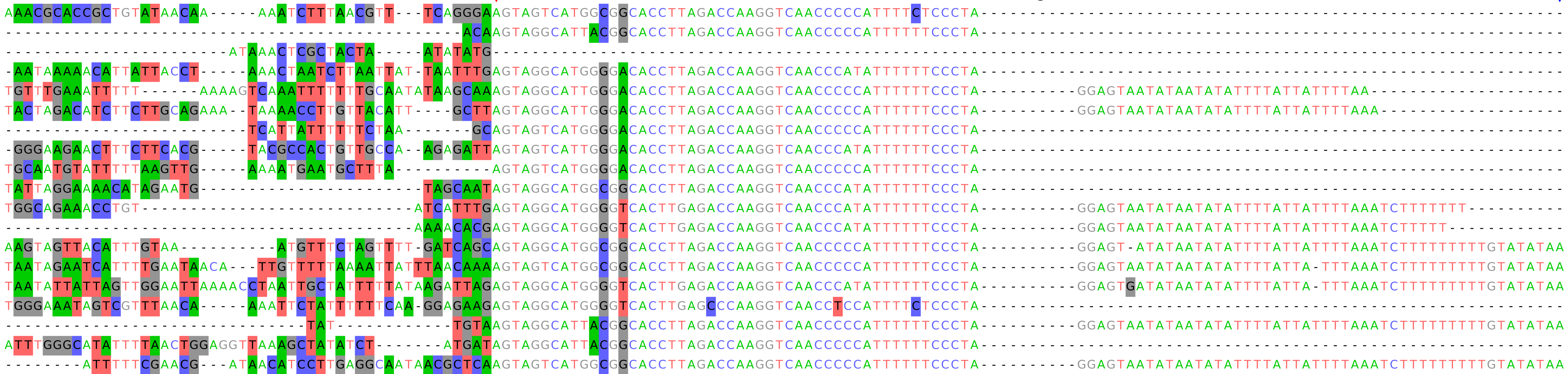
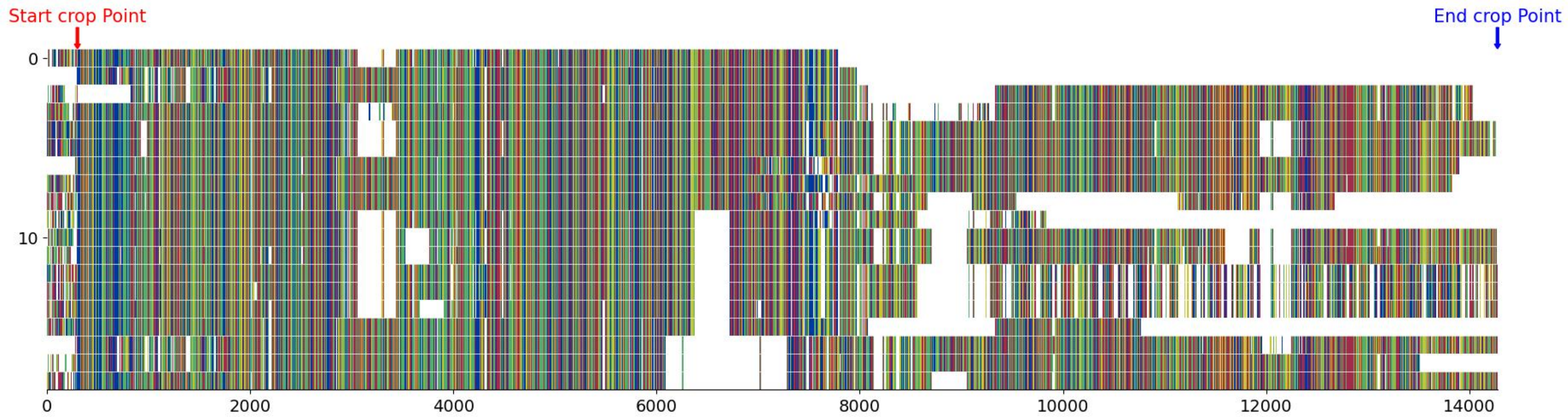


Start crop Point

End crop Point

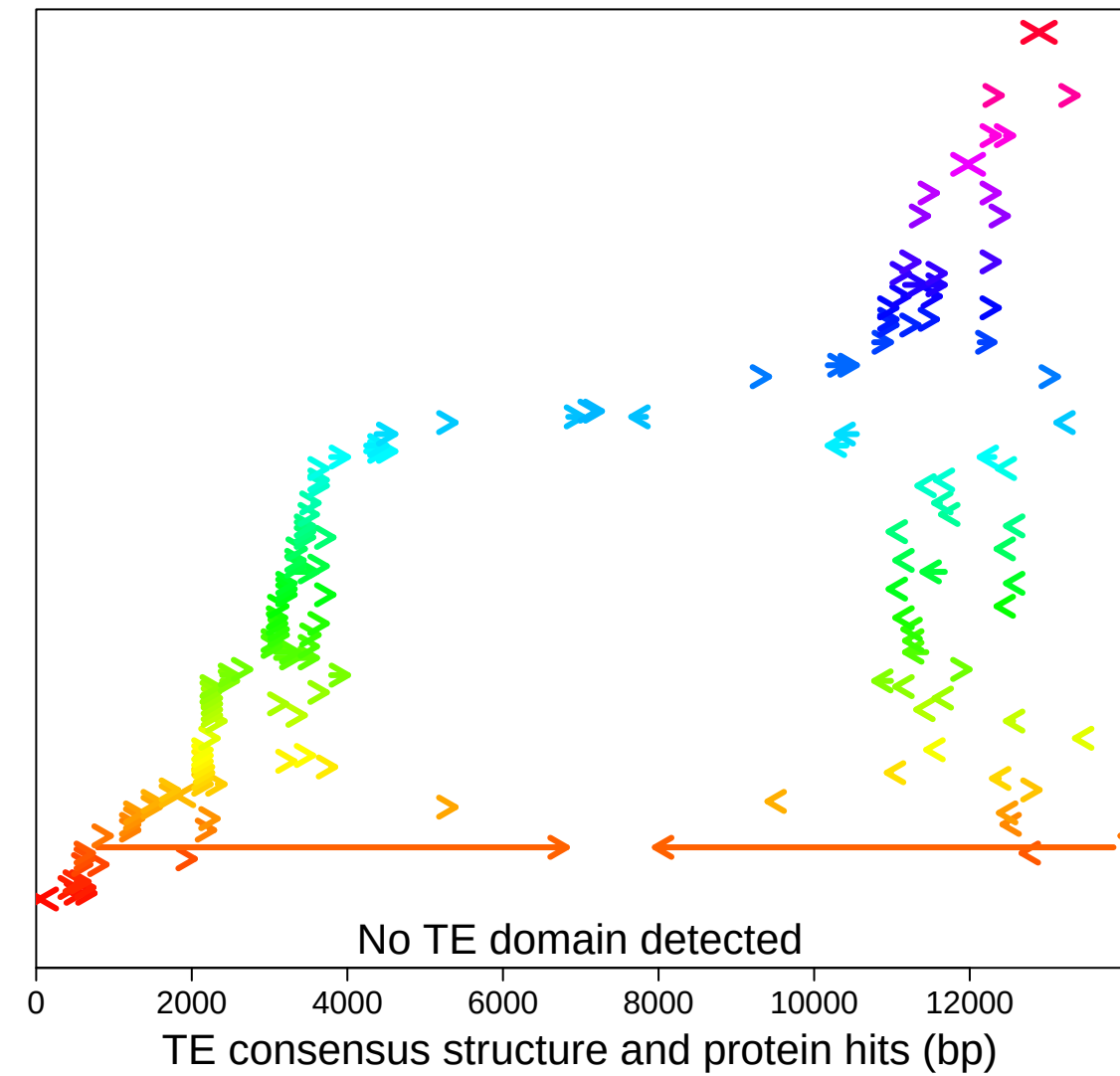
