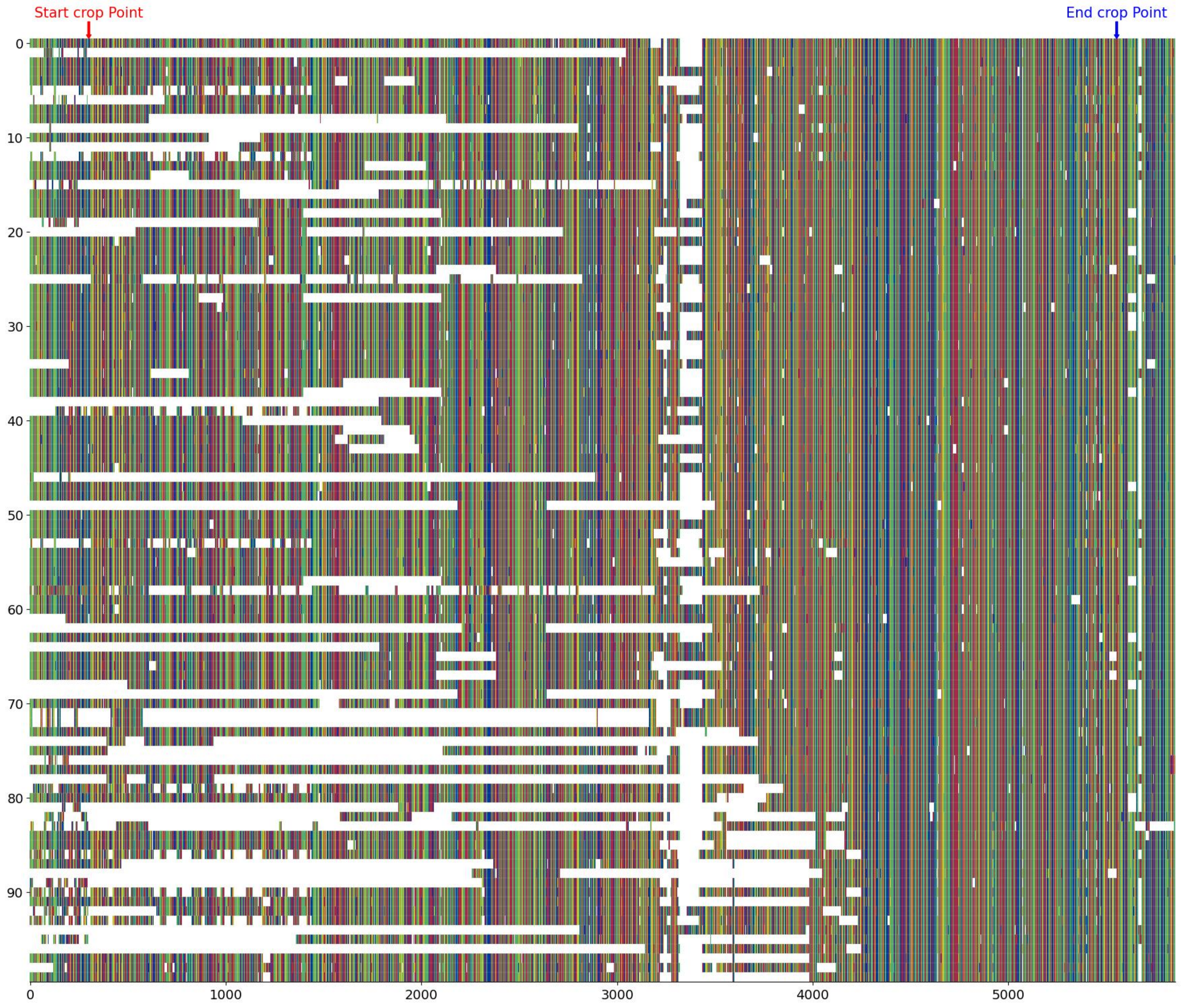


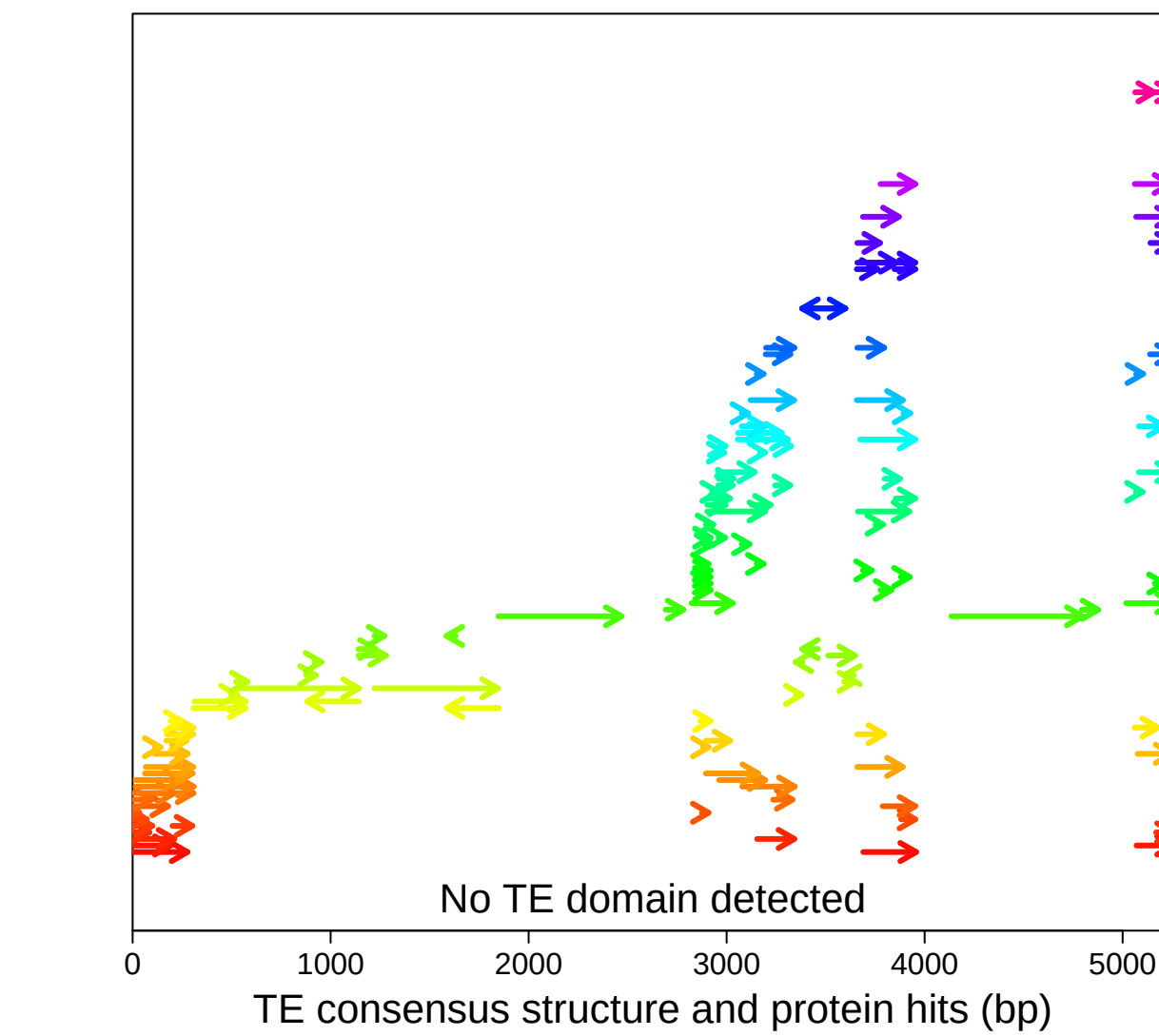
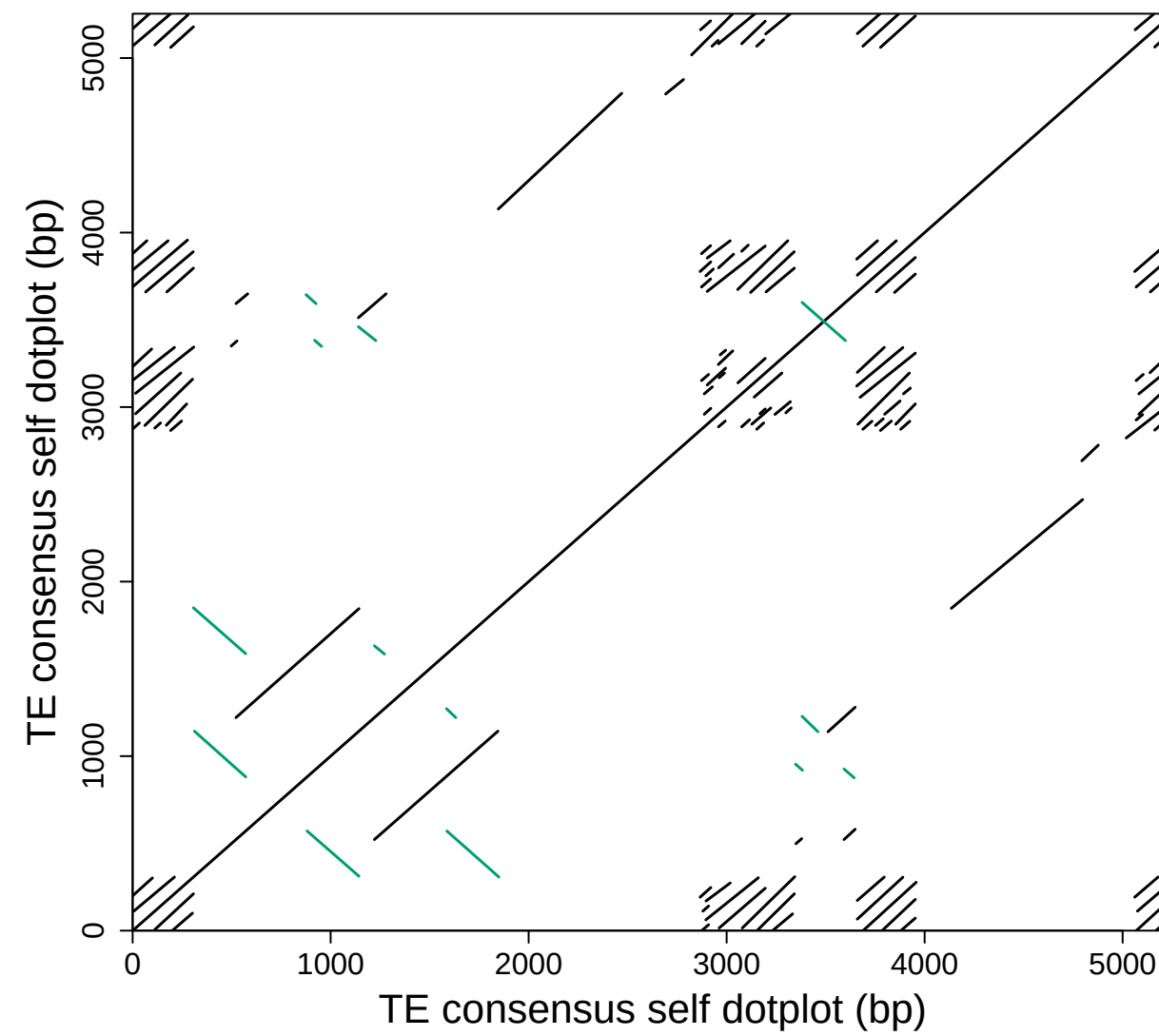
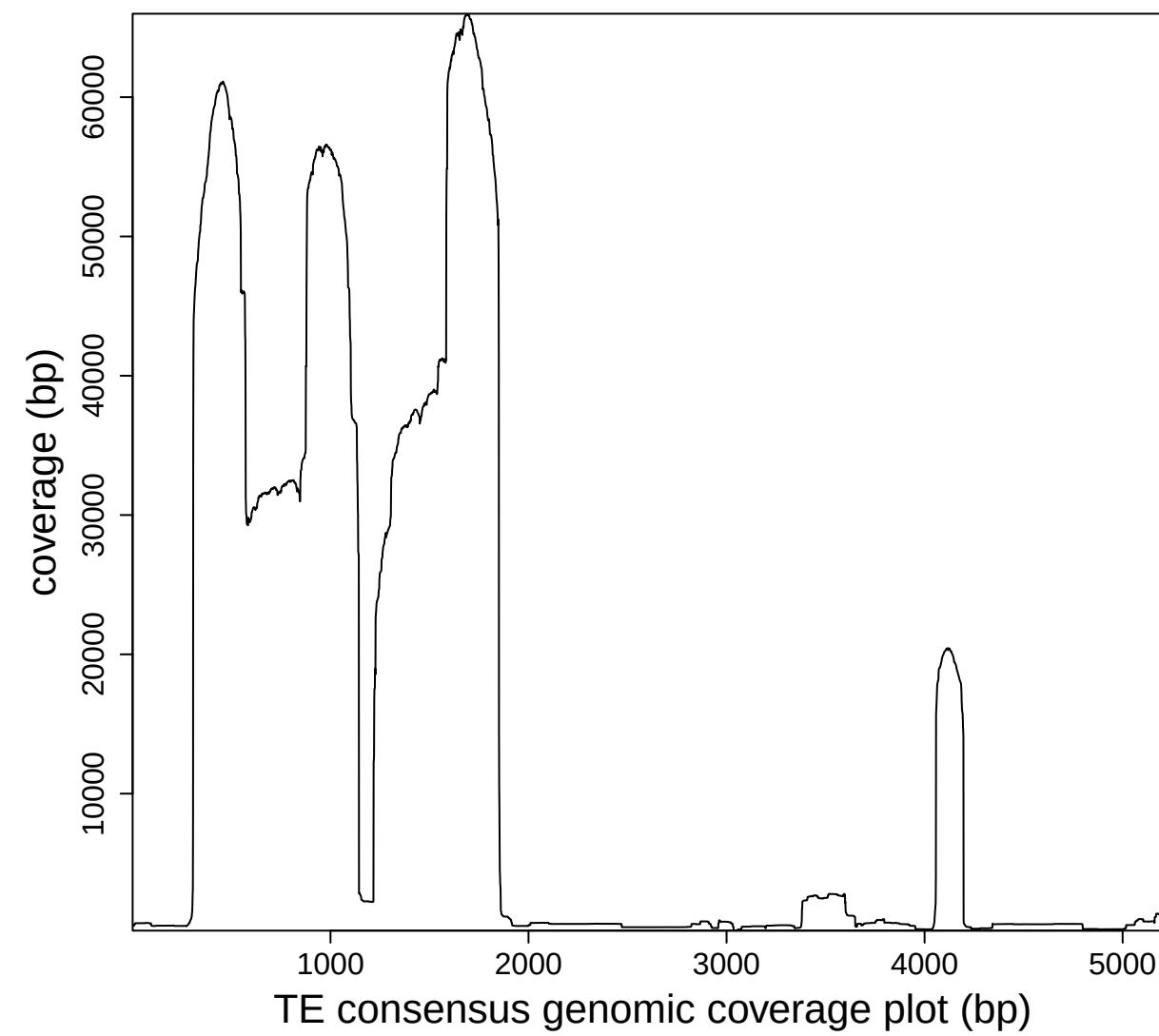
