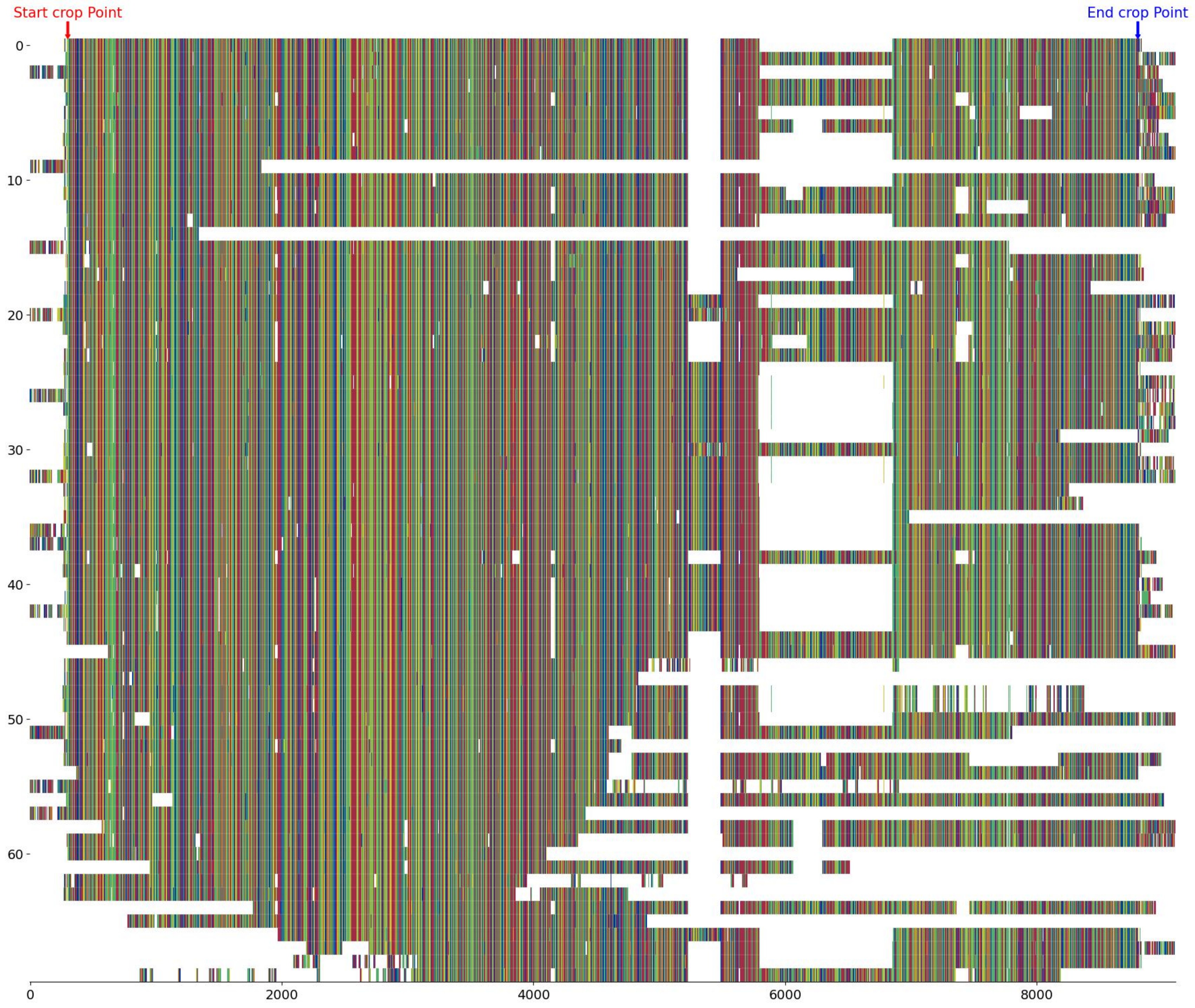
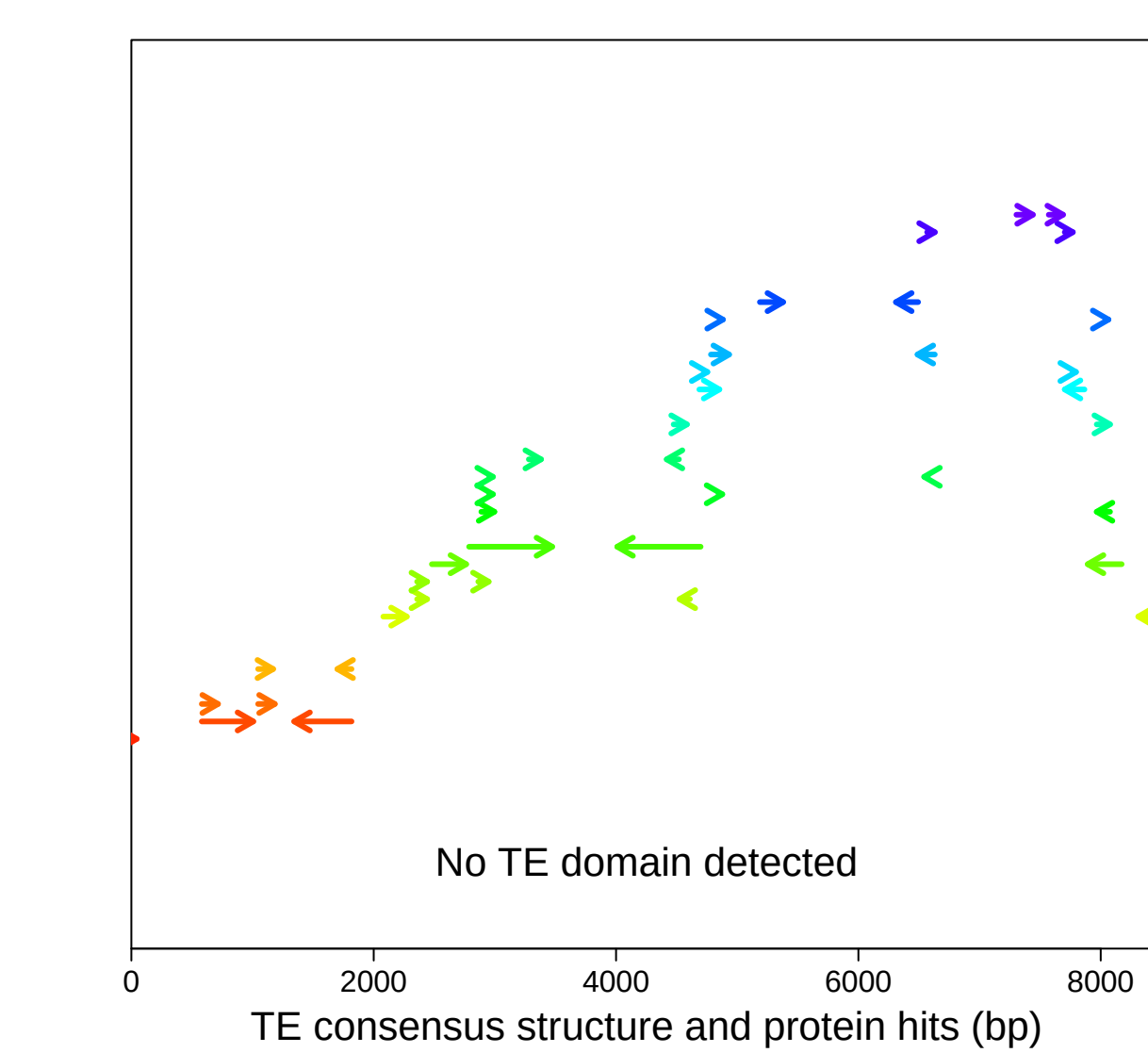
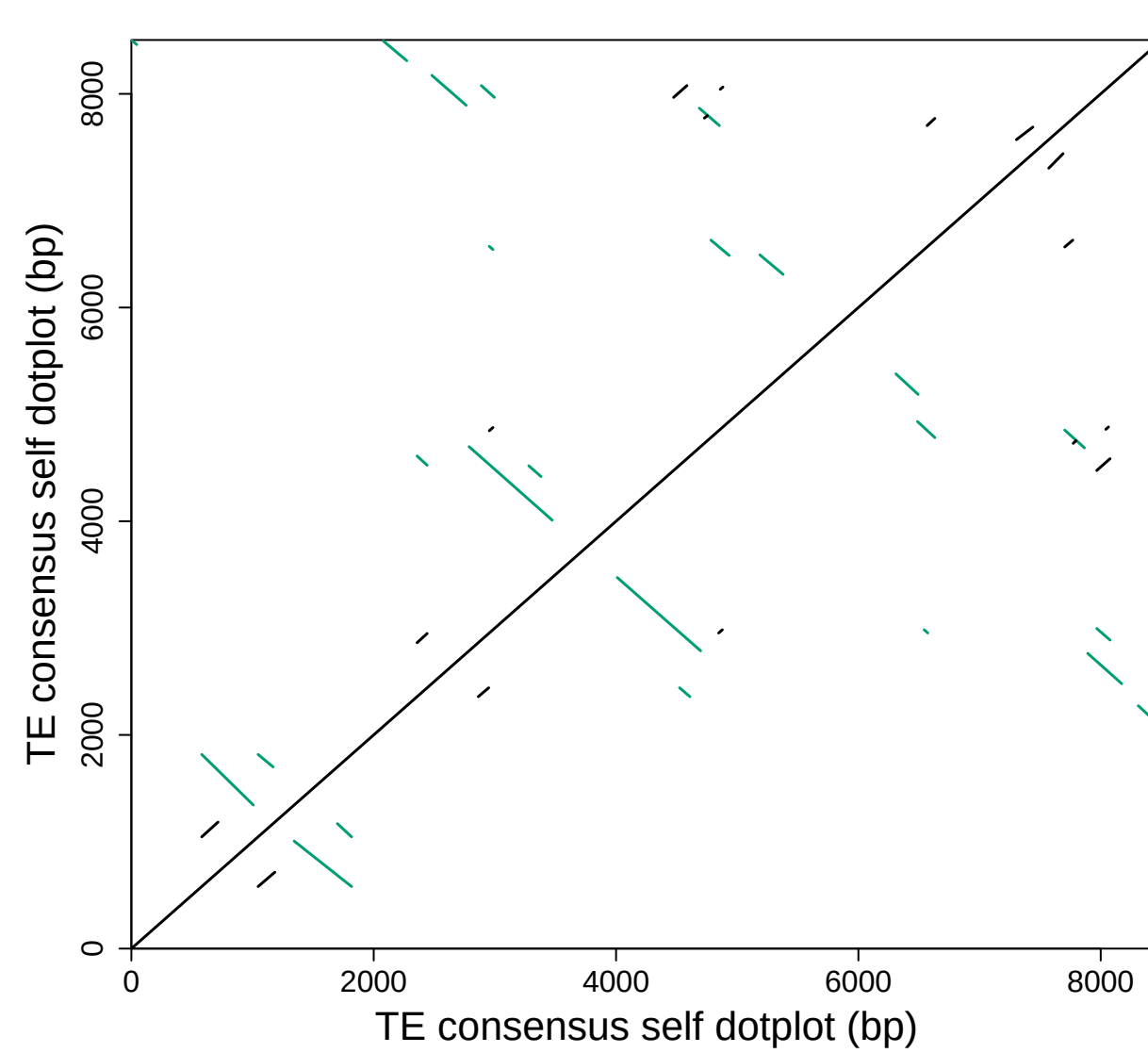
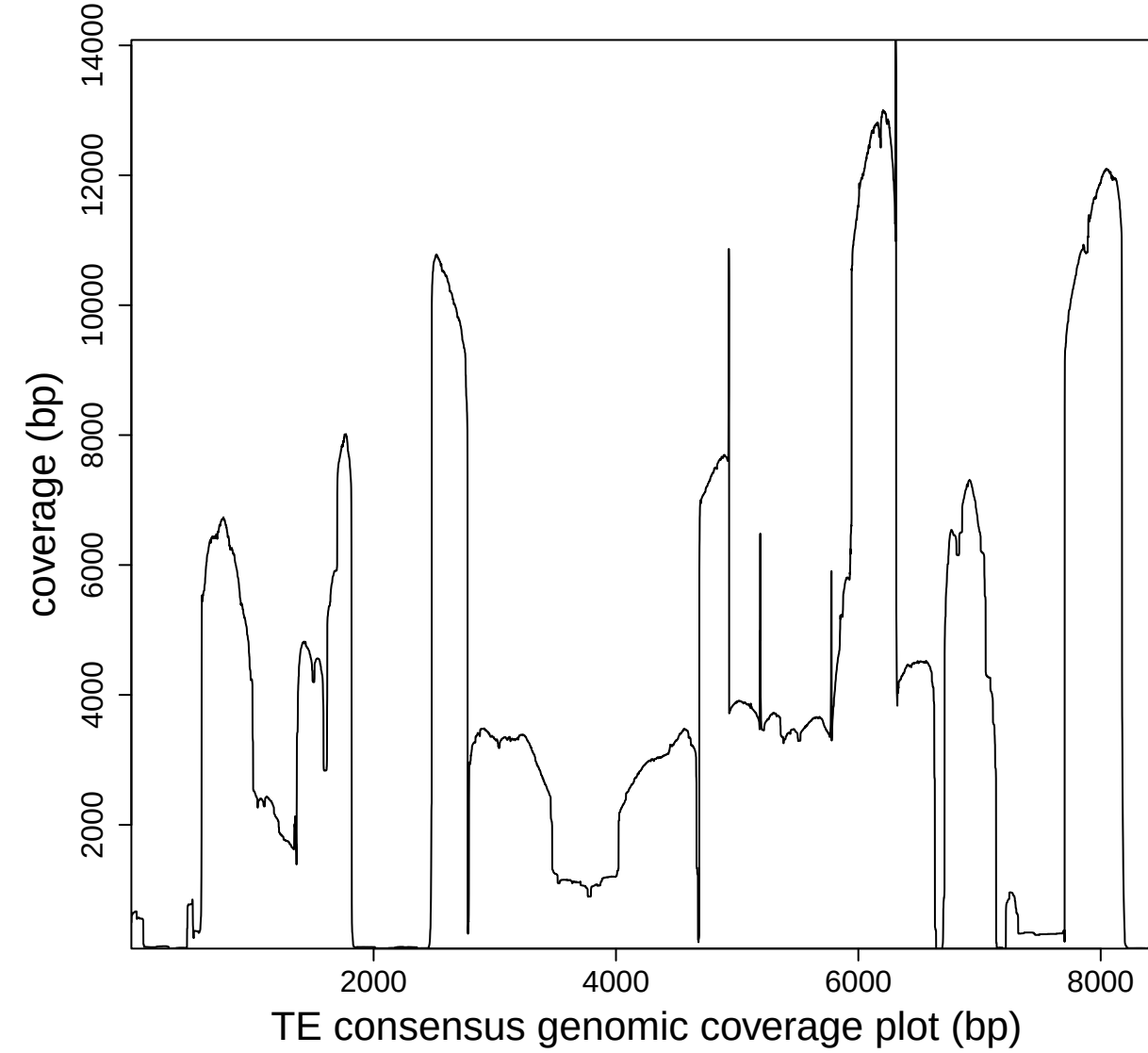