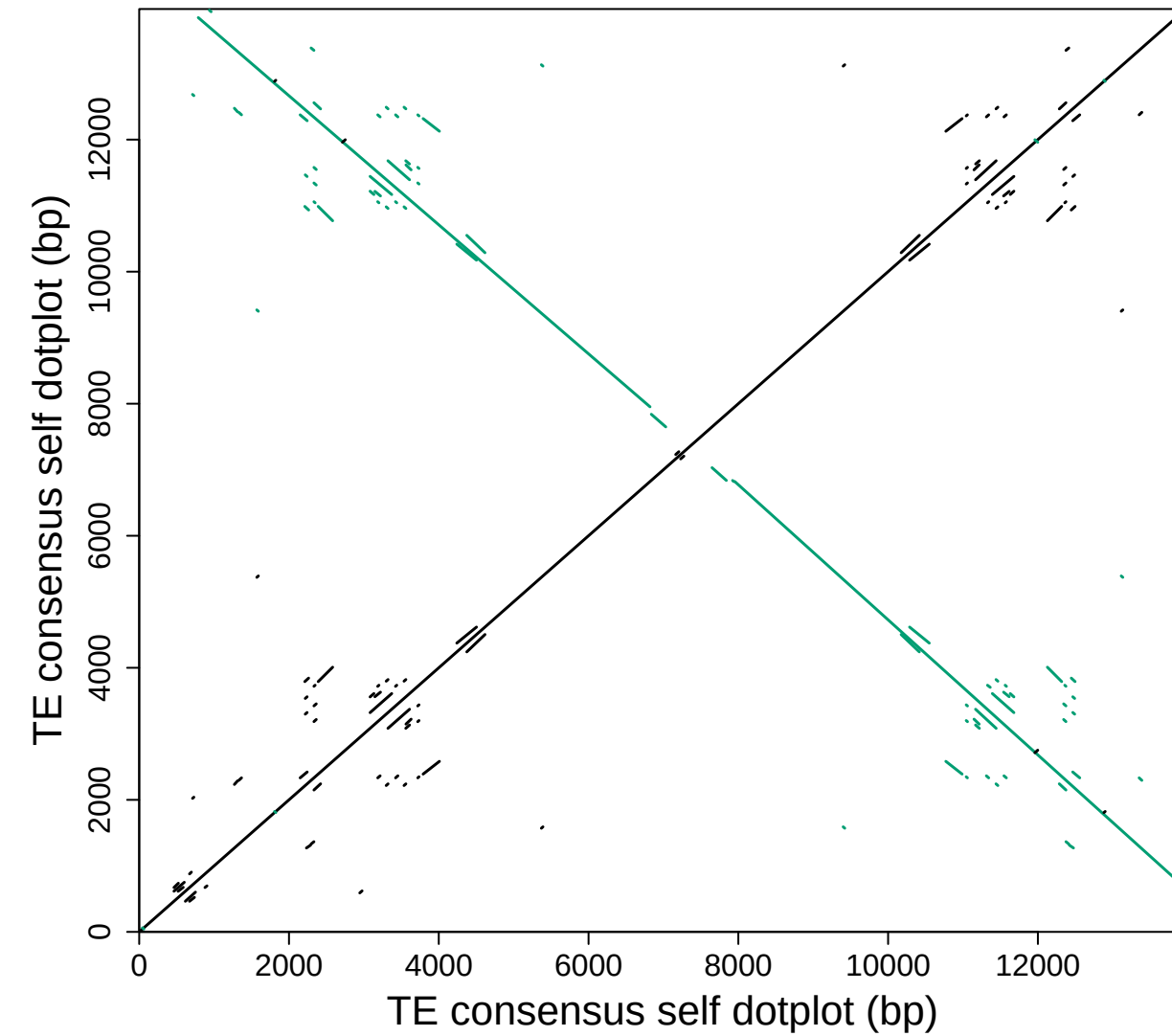
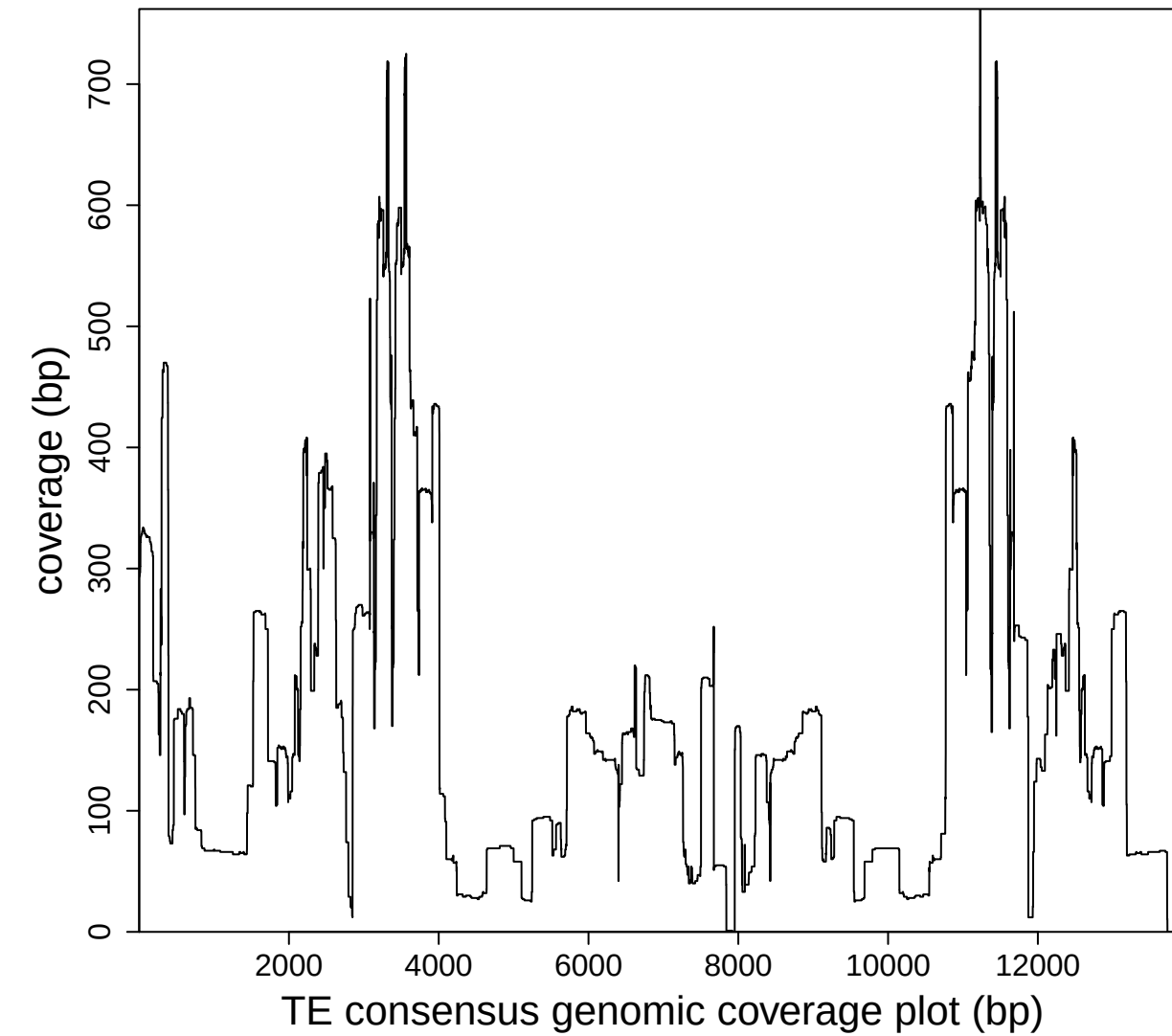
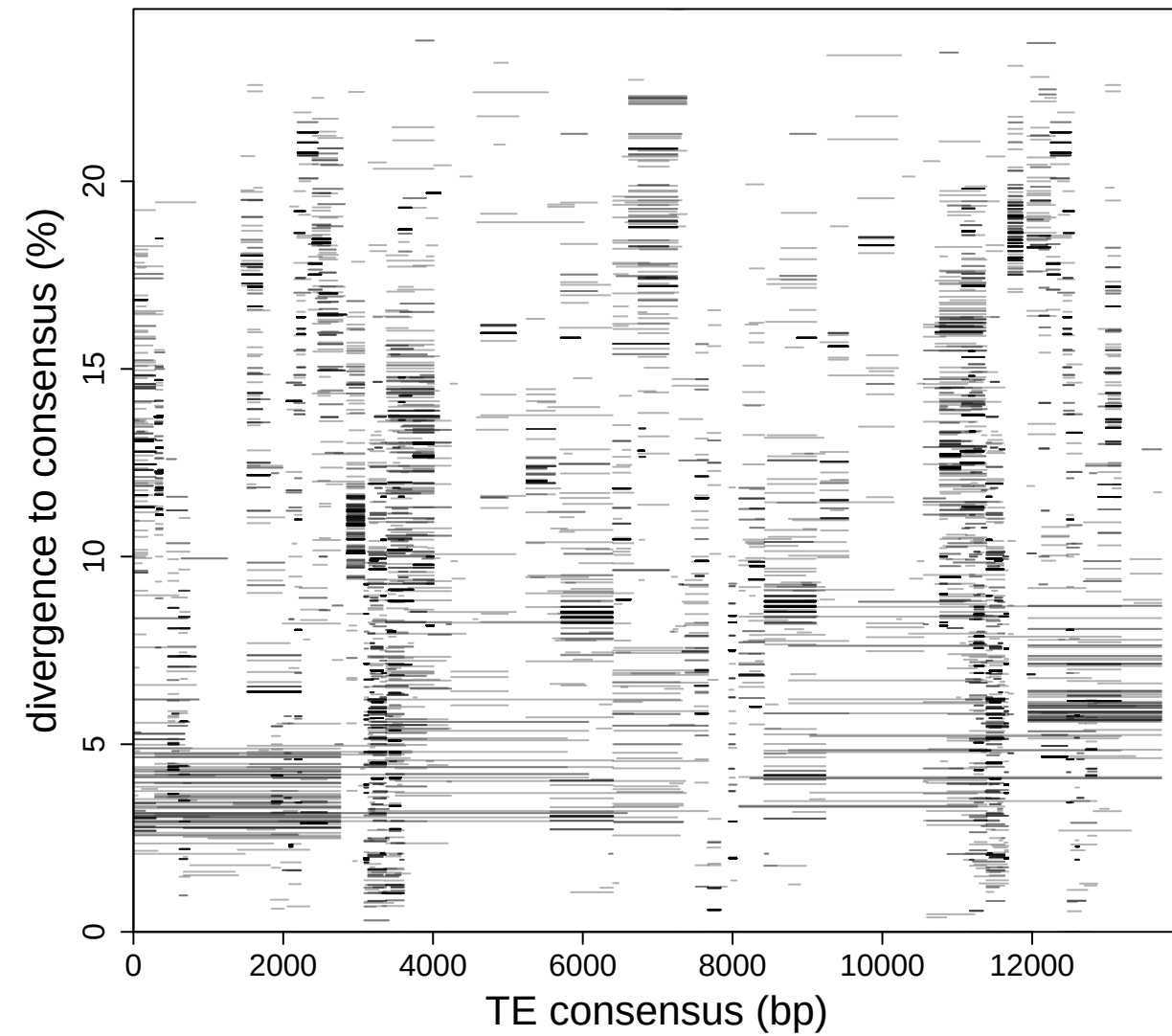
MSA length = 13978





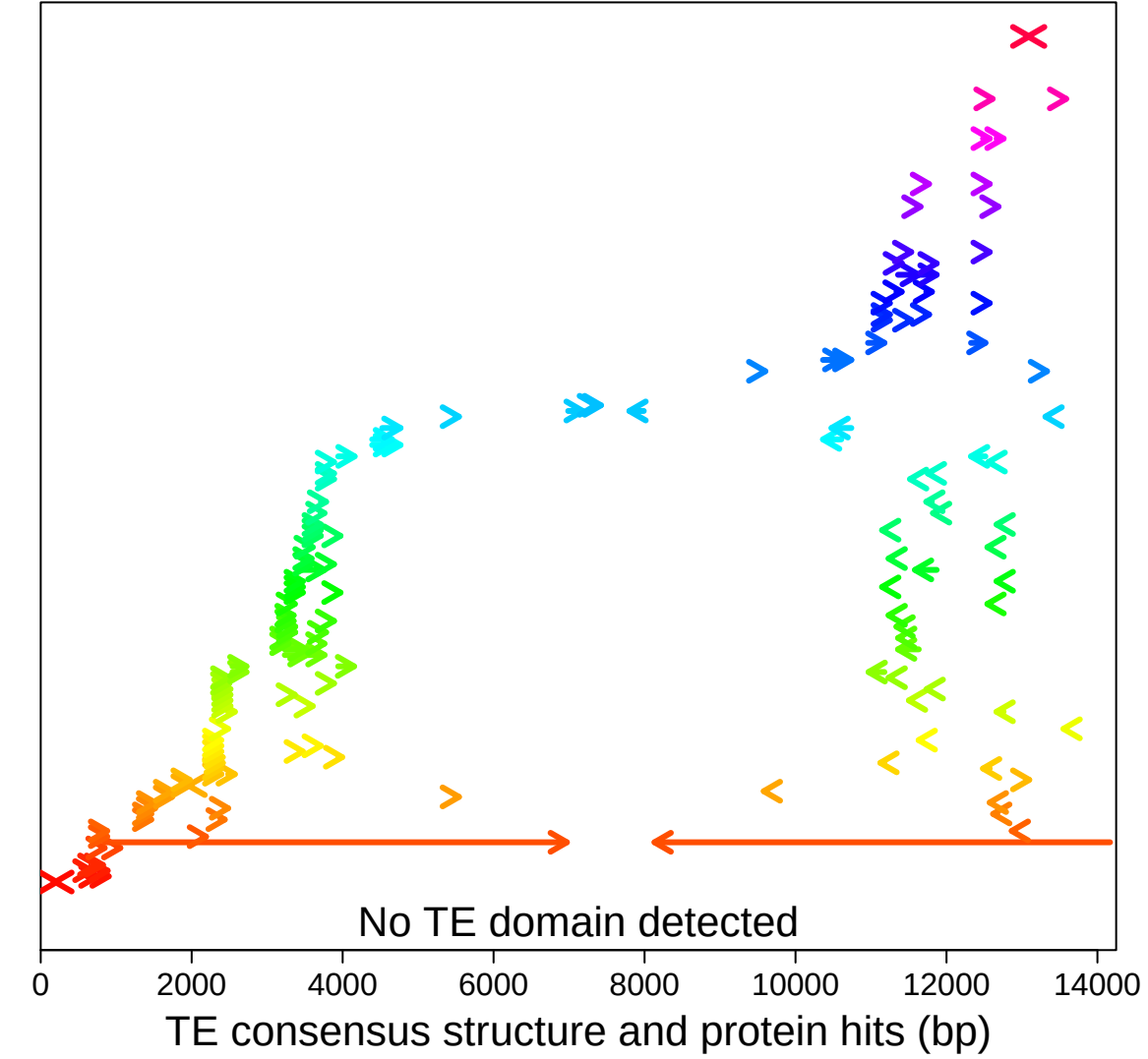