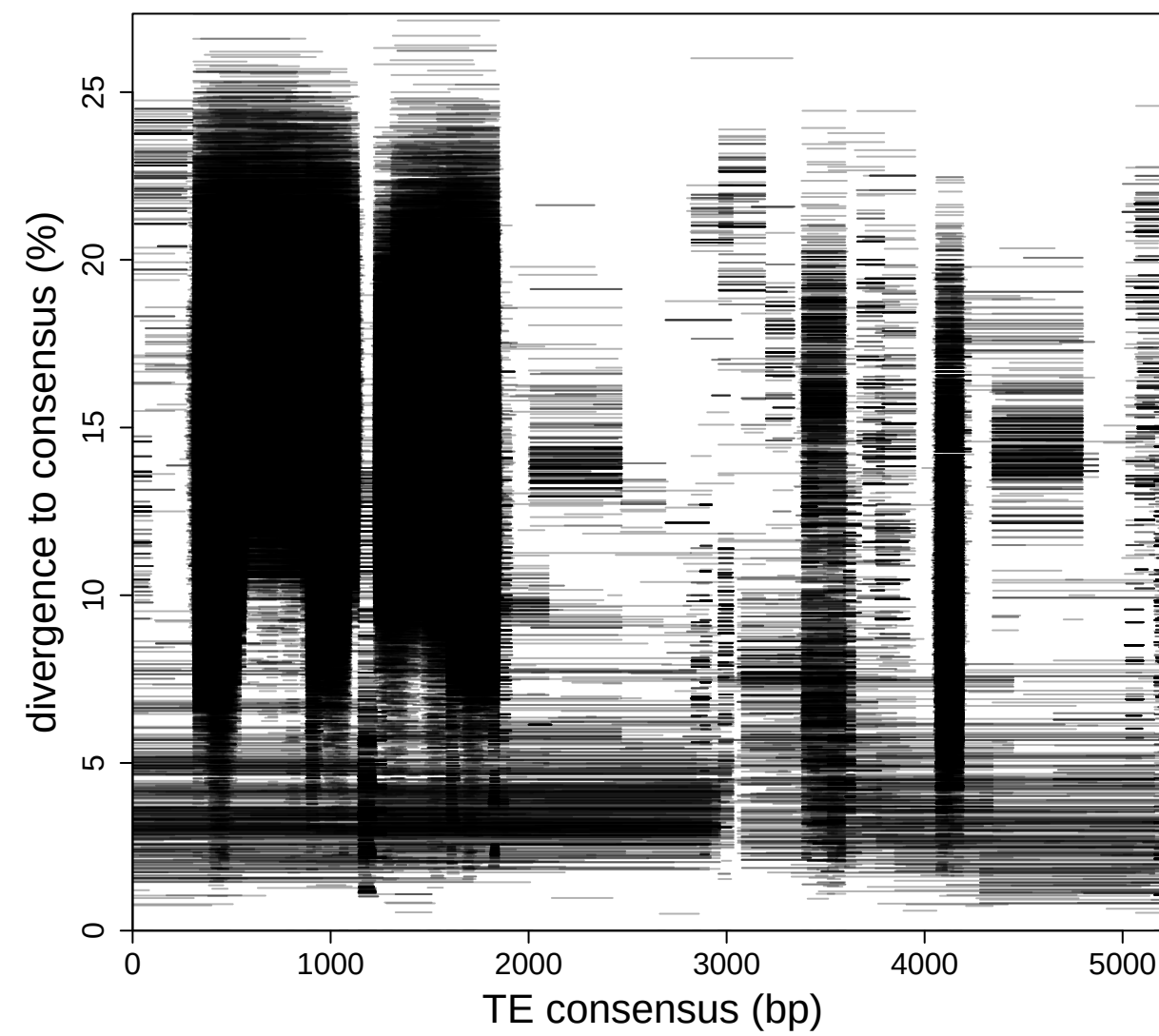
Start crop Point

End crop Point



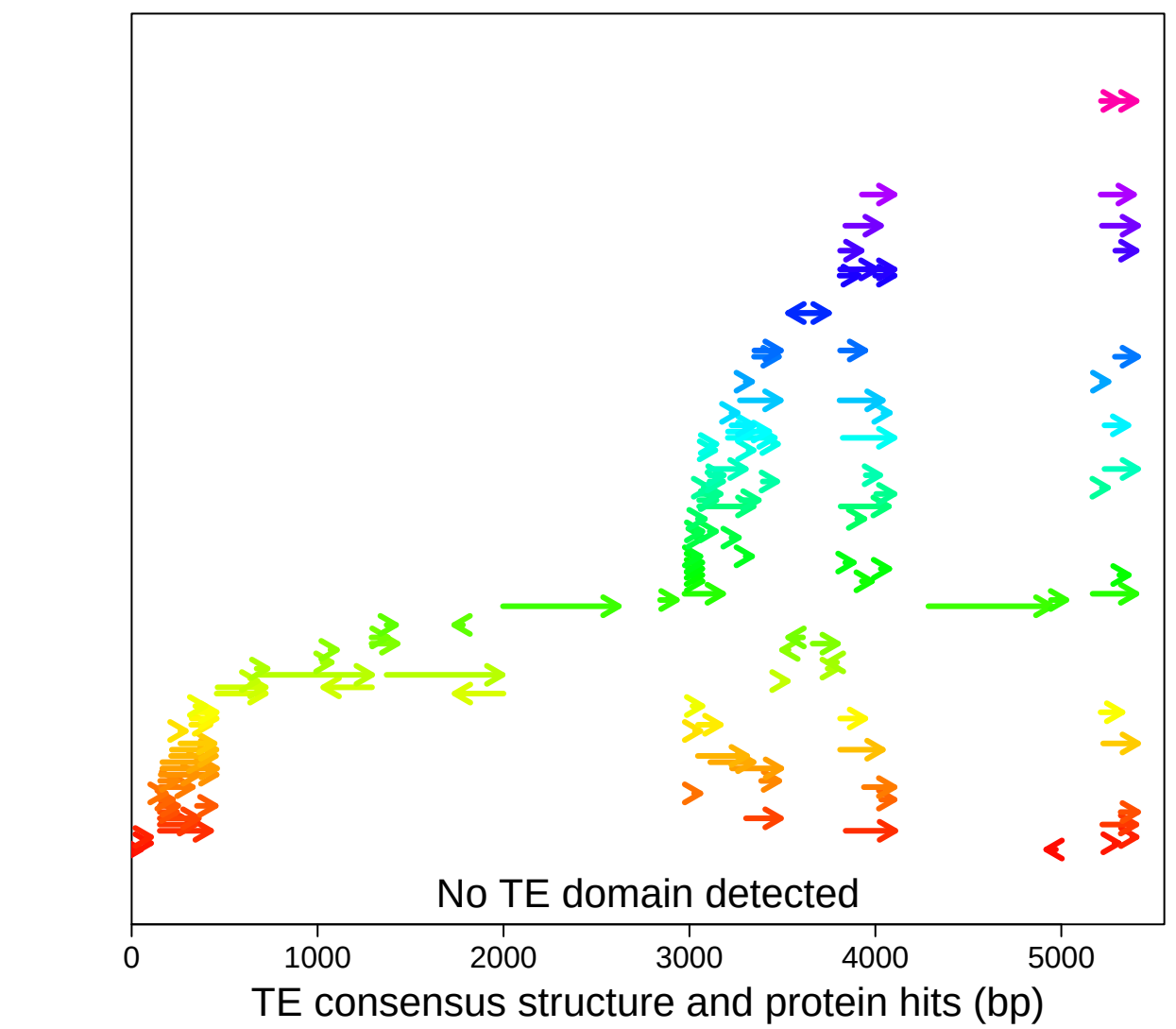
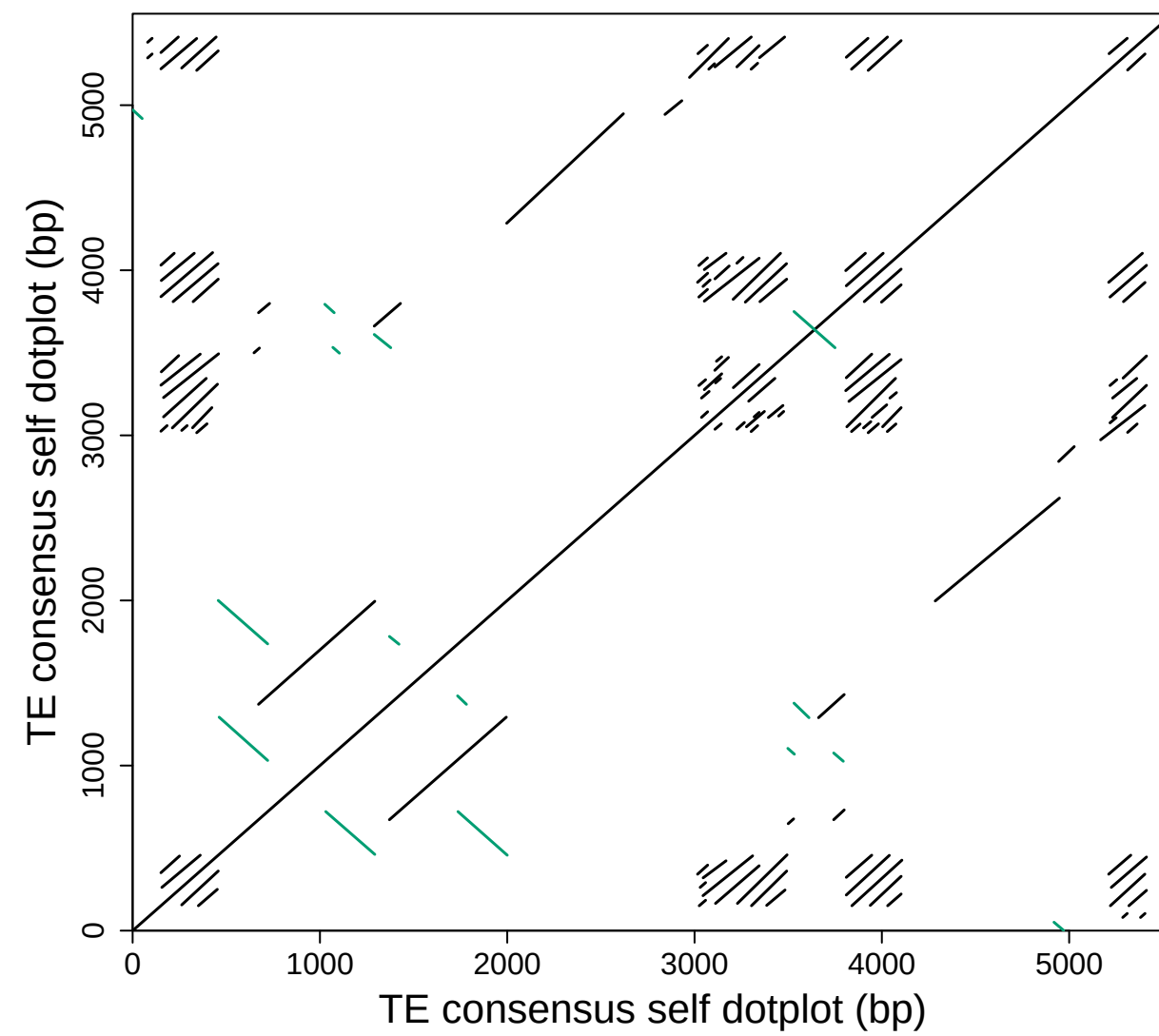
