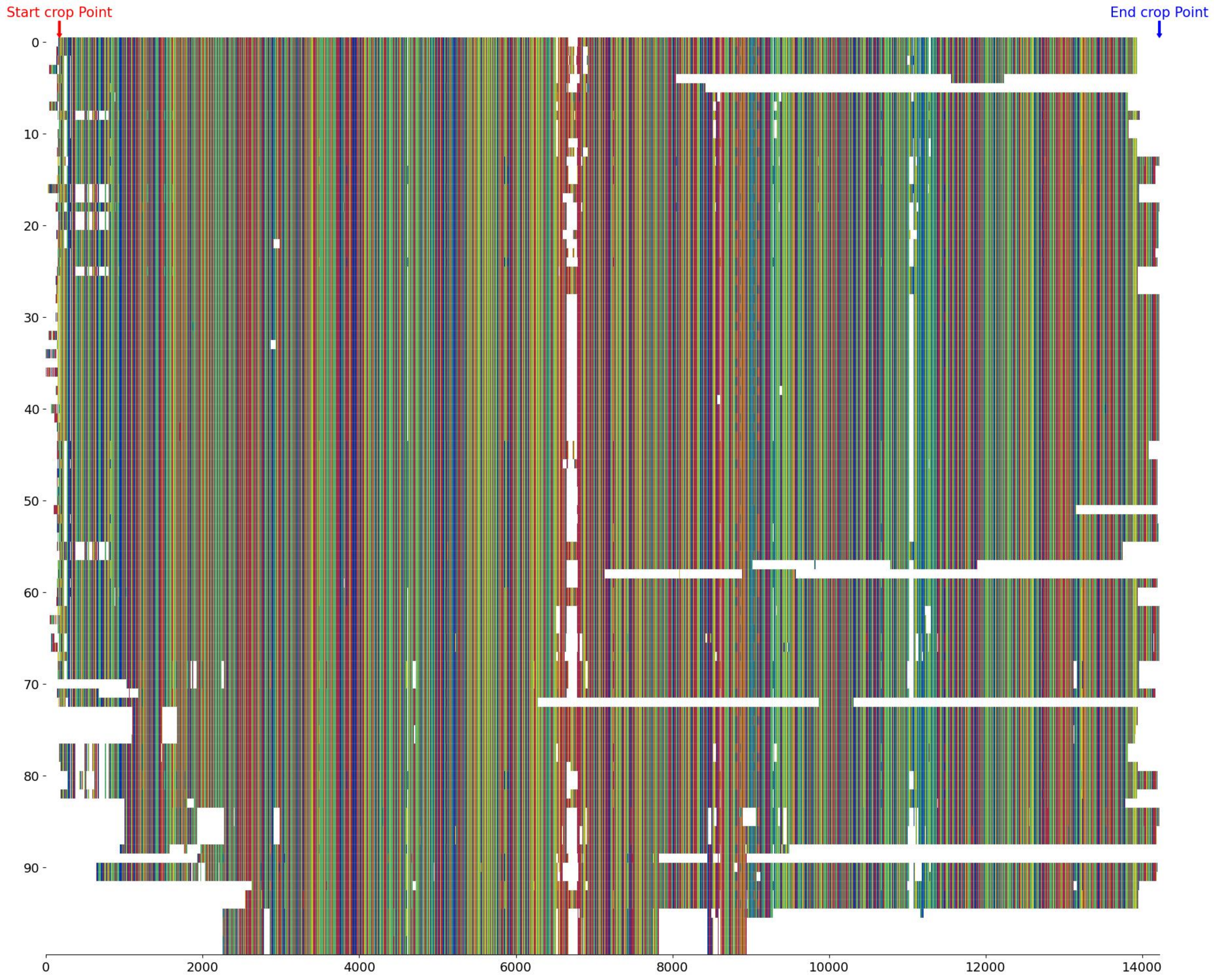


