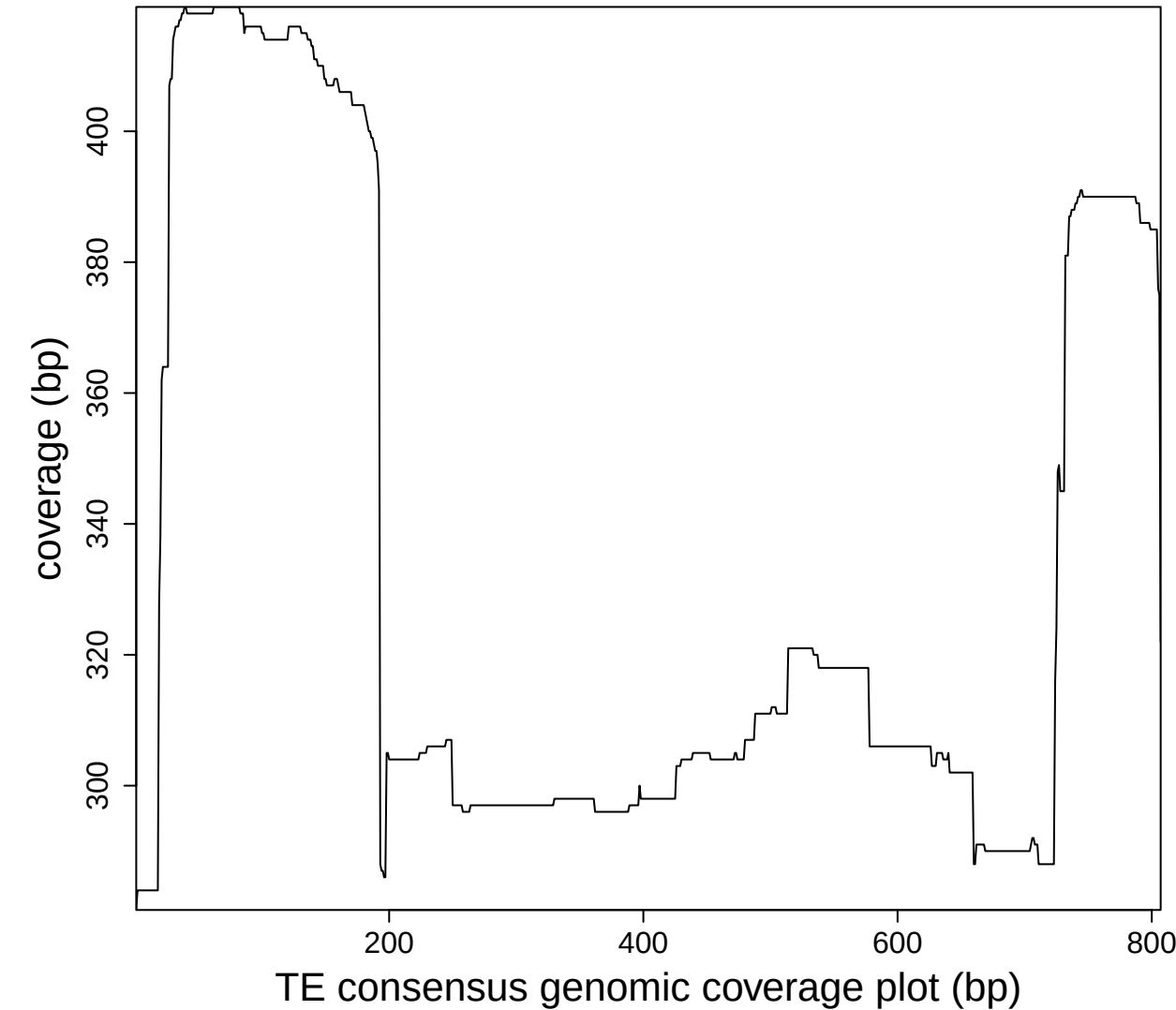