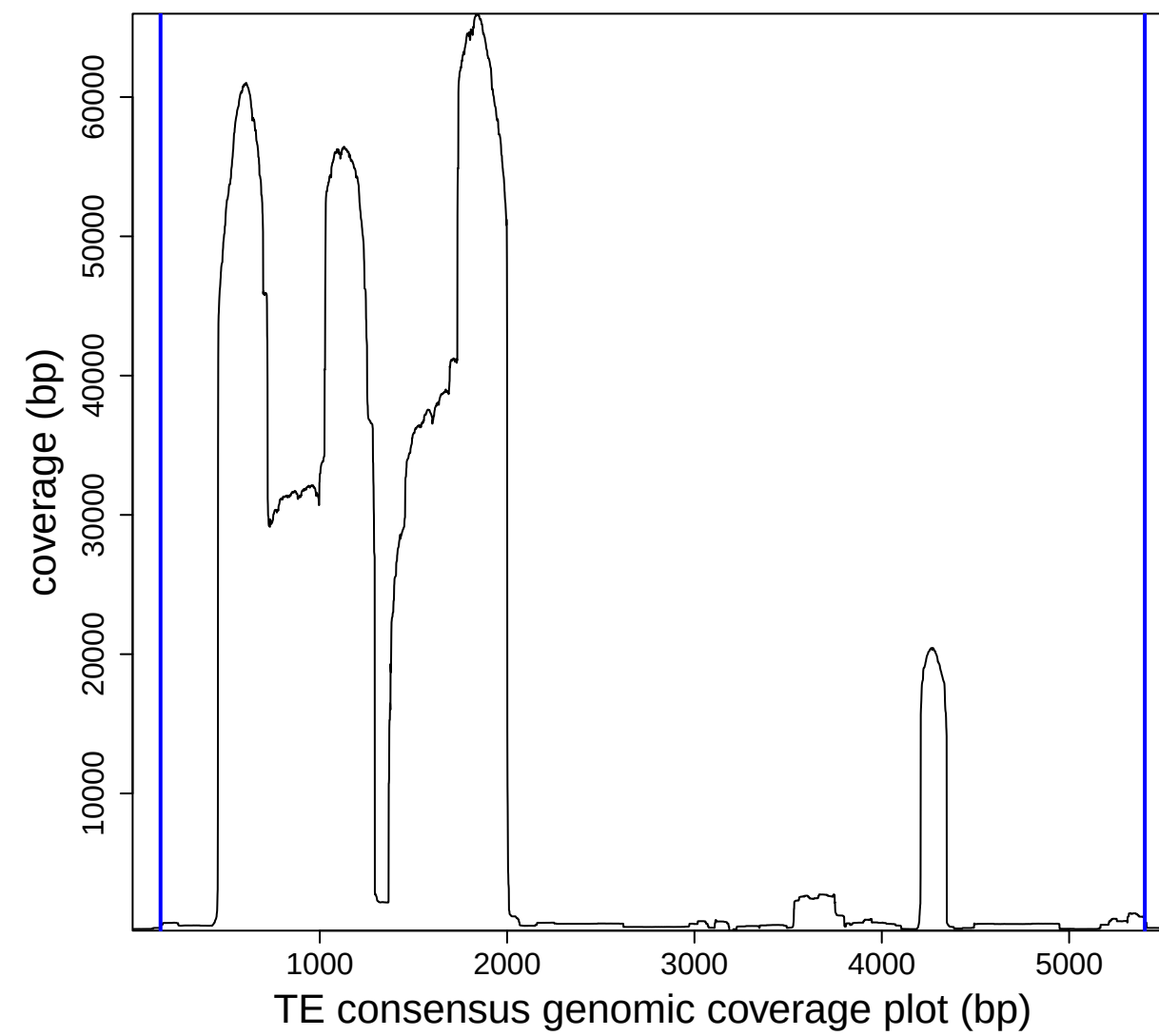
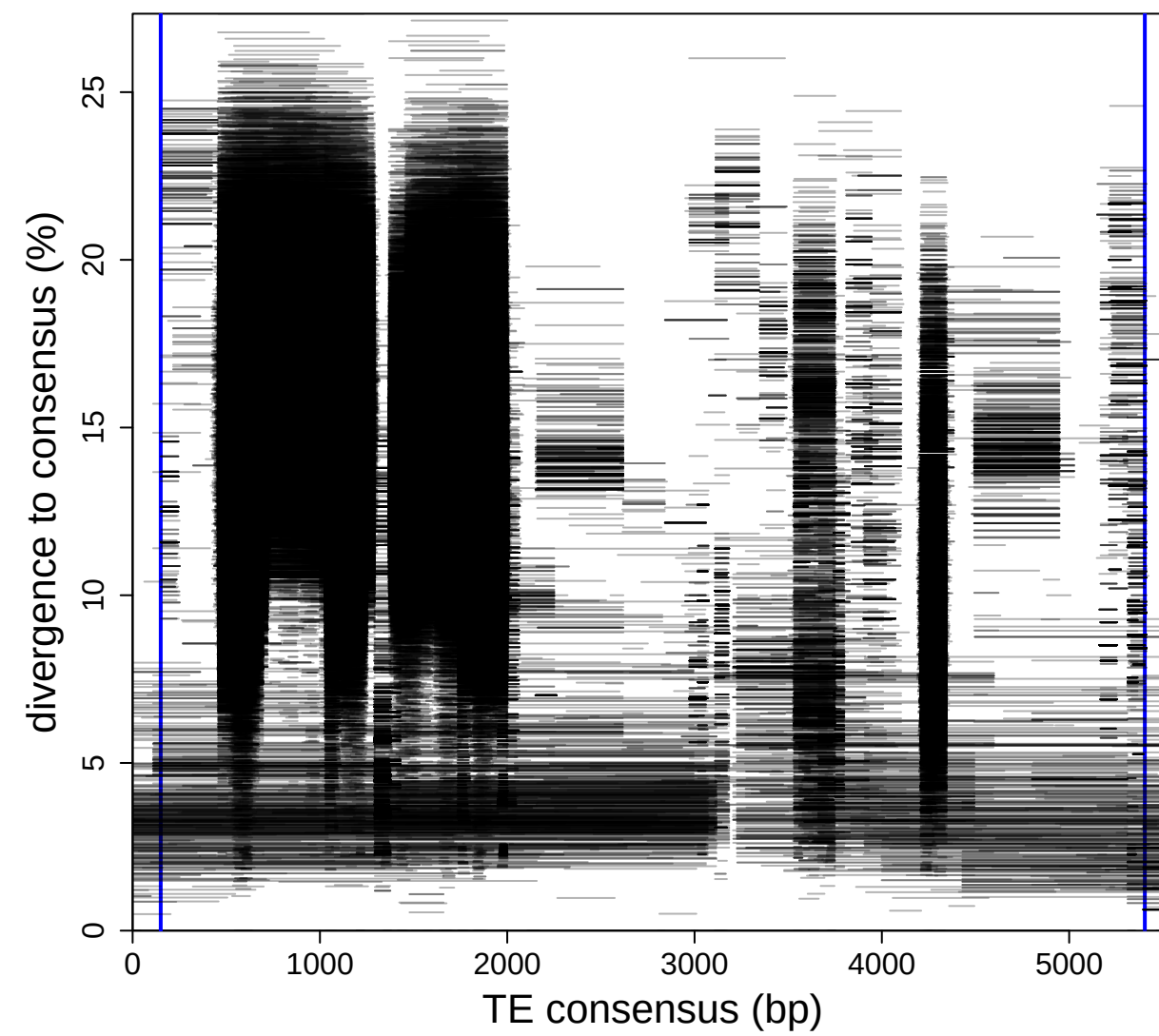
bed_uf.bed g_1.bed fm_1.bed 0 0 bcln.fa aln.fa cl.fa gs.fa c