Start crop Point

End crop Point

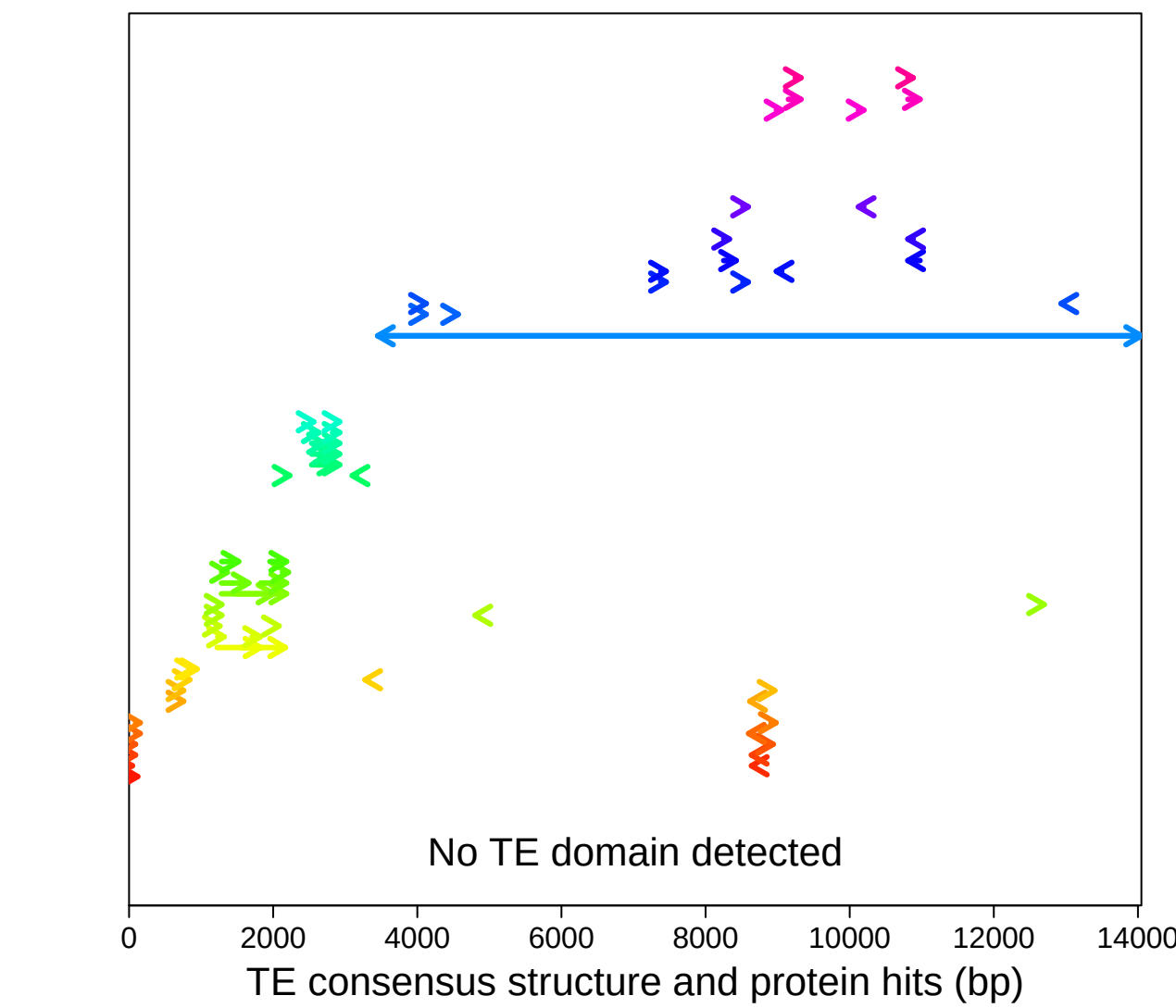