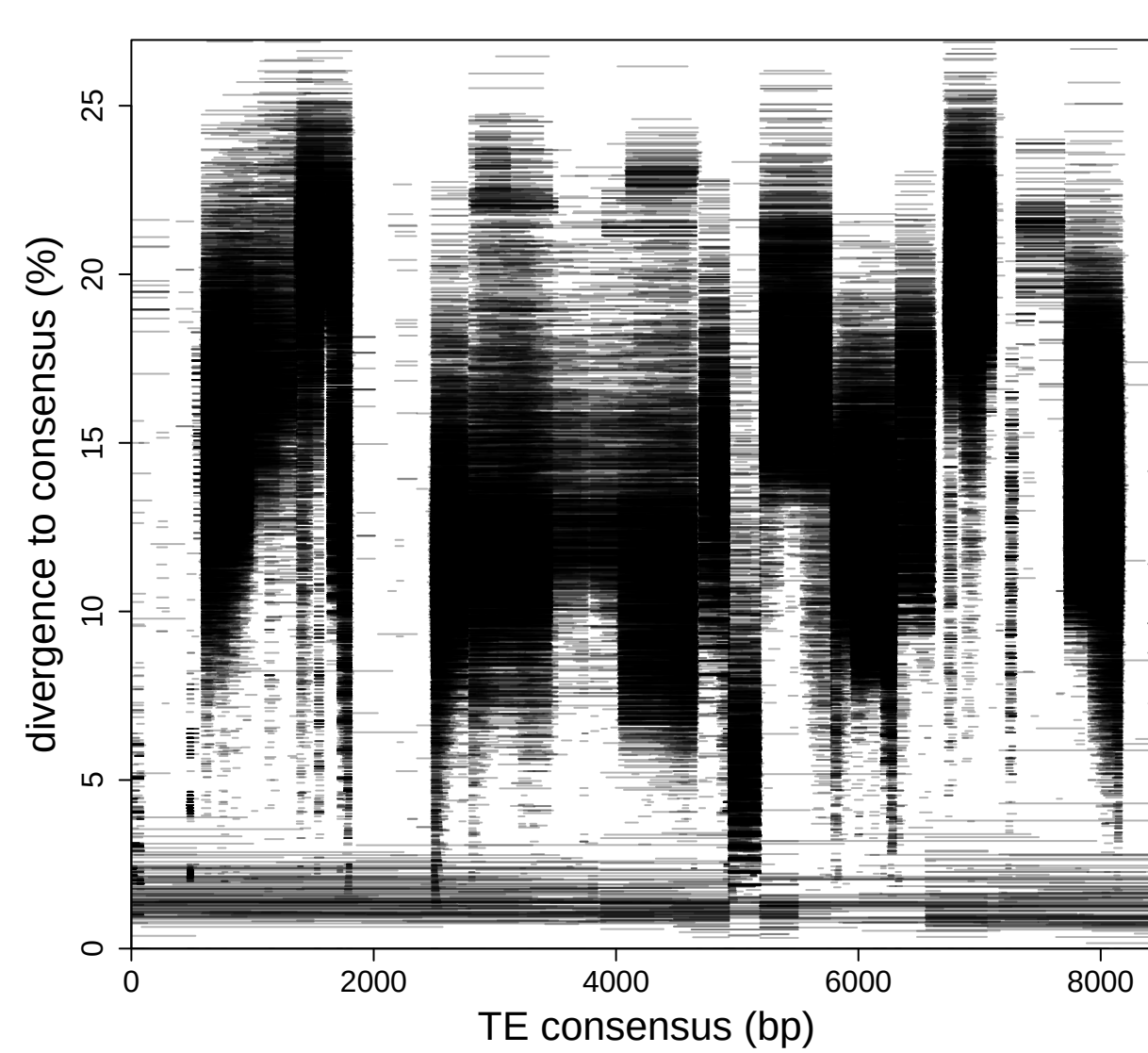






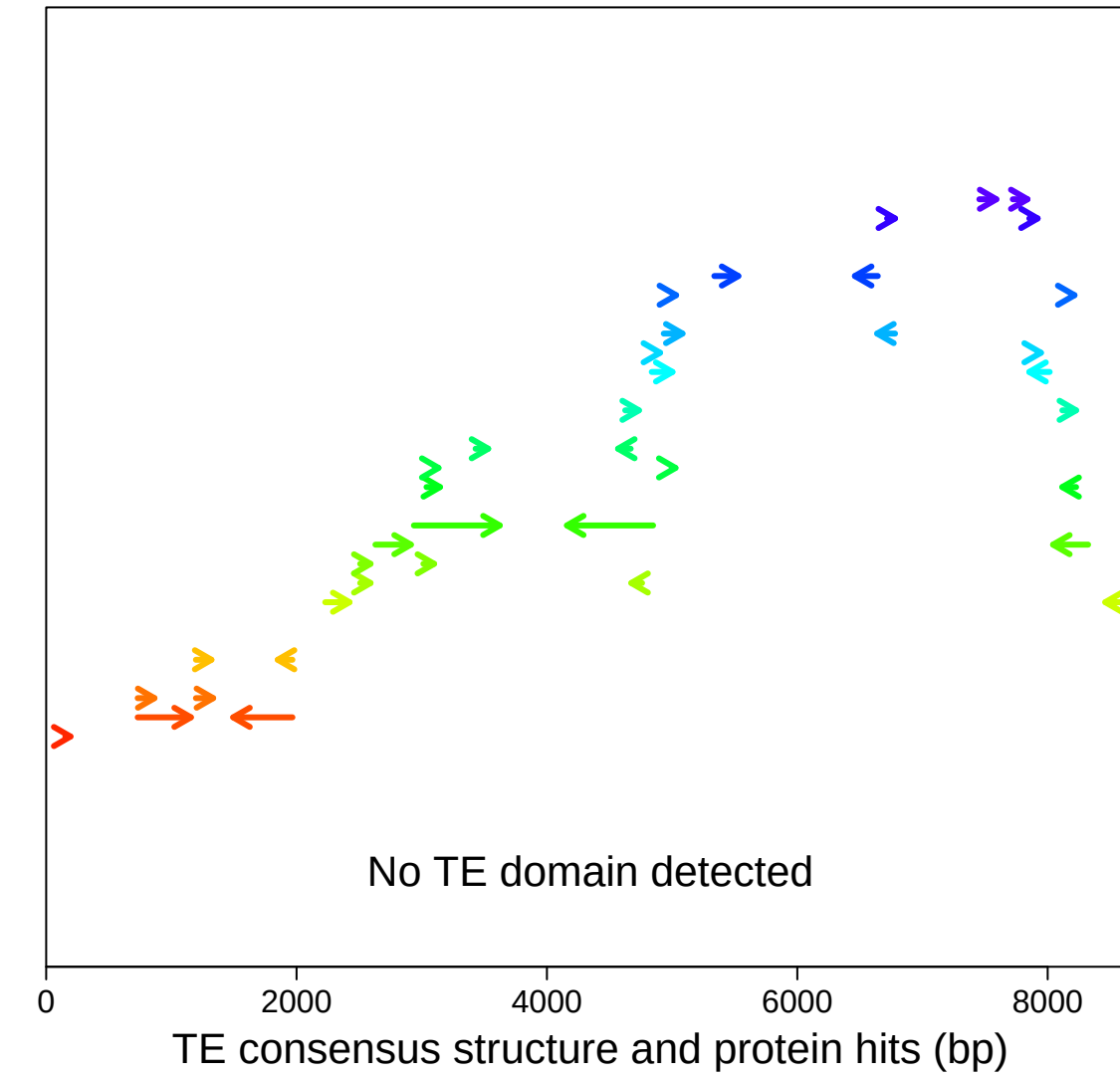
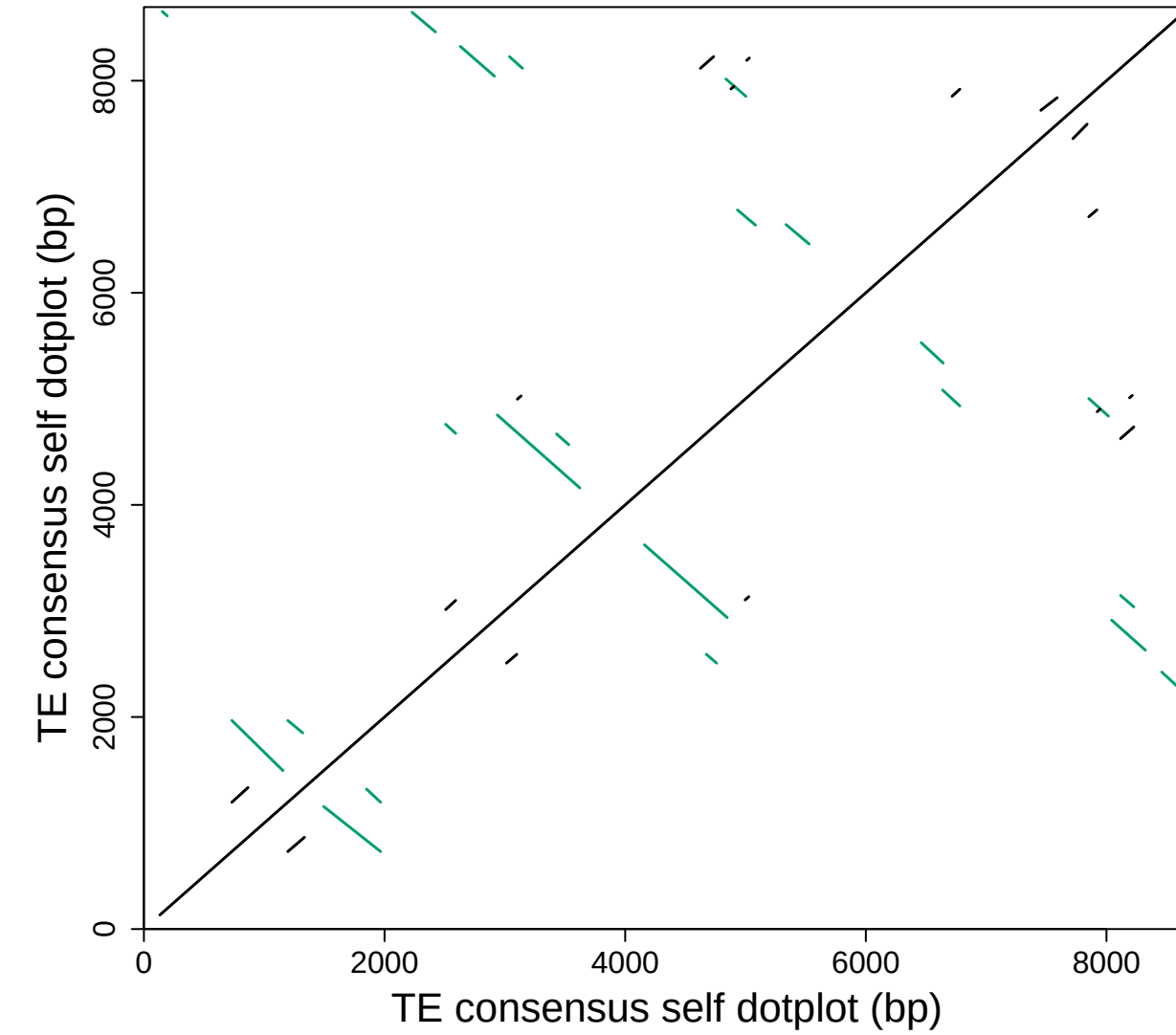