size: 5254bp; fragments: 223410; full length: 0 (>=4728.6bp)

After TEtrimmer 5254 bp



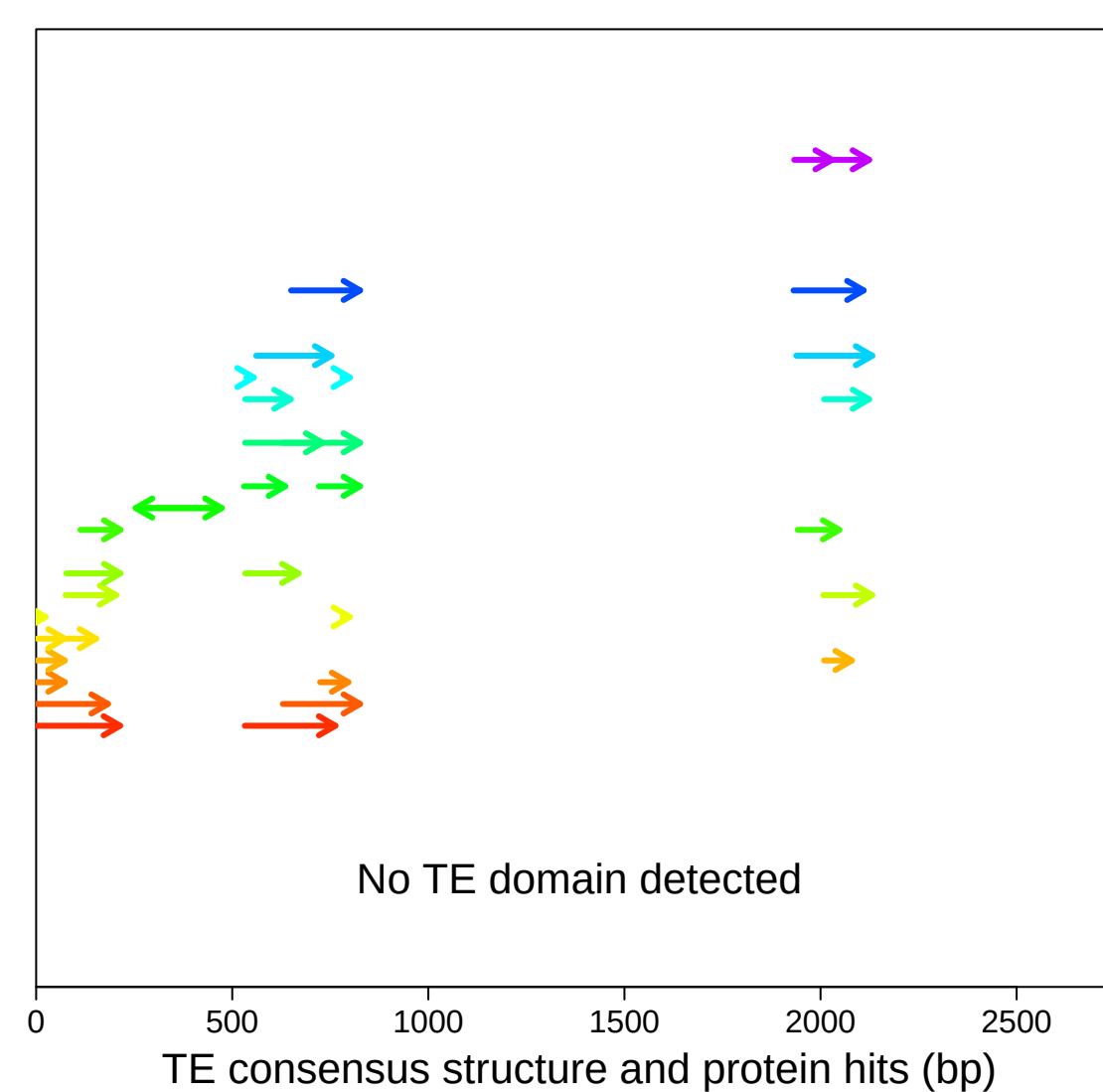