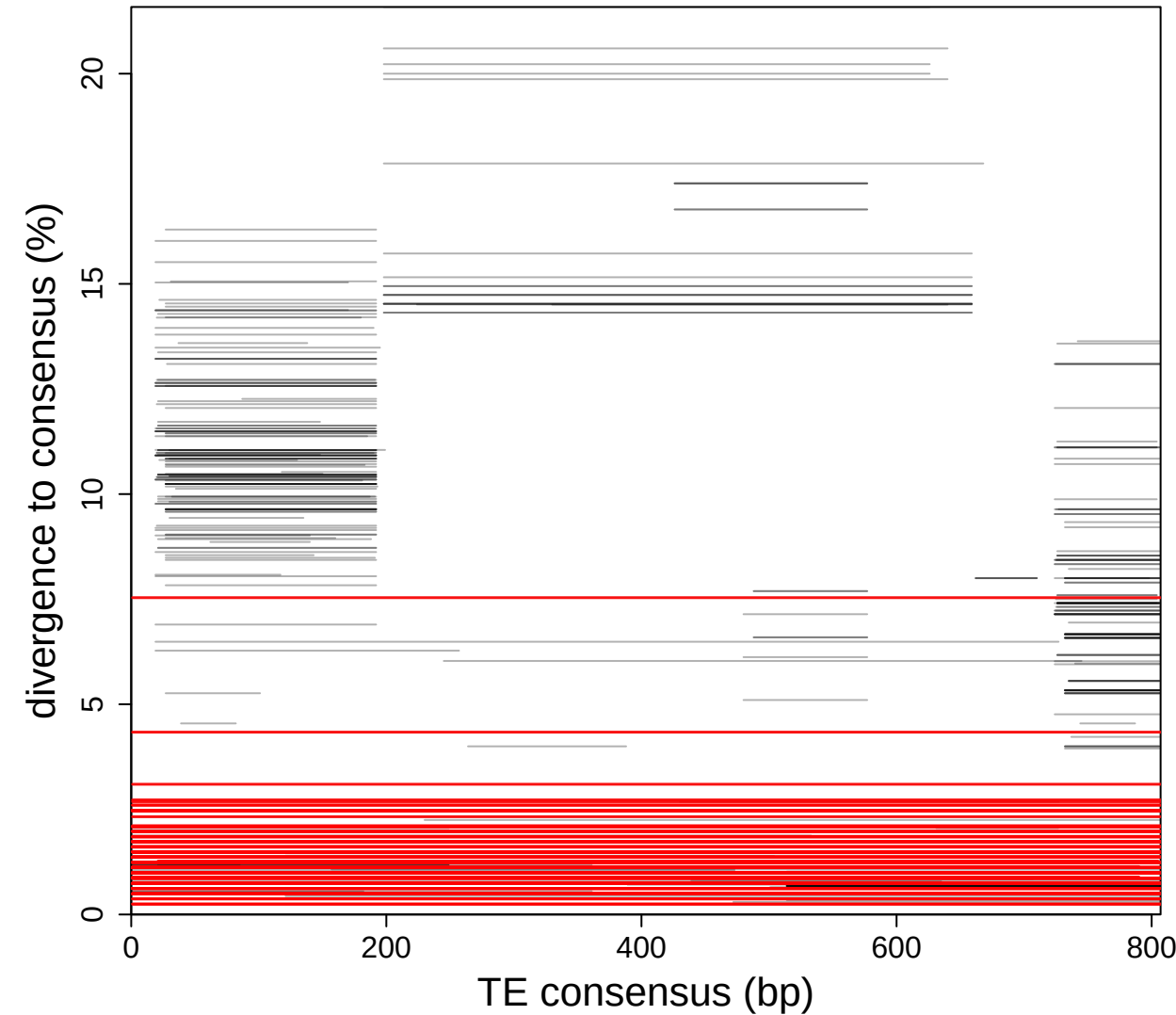