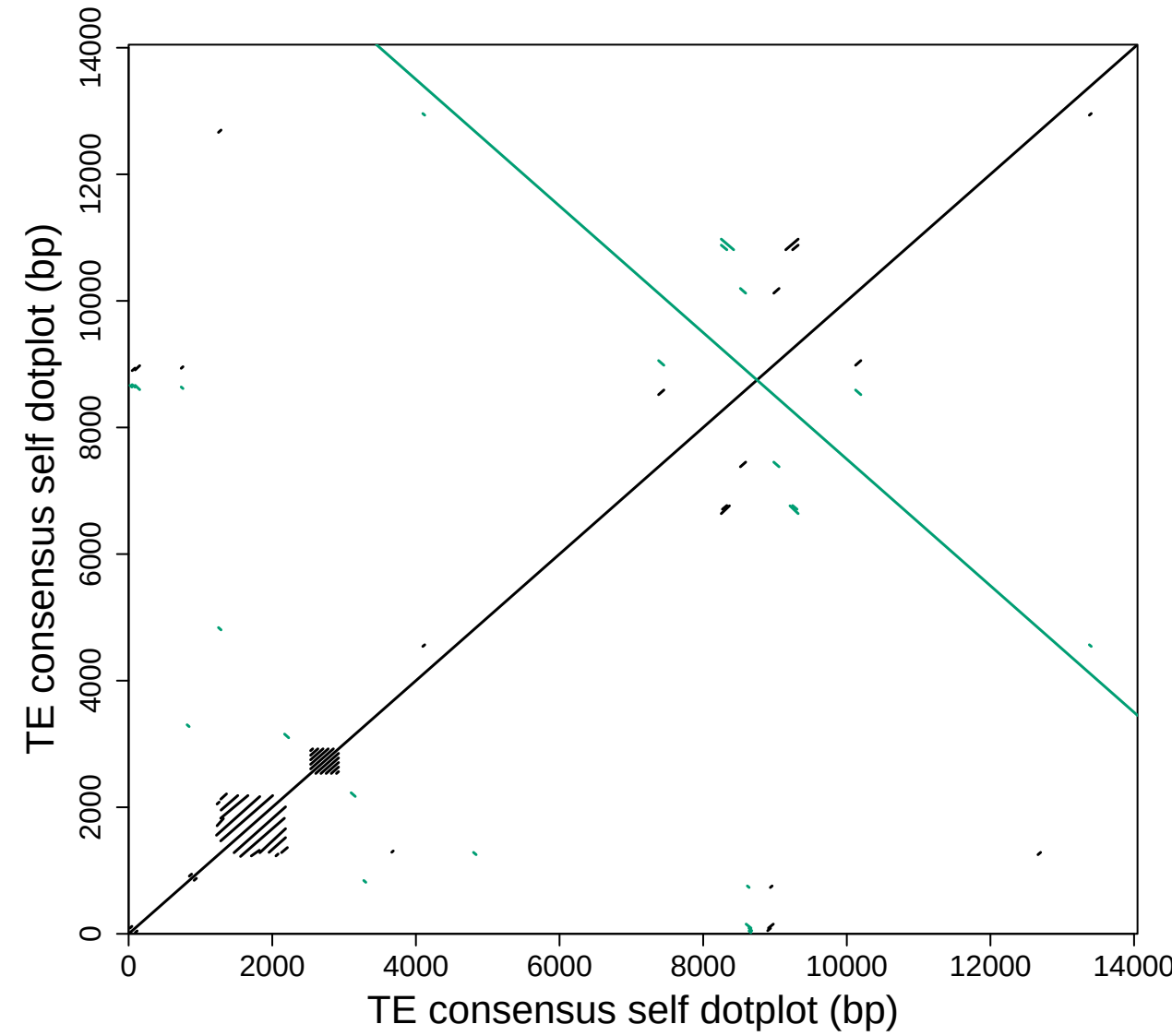