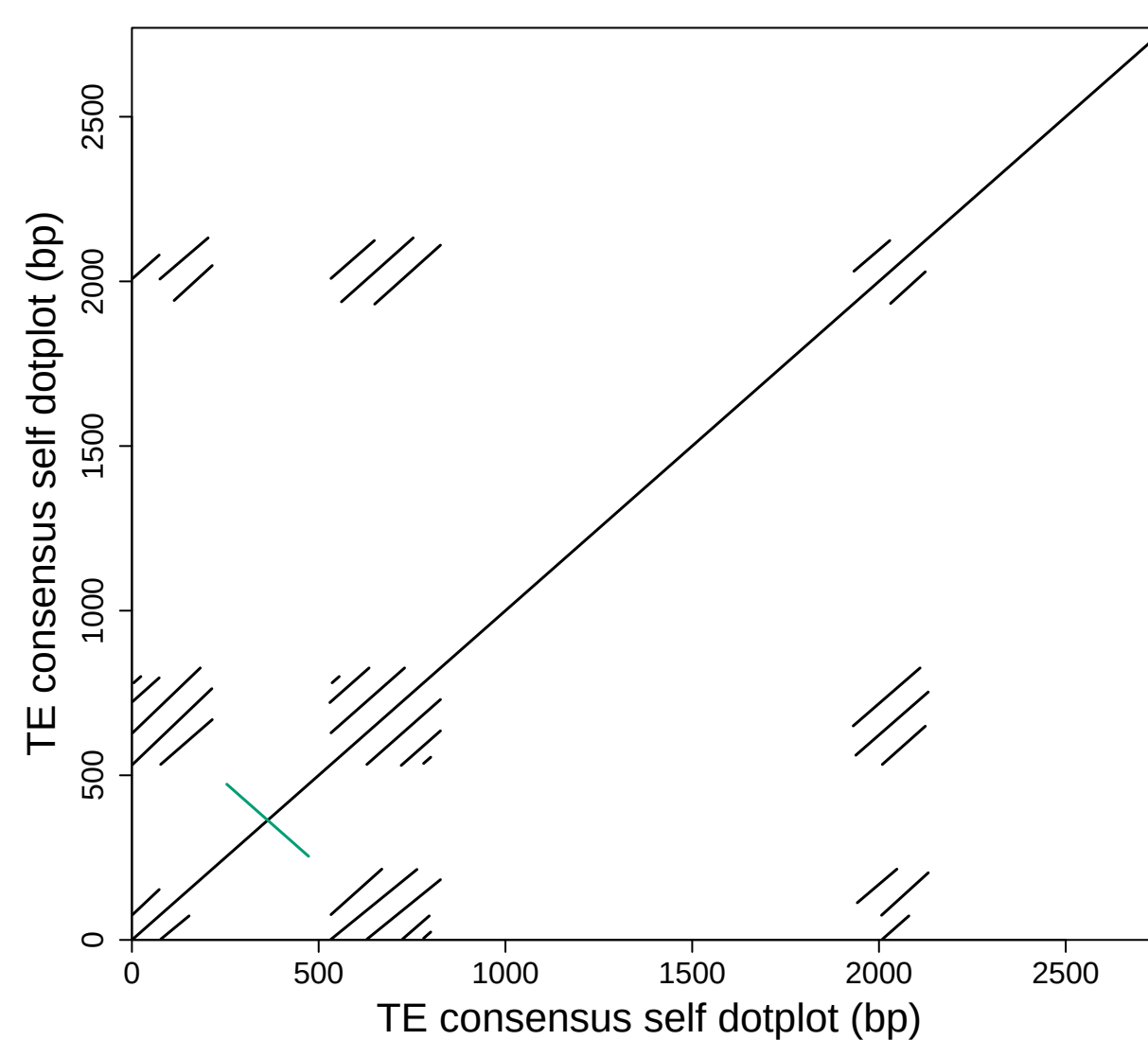
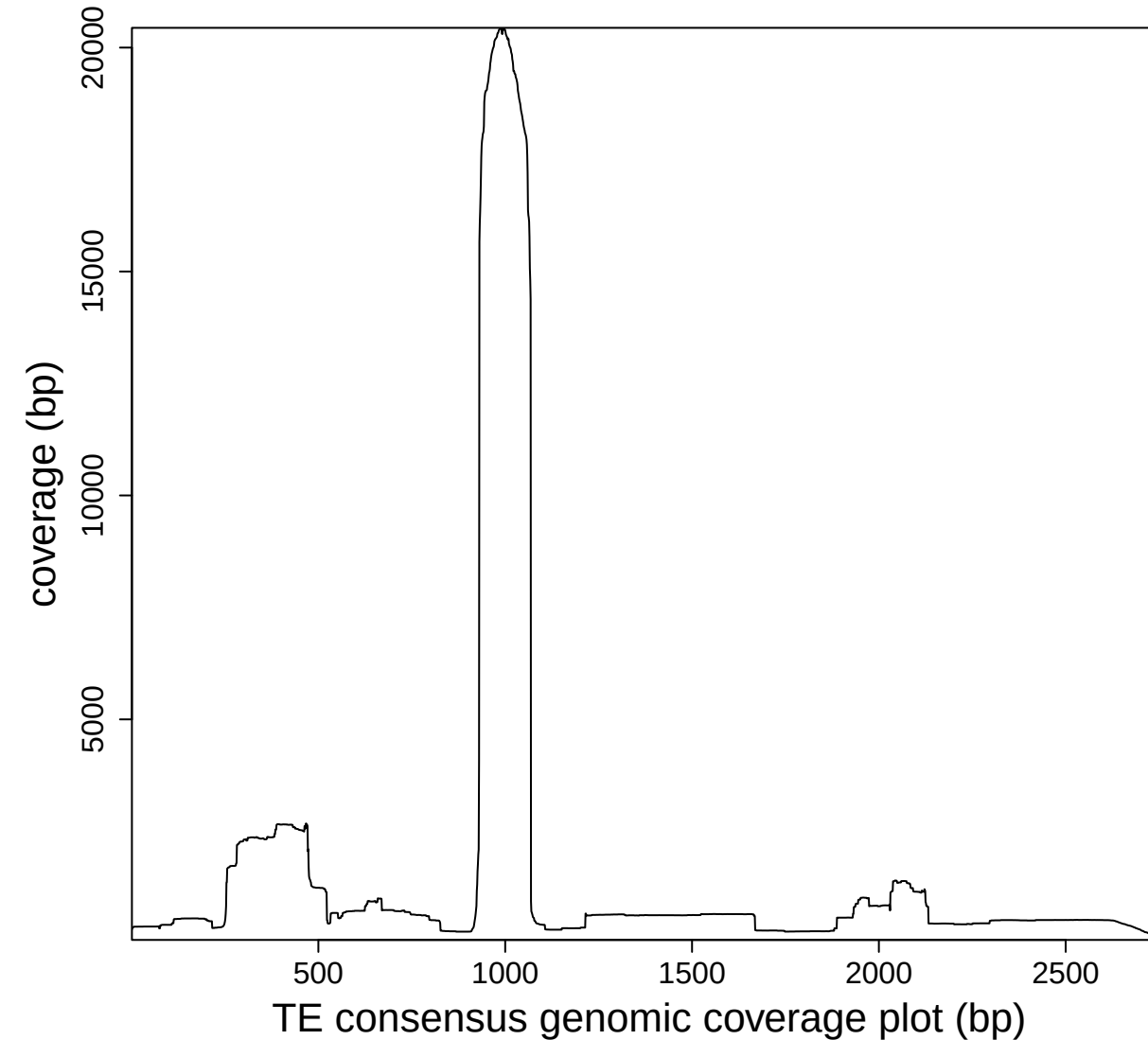
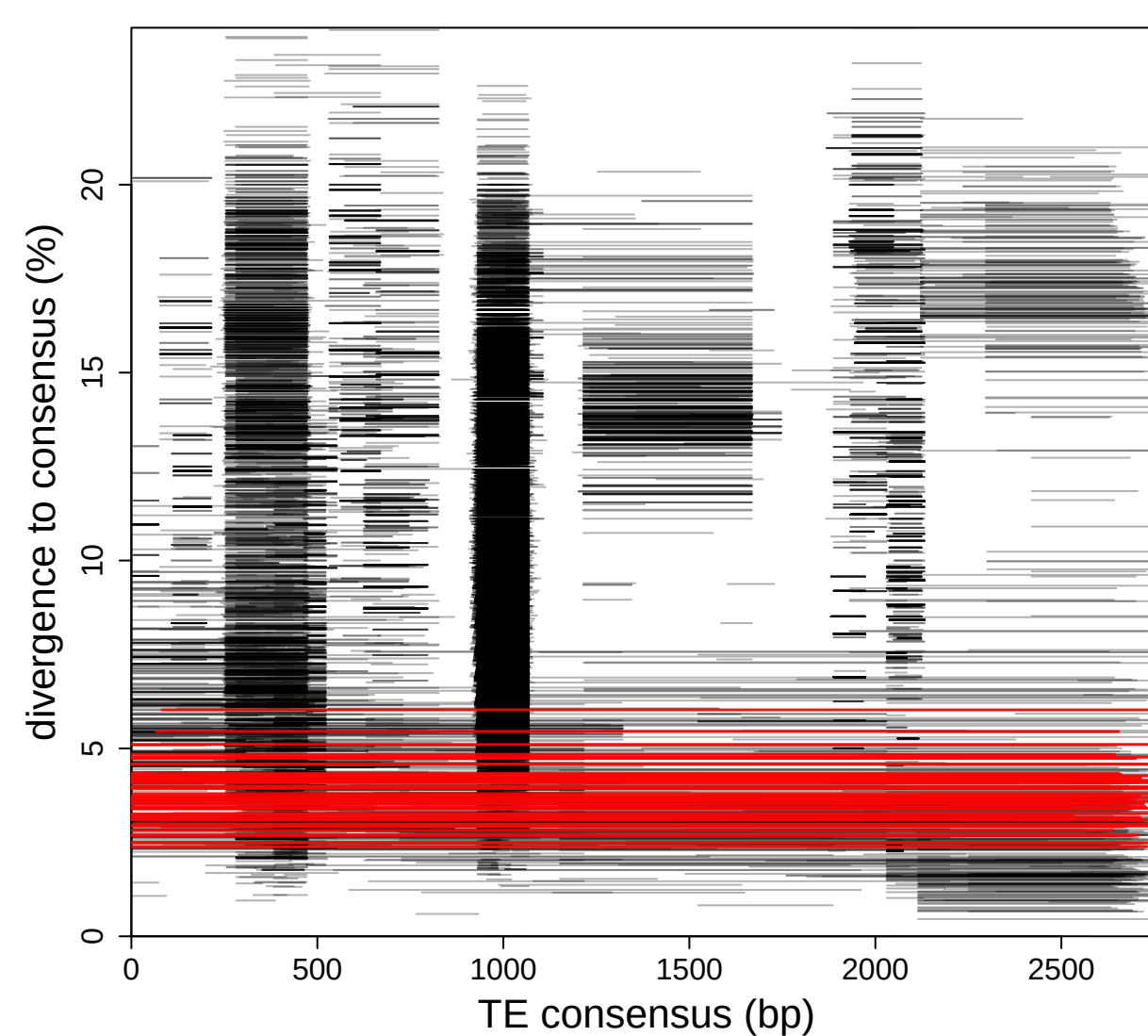
ly_4680.fasta.b.bed uf.bed g_1.bed fm_1.bed_0_0 bcln.fa aln.f