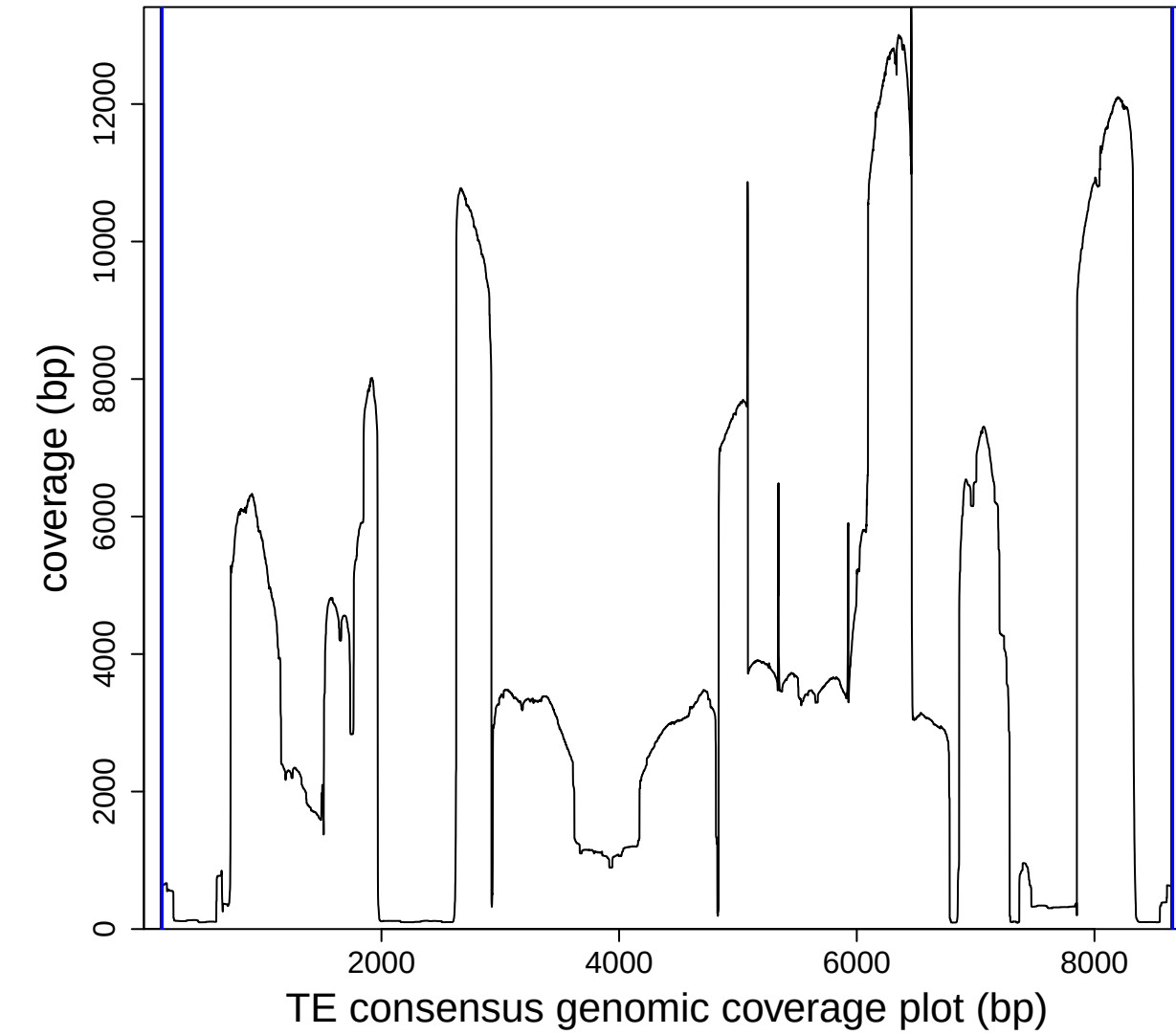
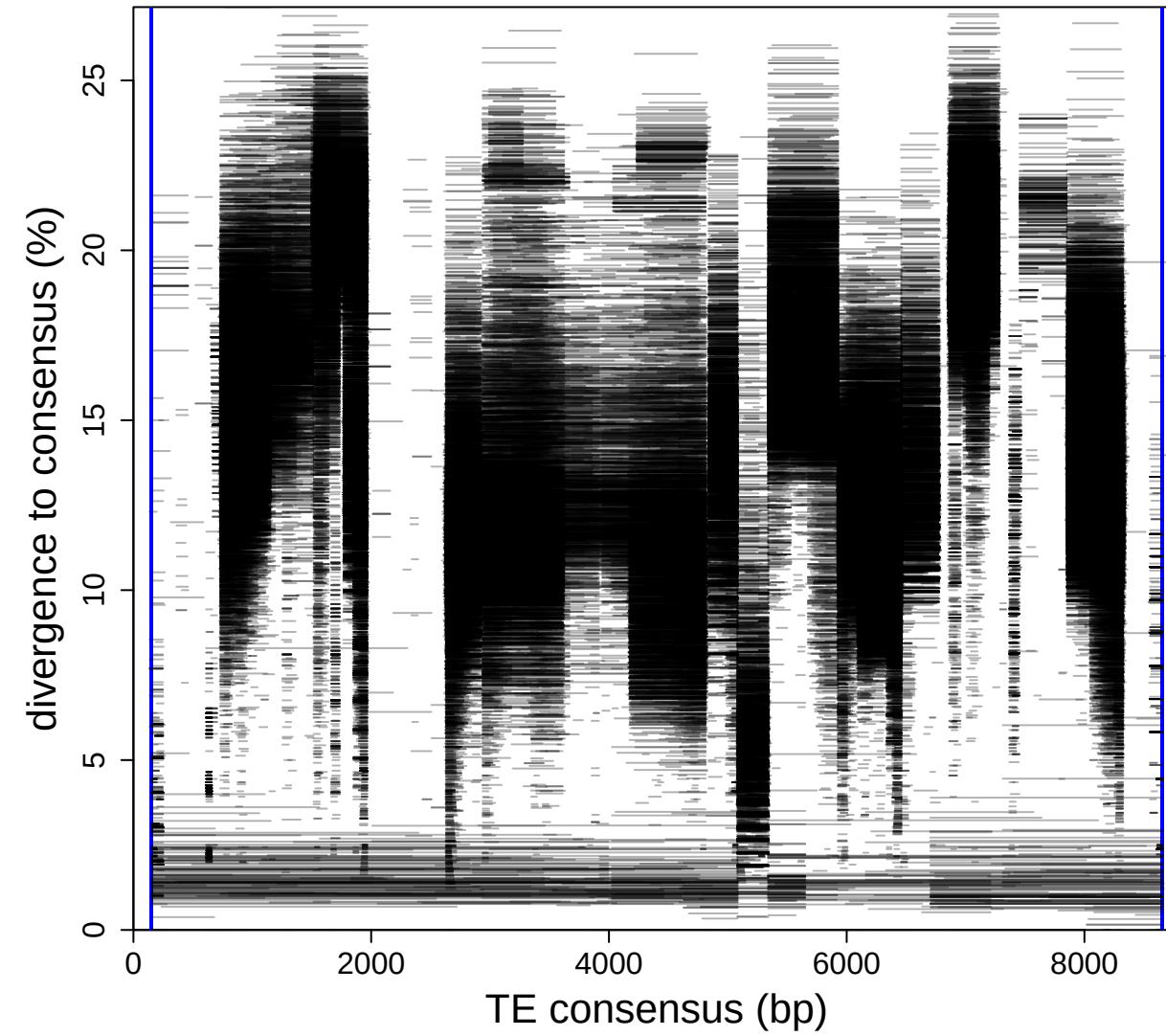
a.b.bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.f