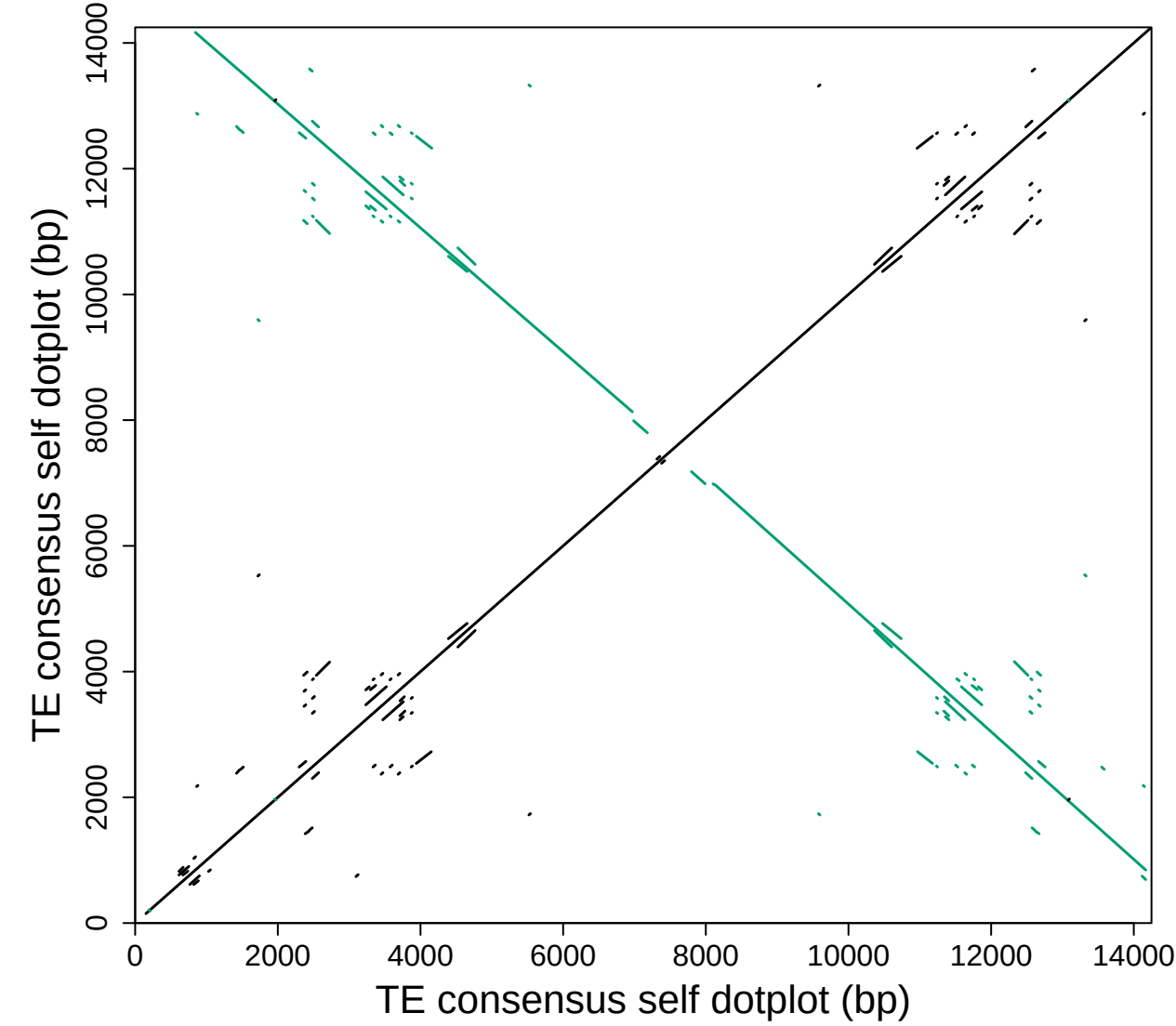
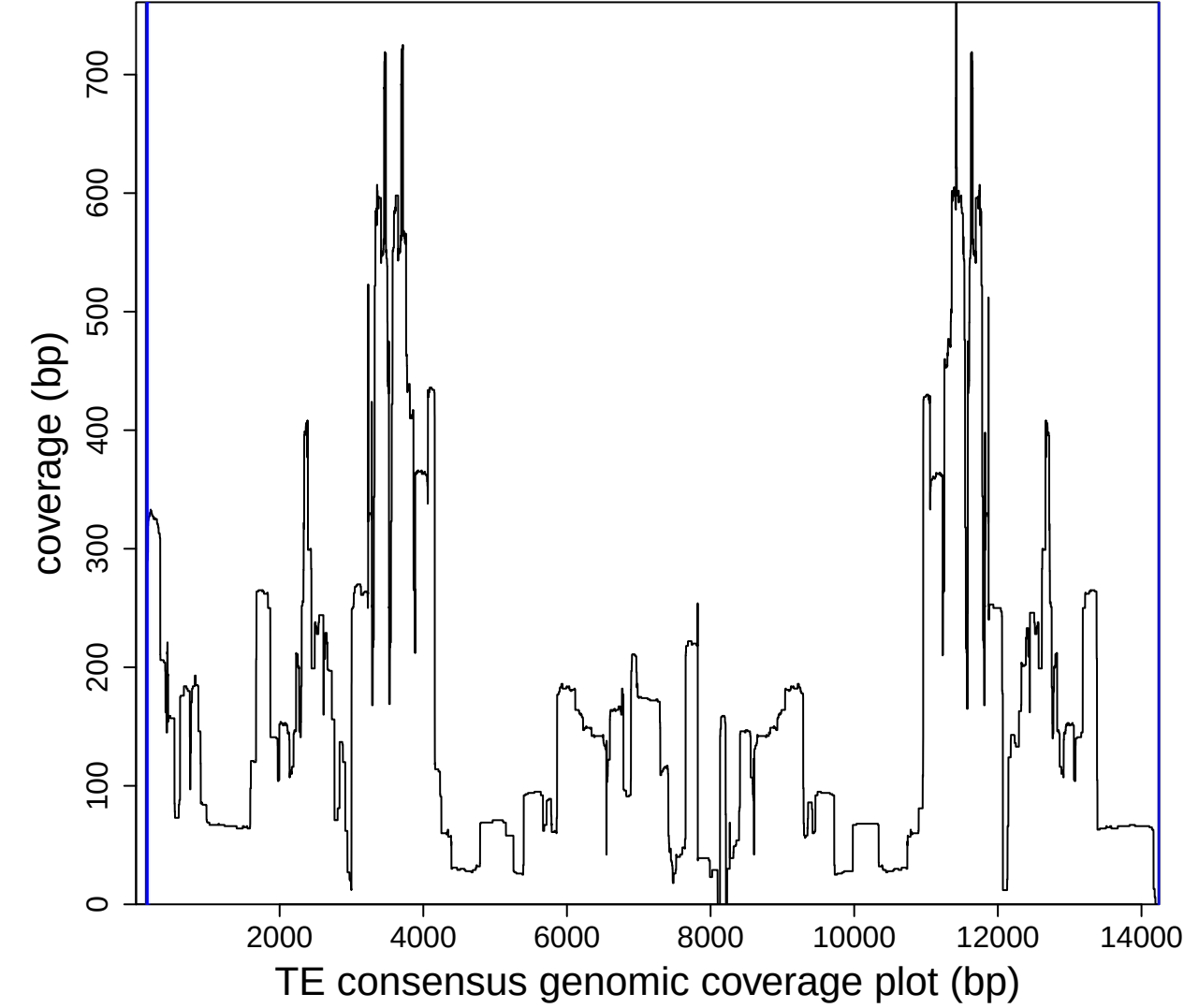
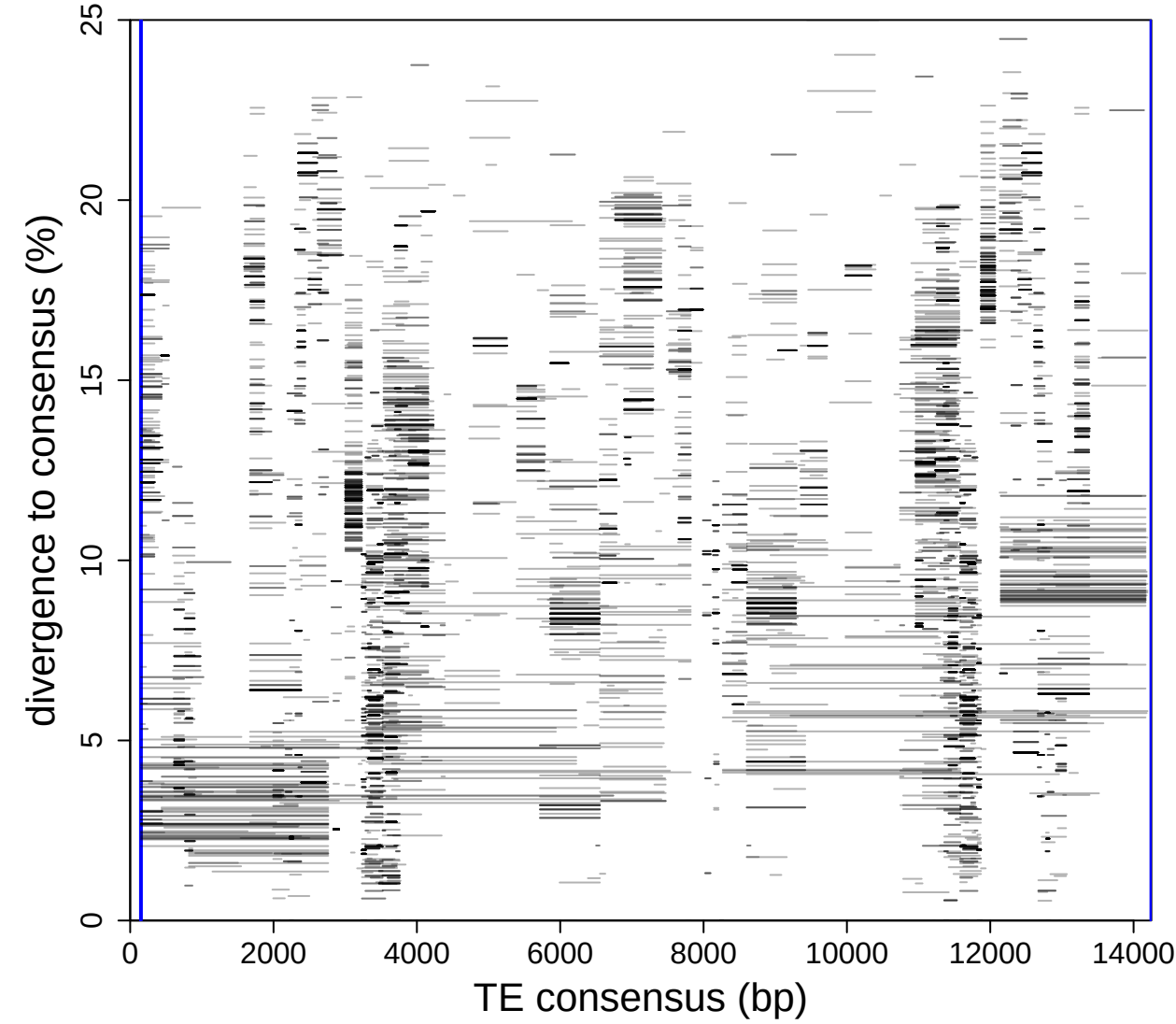