size: 5554bp; fragments: 222949; full length: 0 (>=4998.6bp)

After TEtrimmer Extended plot Blue lines are boundaries

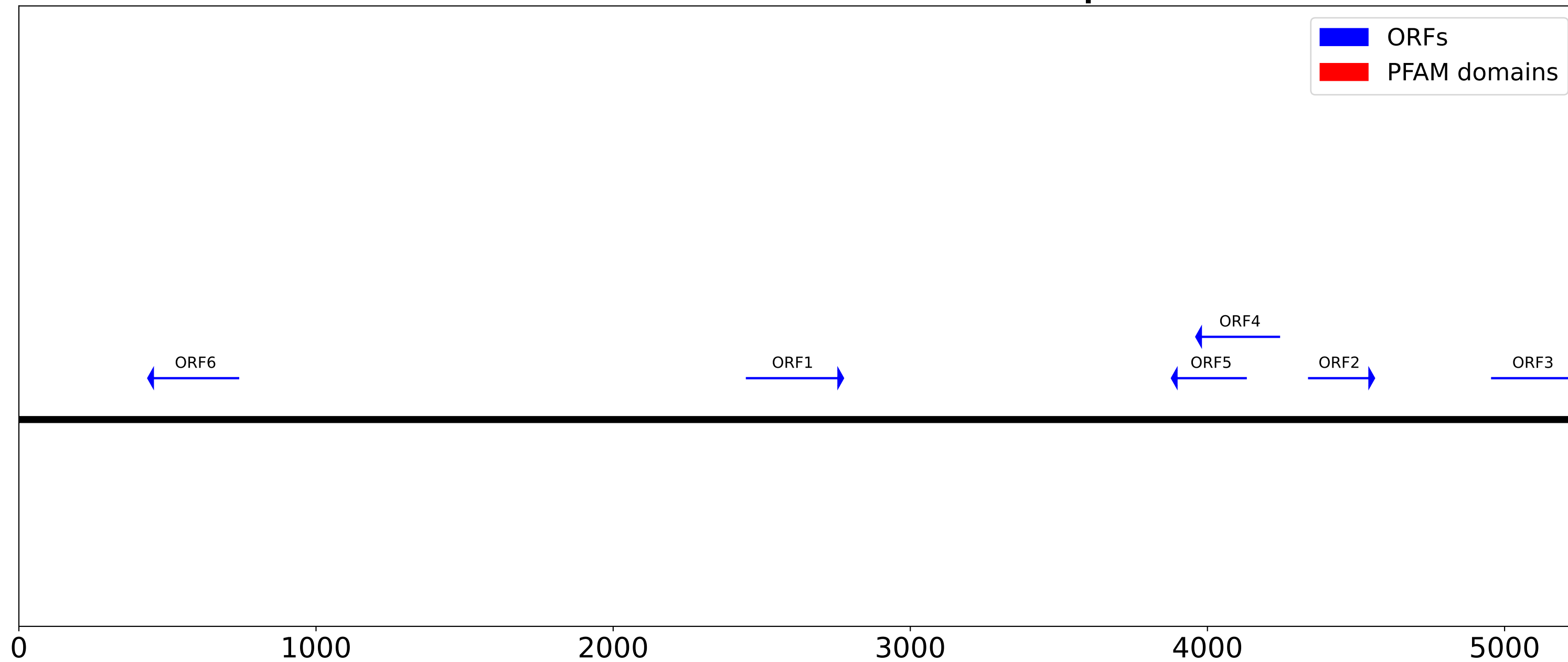