size: 8504bp; fragments: 100151; full length: 0 (>=7653.6bp)

After TEtrimmer 8504 bp



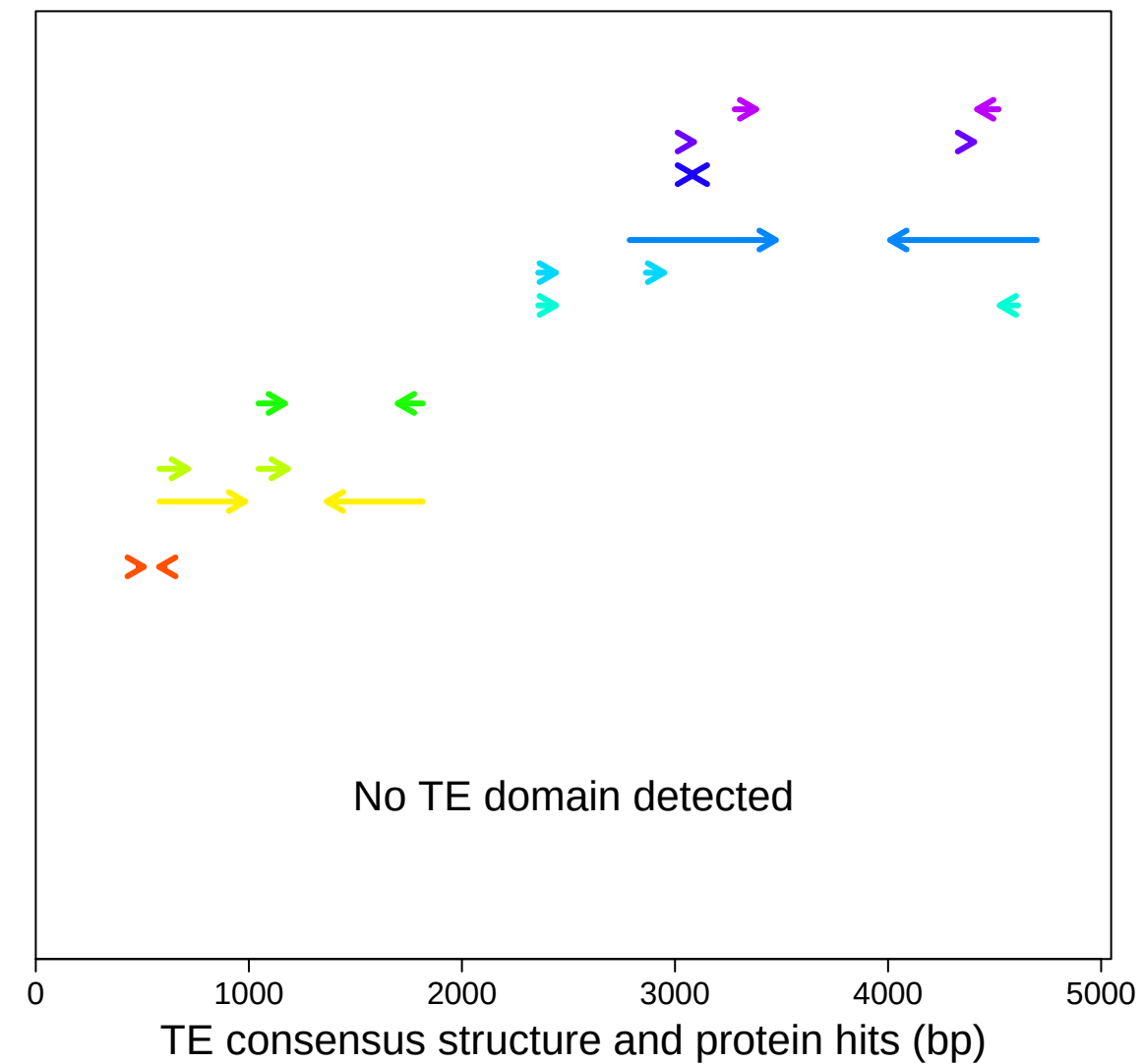
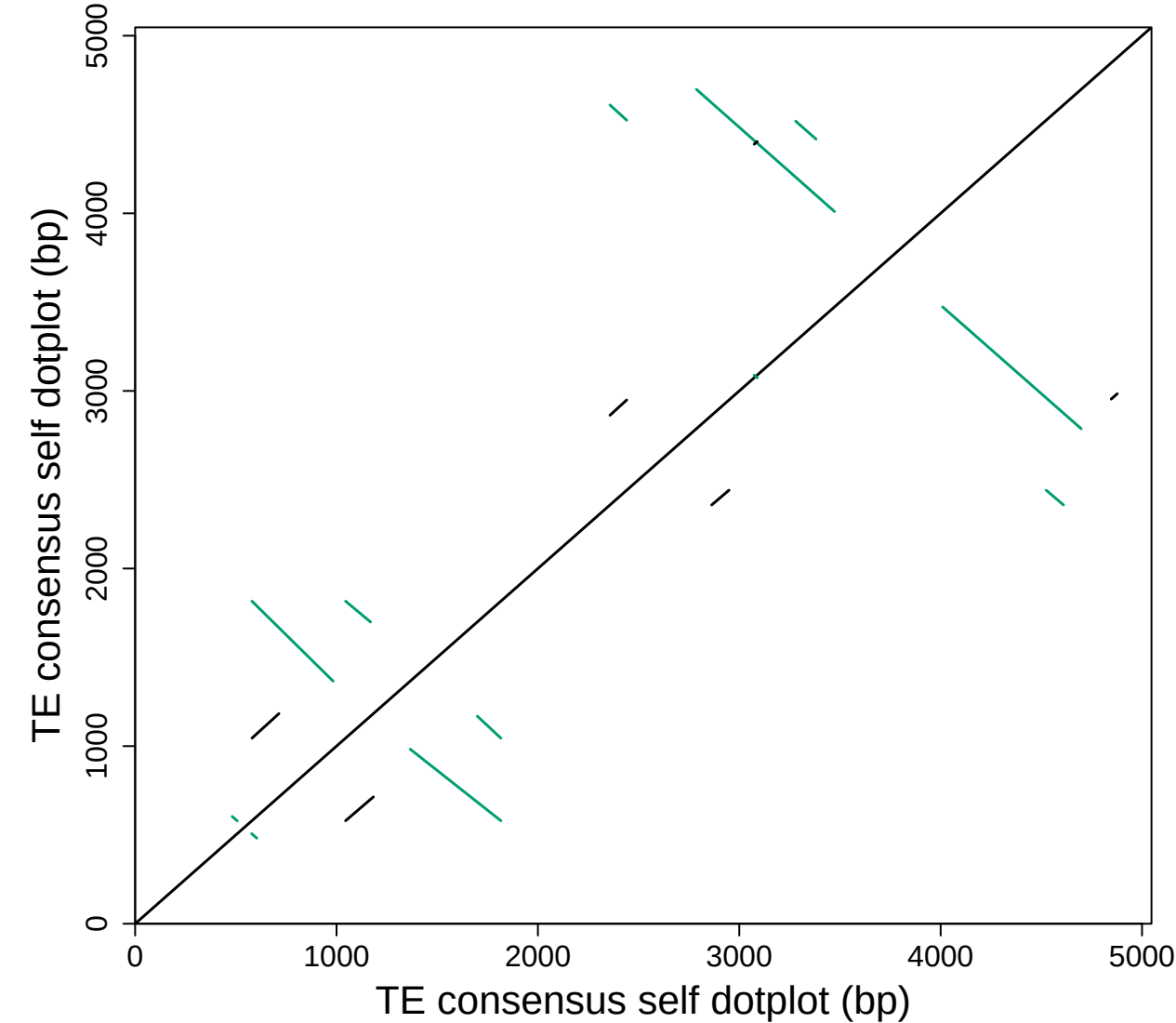