a.b.bed_uf.bed_g_2.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.f
size: 13978bp; fragments: 9161; full length: 0 (>=12580.2bp)

After TEtrimmer 13978 bp



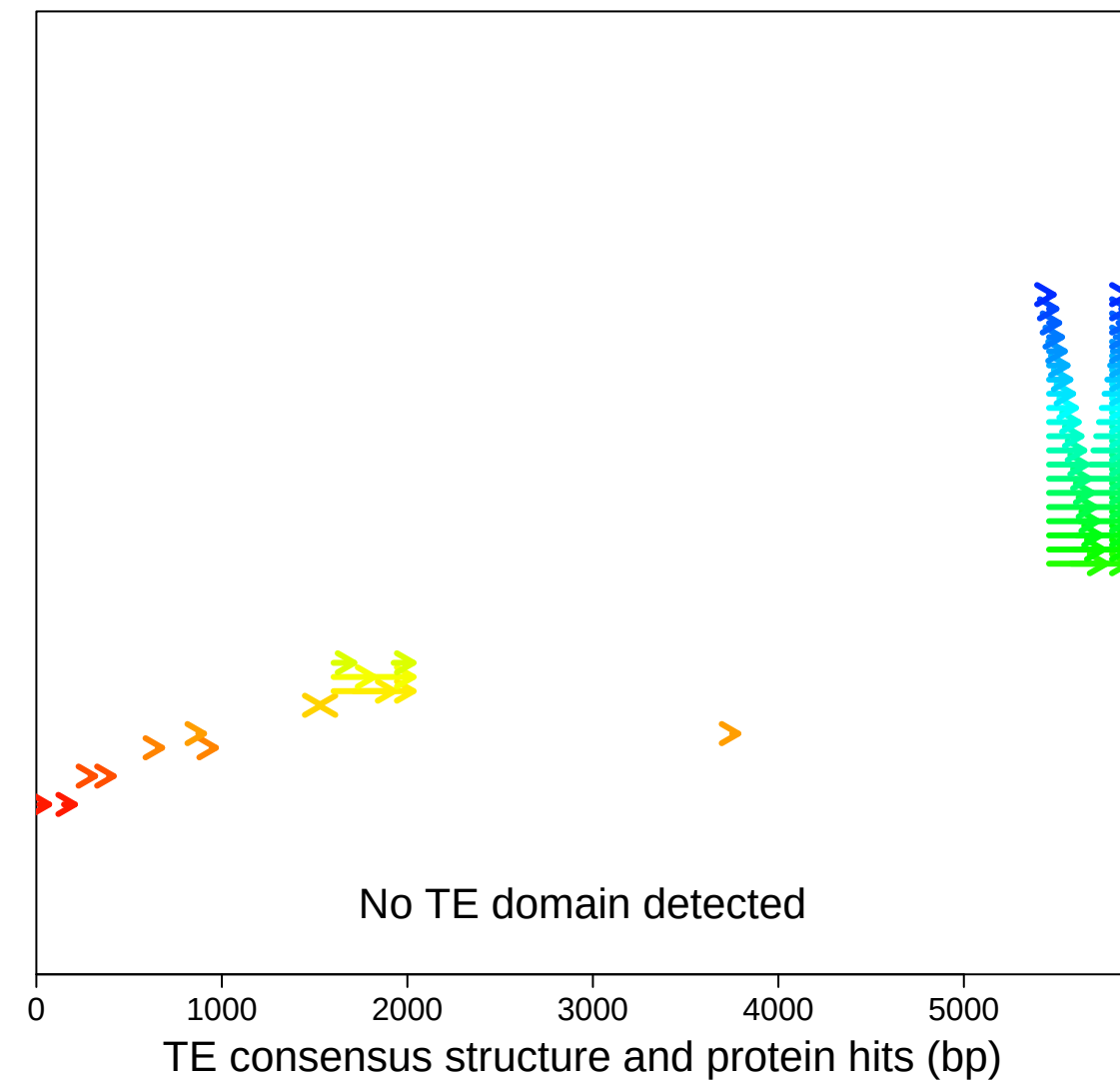