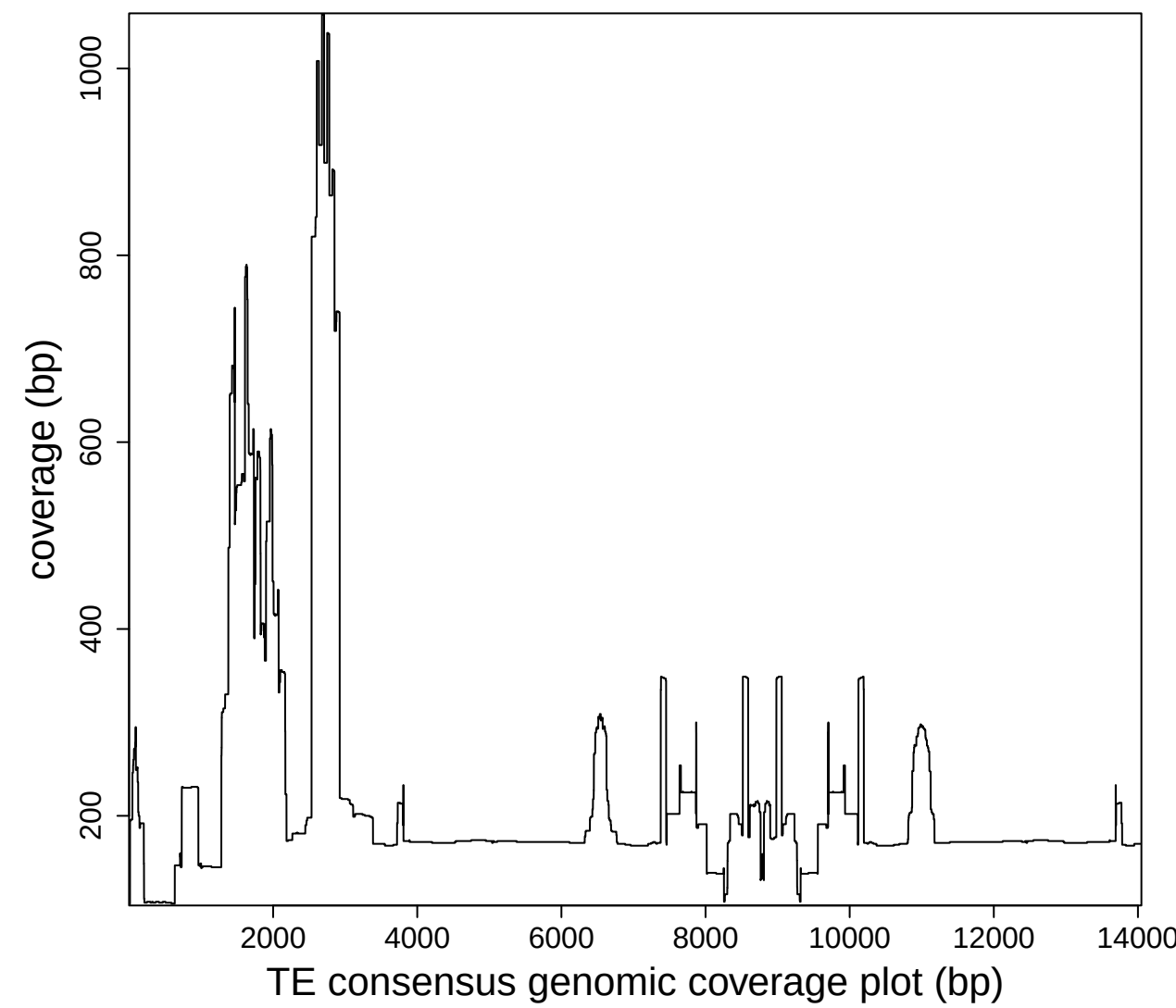