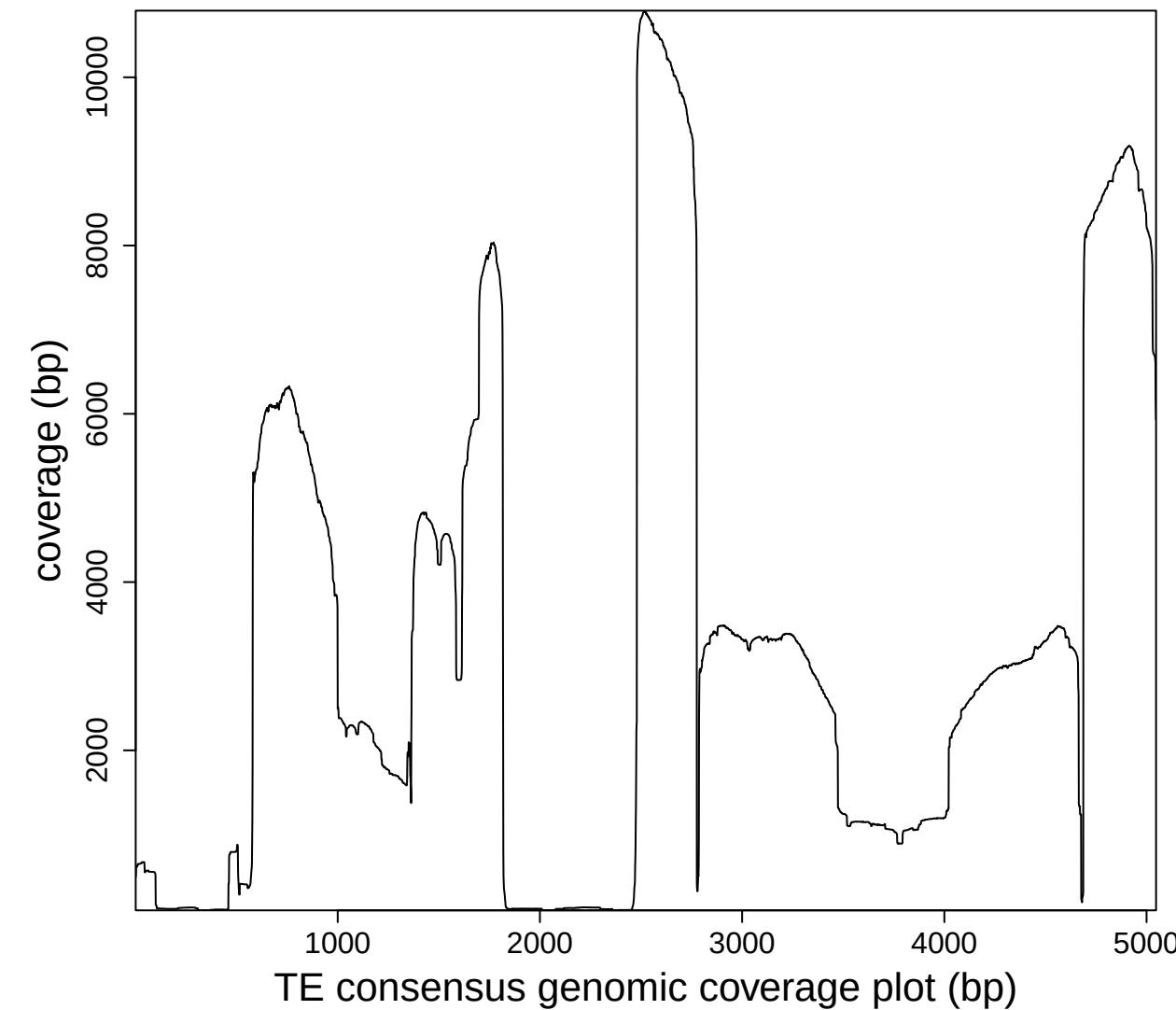
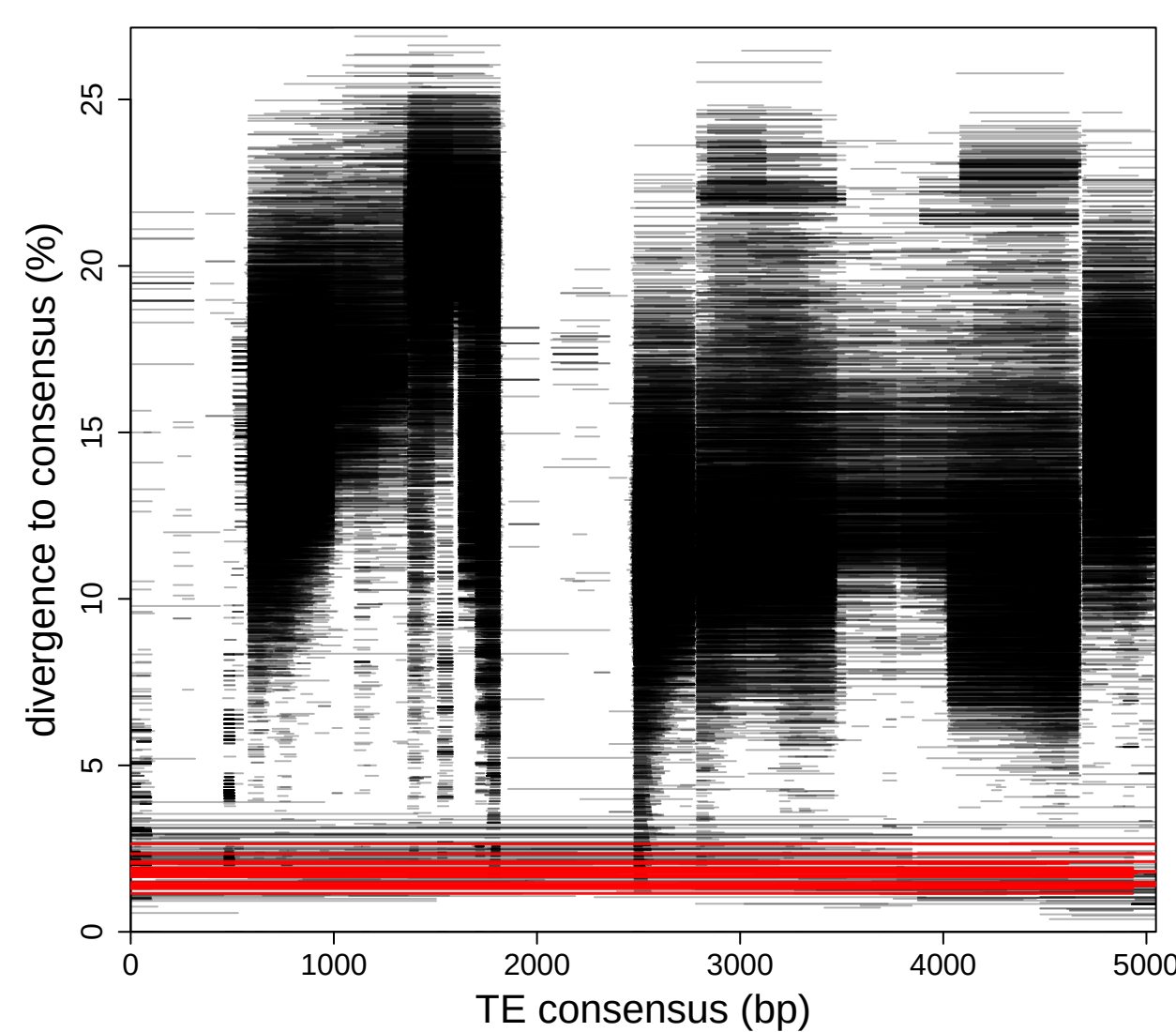
9.fasta.b.bed uf.bed g_1.bed fm_1.bed 0 0 bcln.fa aln.fa cl.f