ly_6942.fasta.b.bed uf.bed g_1.bed fm_1.bed_0_0 bcln.fa aln.f
size: 18035bp; fragments: 5720; full length: 0 (>=16231.5bp)

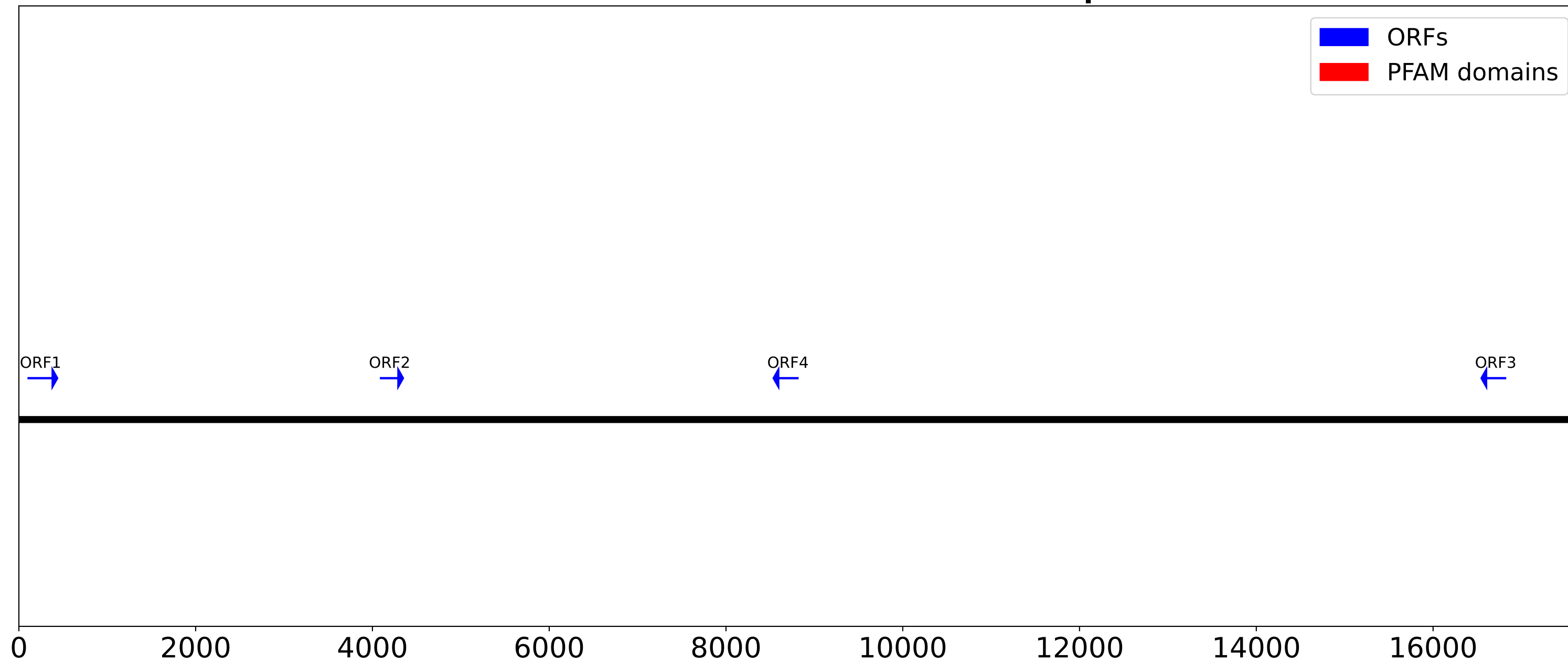
After TEtrimmer Extended plot Blue lines are boundaries



TE: rnd_5_family_6942
size: 807bp; fragments: 595; full length: 258 (≥ 726.3 bp)



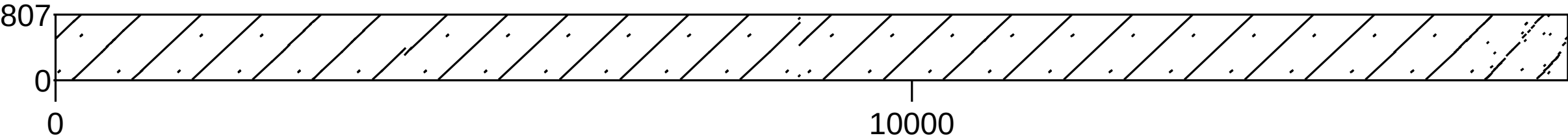
After TEtrimmer ORF and PFAM domain plot



Dotplot

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

nsus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)