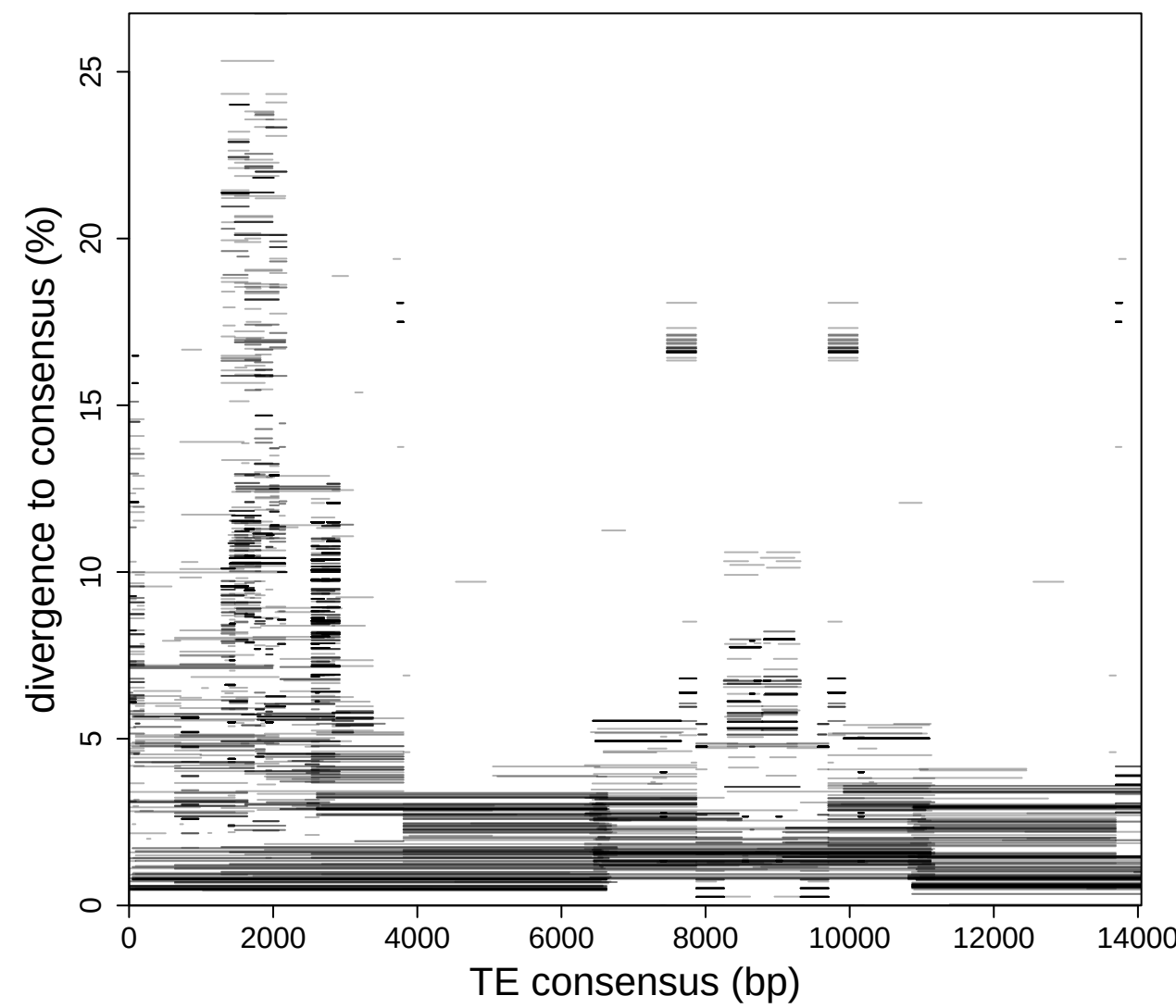