size: 8804bp; fragments: 98078; full length: 0 (>=7923.6bp)

After TEtrimmer Extended plot Blue lines are boundaries

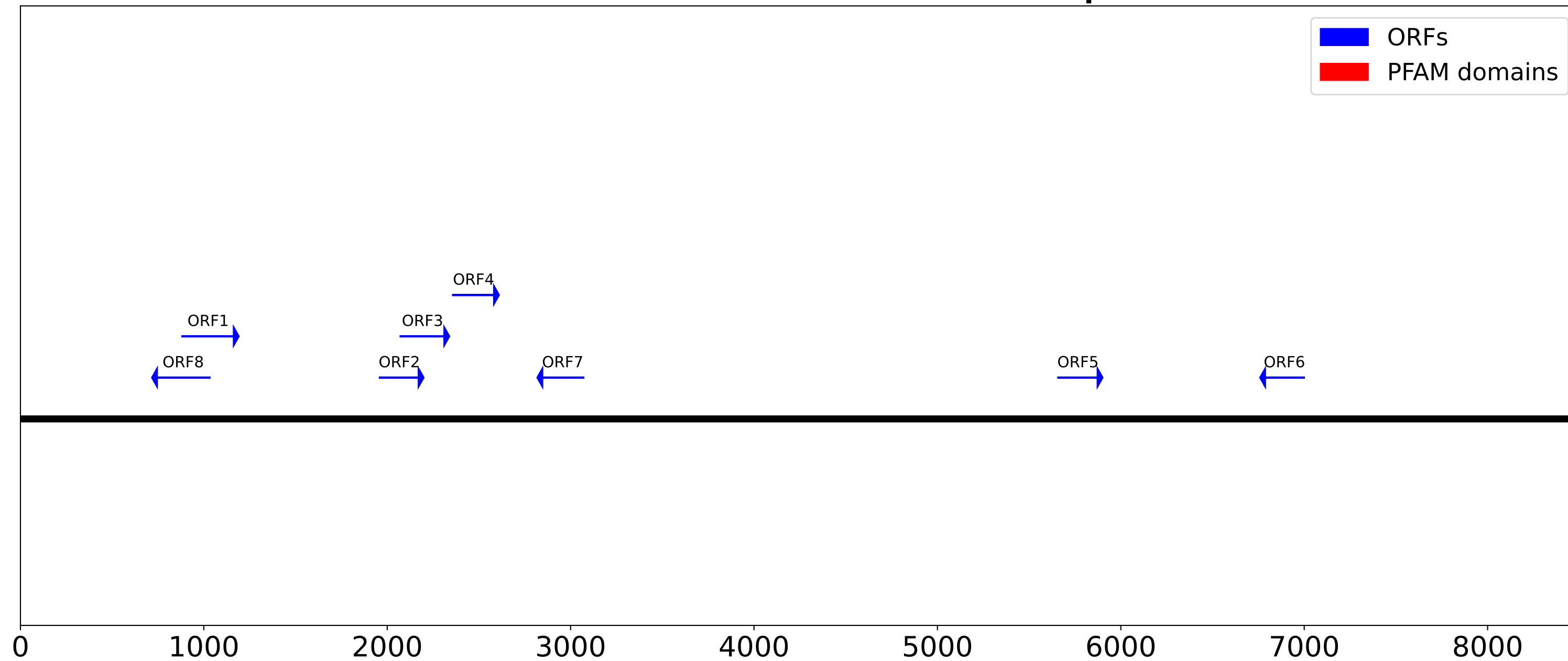