TE: rnd_5_family_4680
size: 2770bp; fragments: 27423; full length: 54 (≥ 2493 bp)

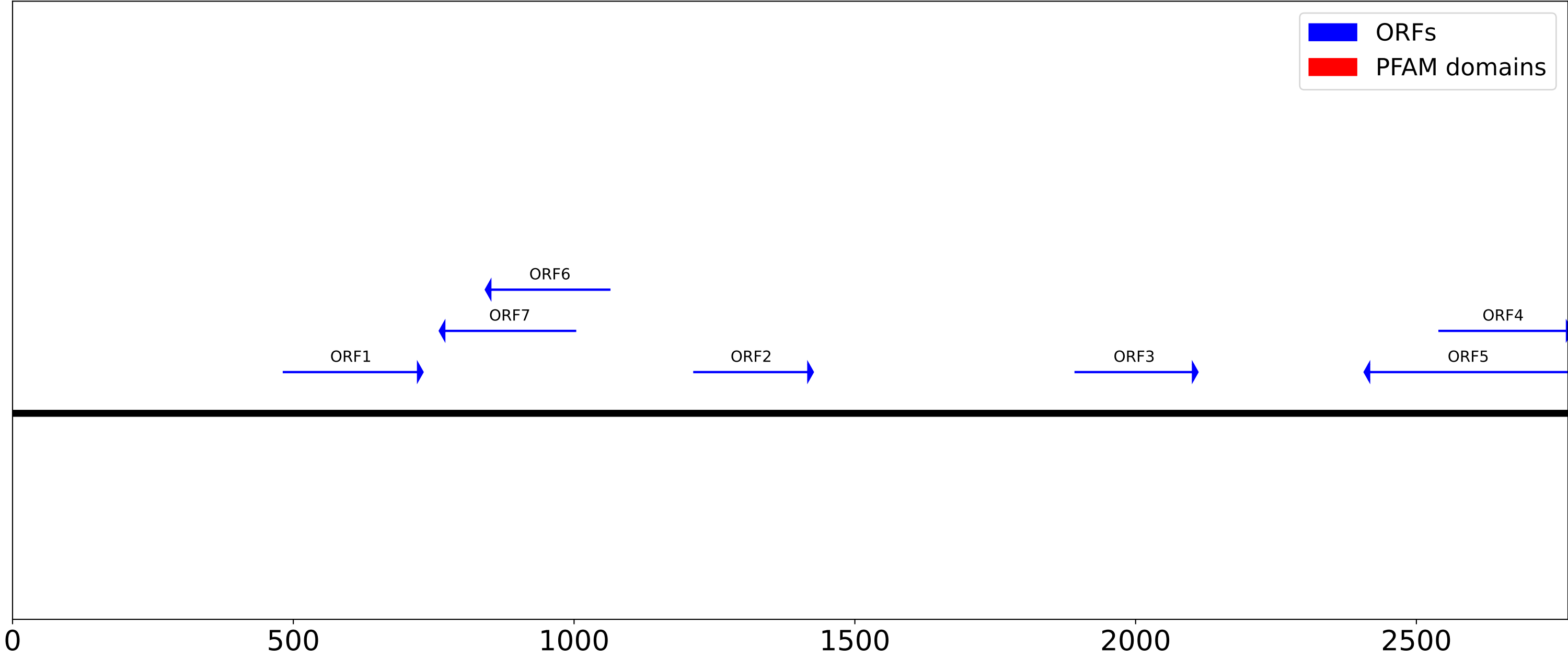