MSA length = 14048





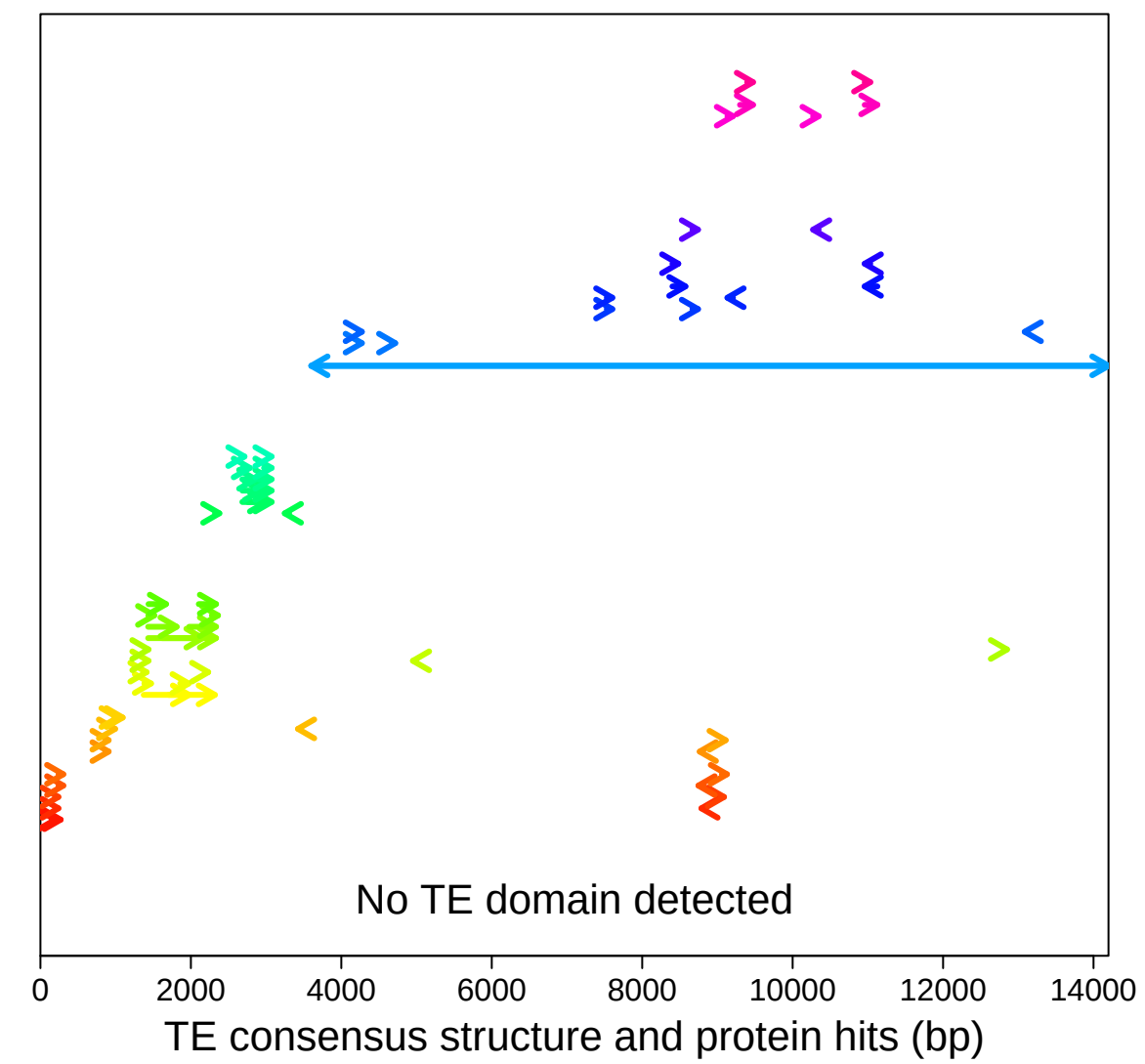