TE: rnd_5_family_3349
size: 5047bp; fragments: 48908; full length: 24 (≥ 4542.3 bp)

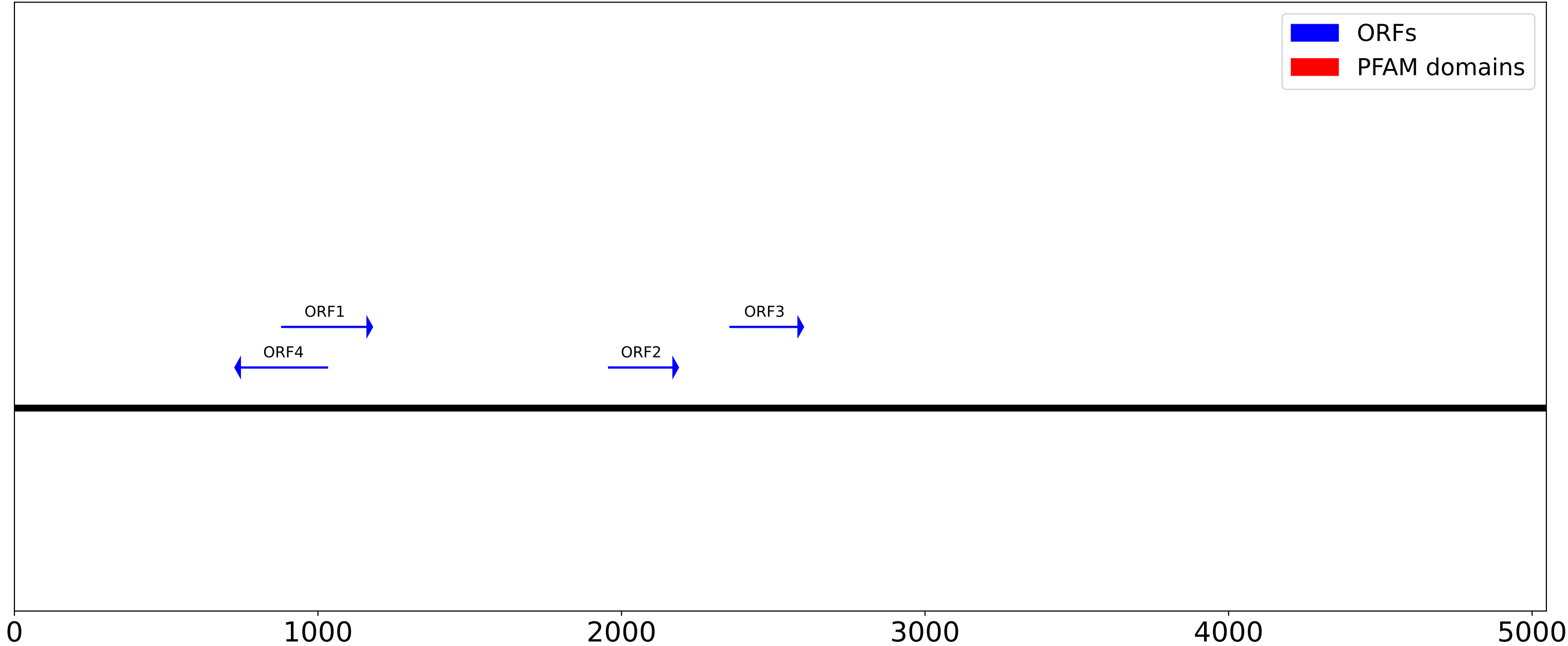
Before TEtrimmer 5047 bp