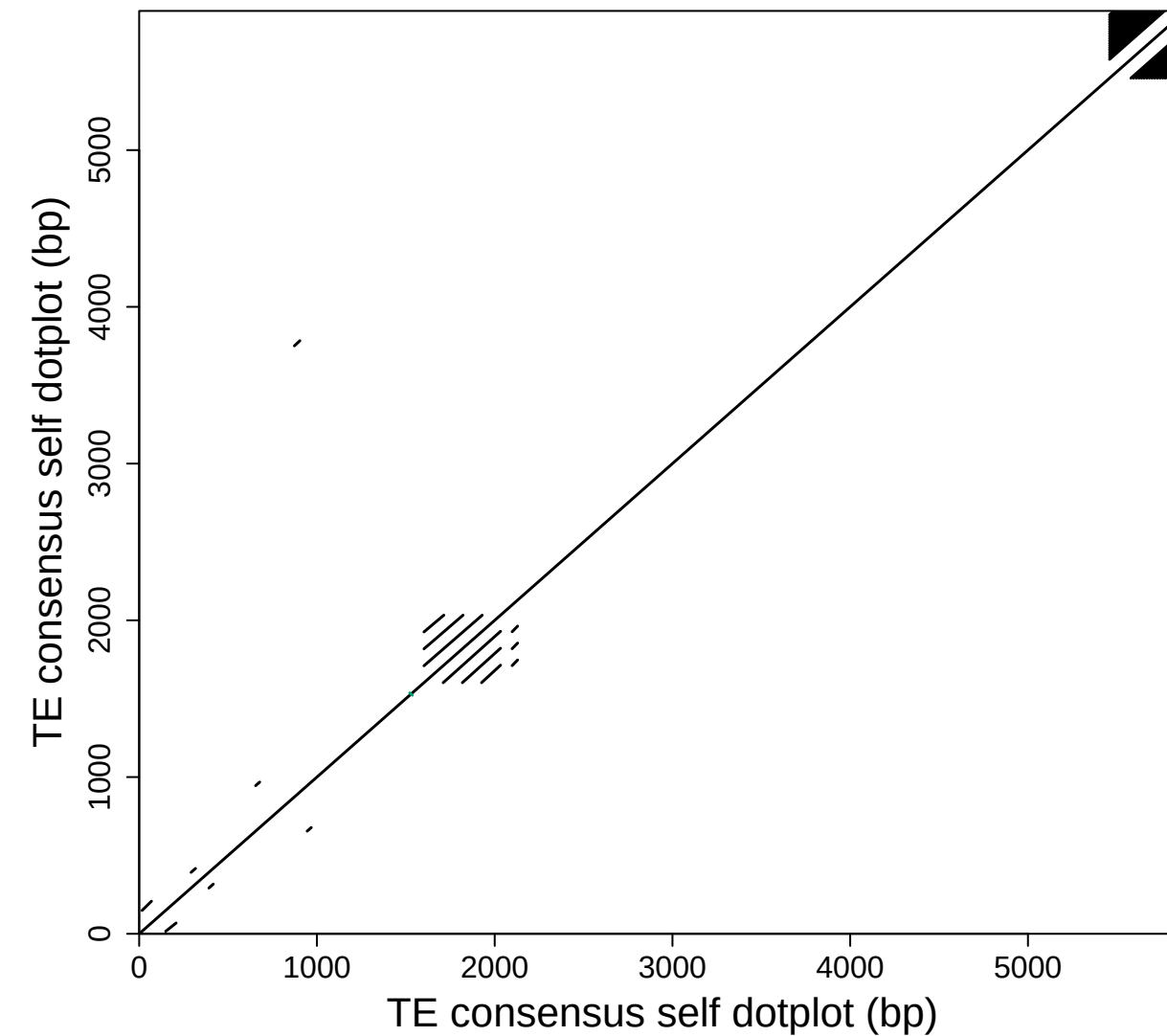
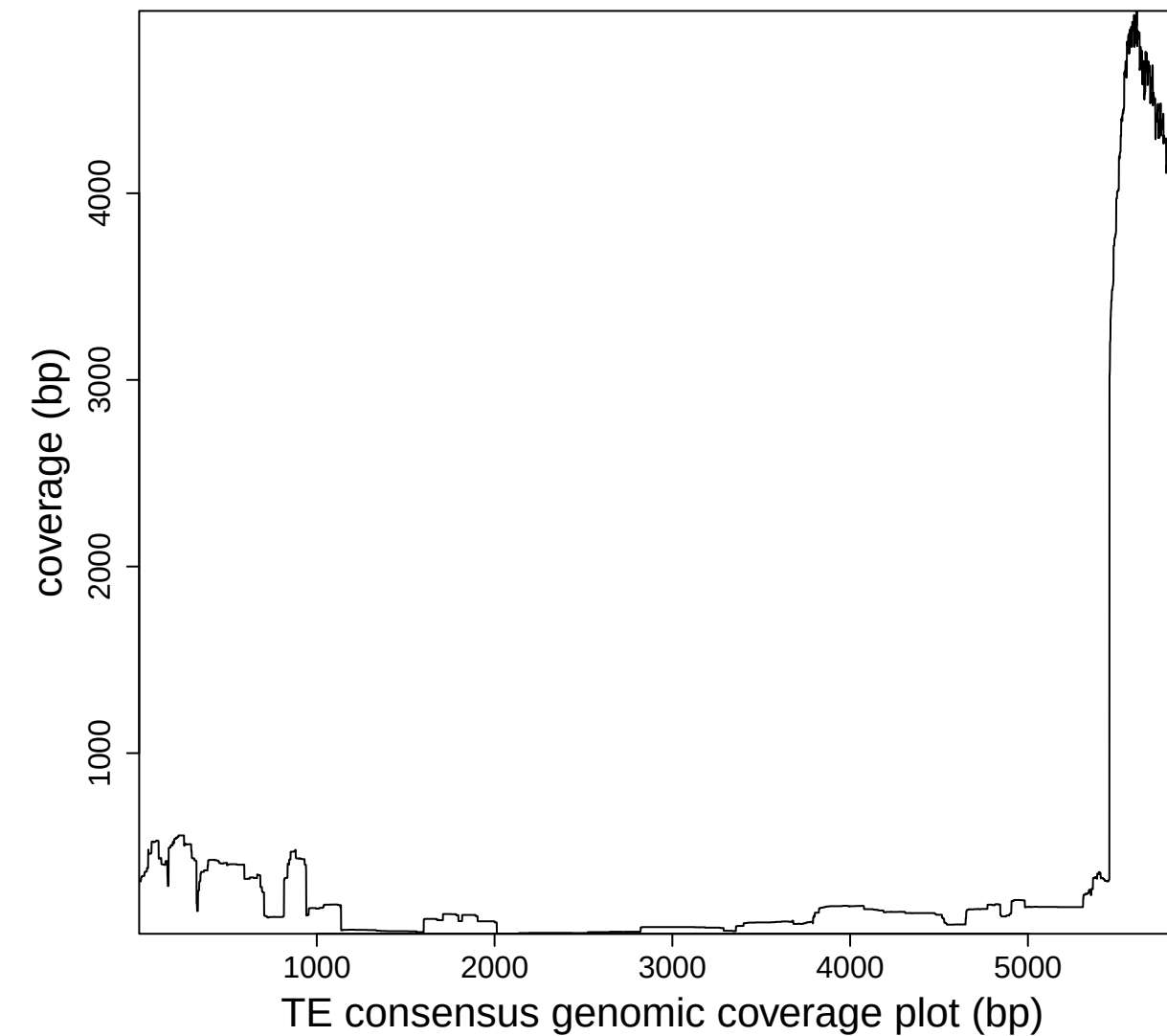
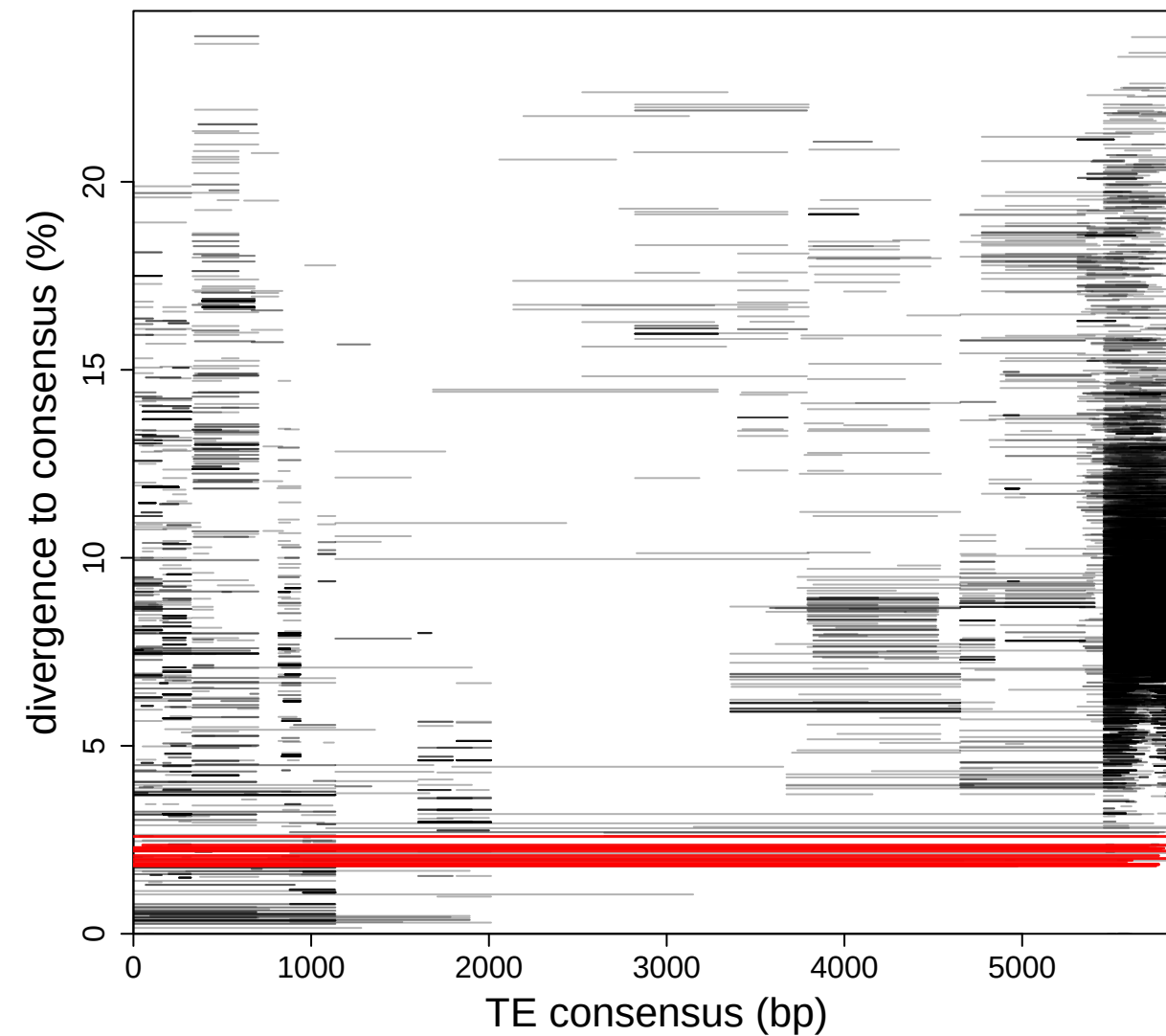