Before TEtrimmer 2770 bp



After TEtrimmer ORF and PFAM domain plot



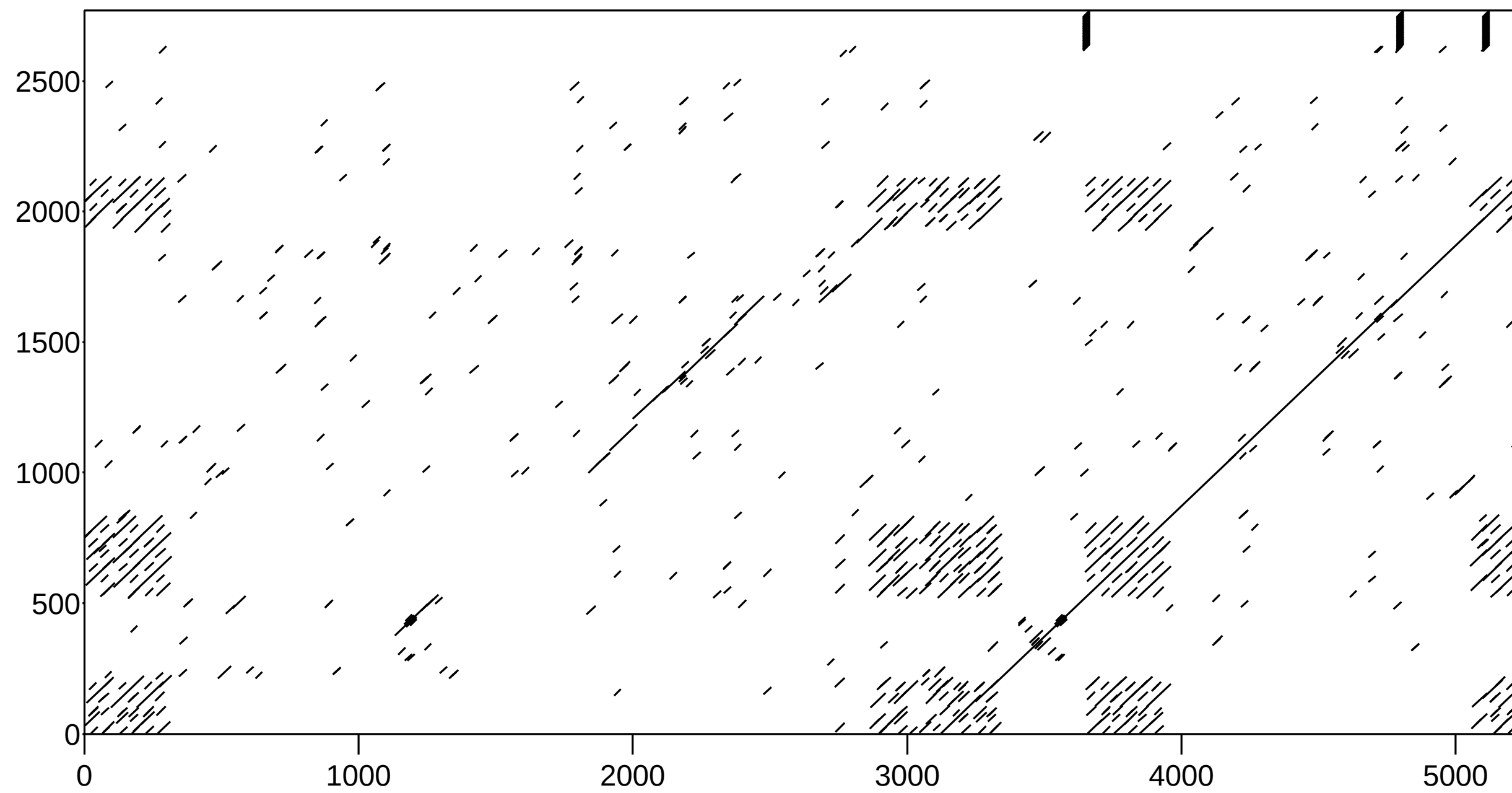
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)