After TEtrimmer ORF and PFAM domain plot



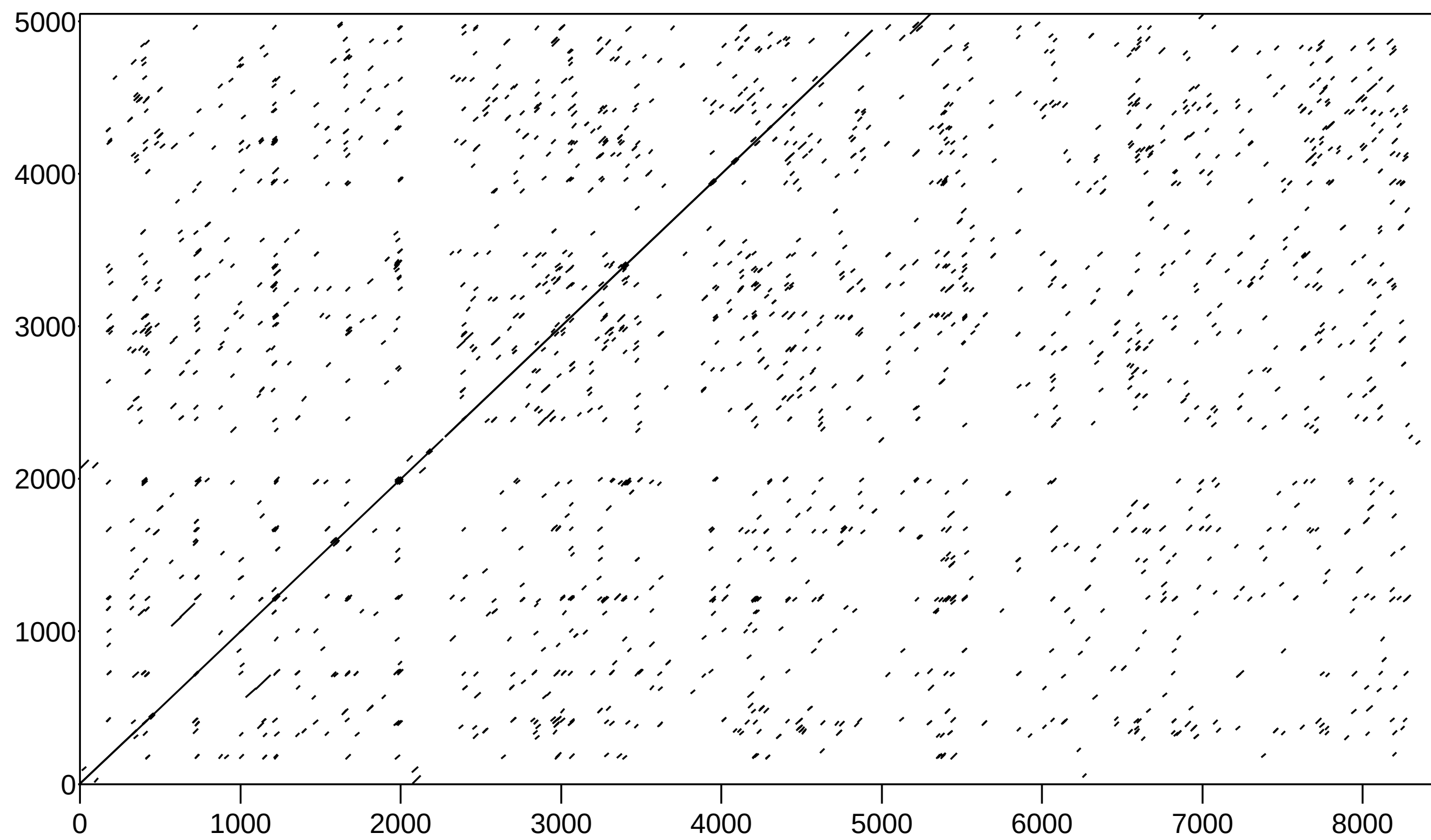
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)