5.fasta.b.bed uf.bed g_2.bed fm_1.bed 0 0 bcln.fa aln.fa cl.f
size: 14248bp; fragments: 8736; full length: 0 (>=12823.2bp)

After TEtrimmer Extended plot Blue lines are boundaries

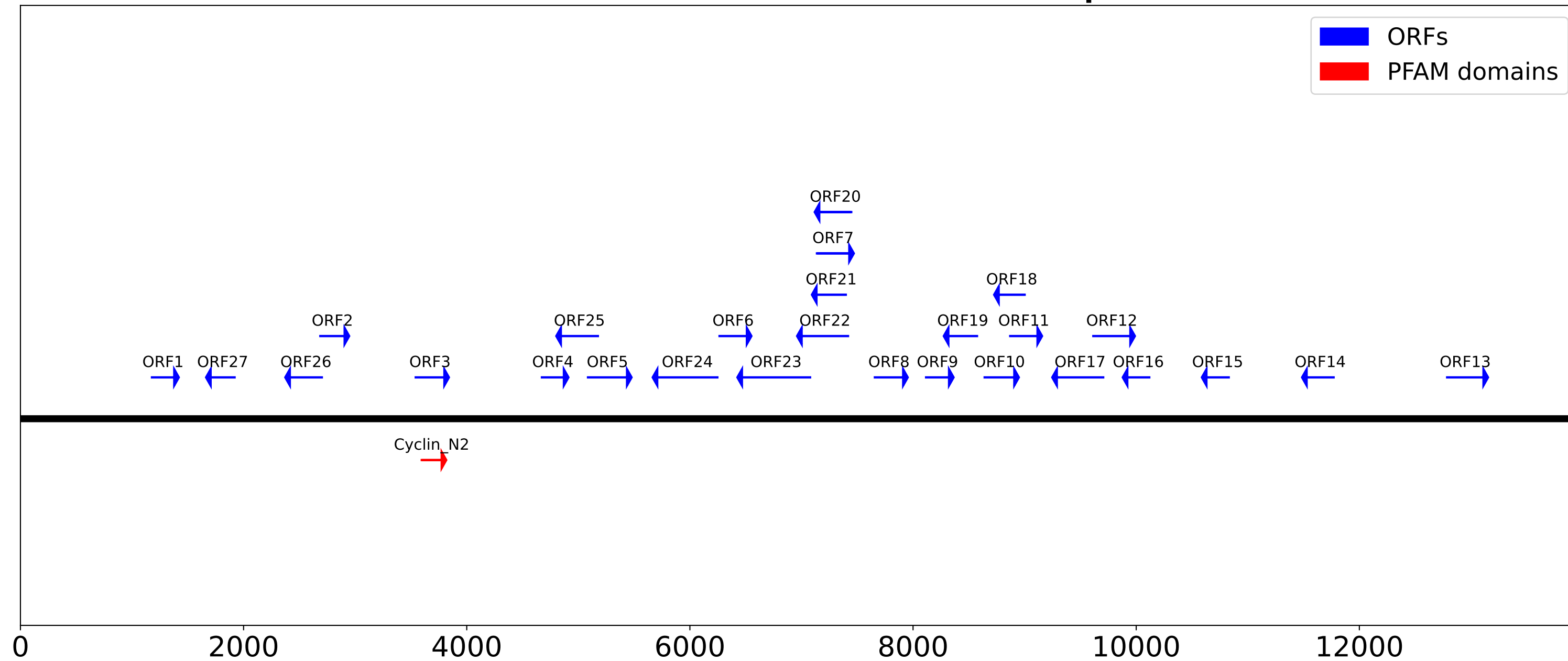


TE: rnd_5_family_7375
size: 5888bp; fragments: 10720; full length: 15 (≥ 5299.2 bp)

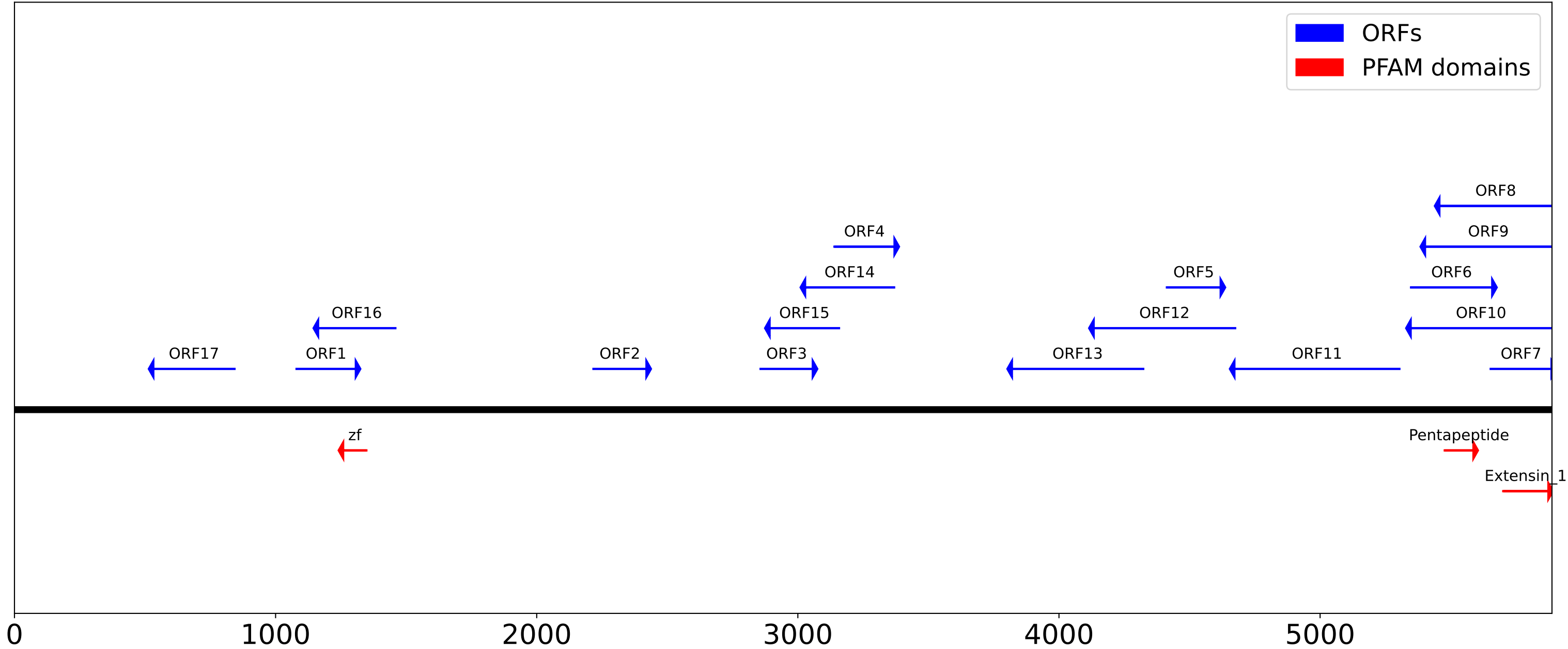
Before TEtrimmer 5888 bp



After TEtrimmer ORF and PFAM domain plot



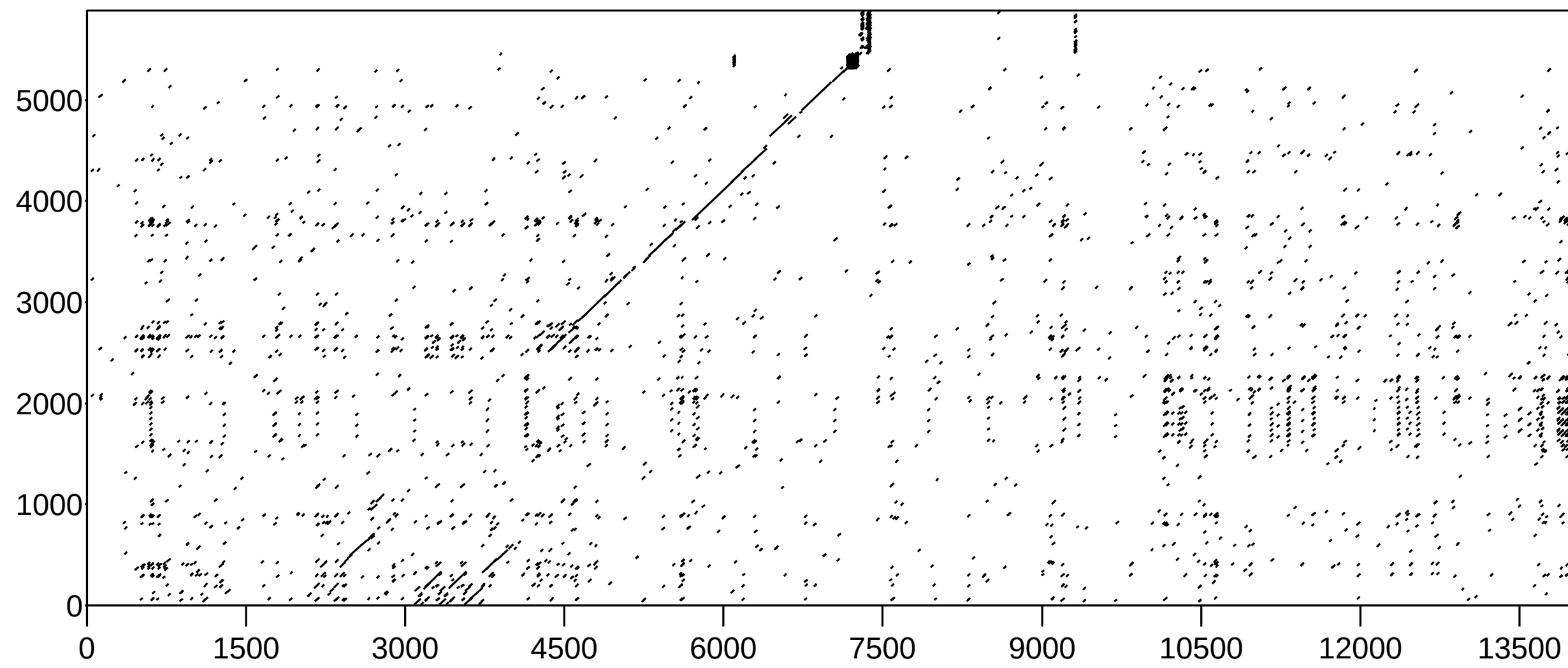
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)