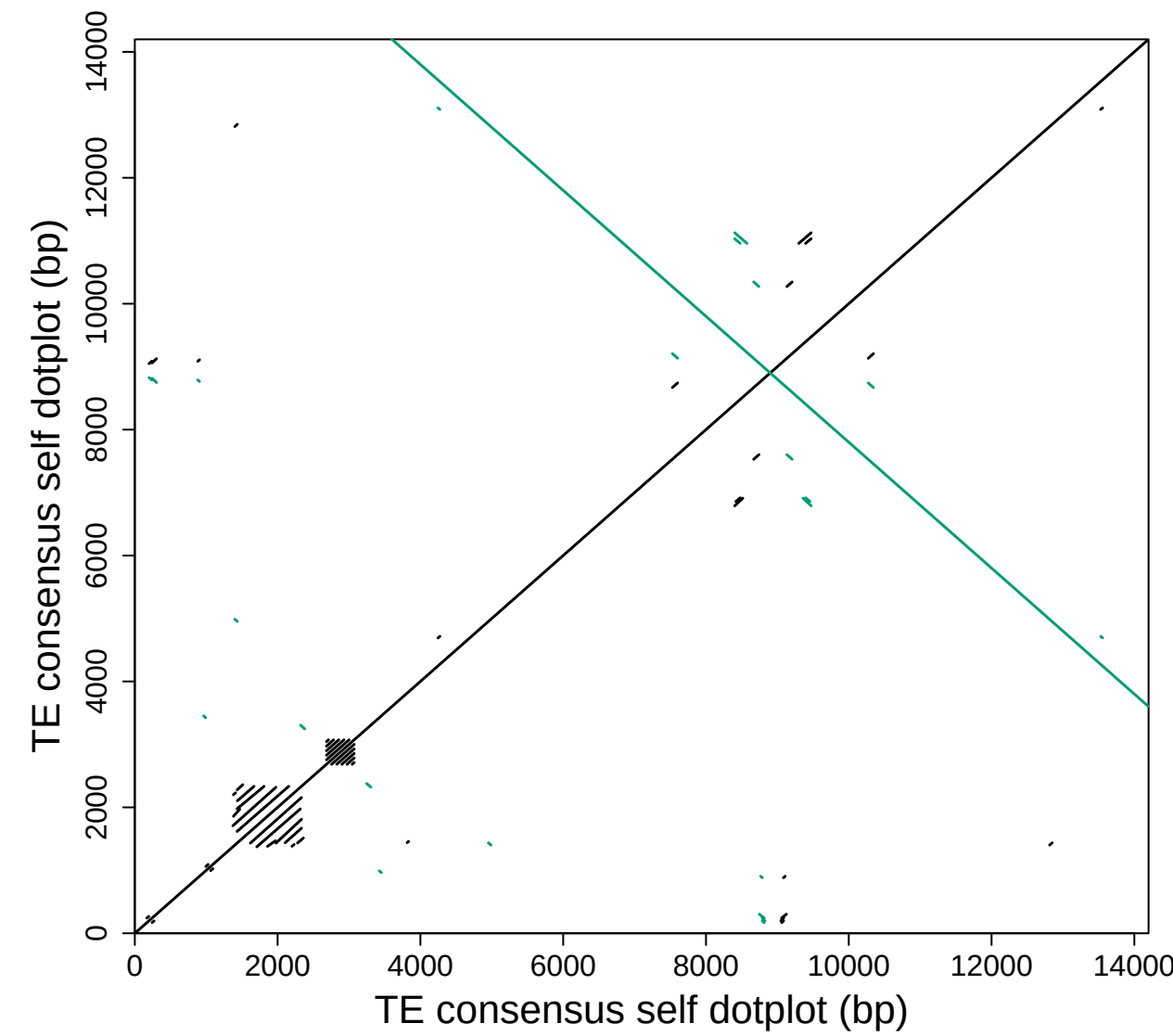
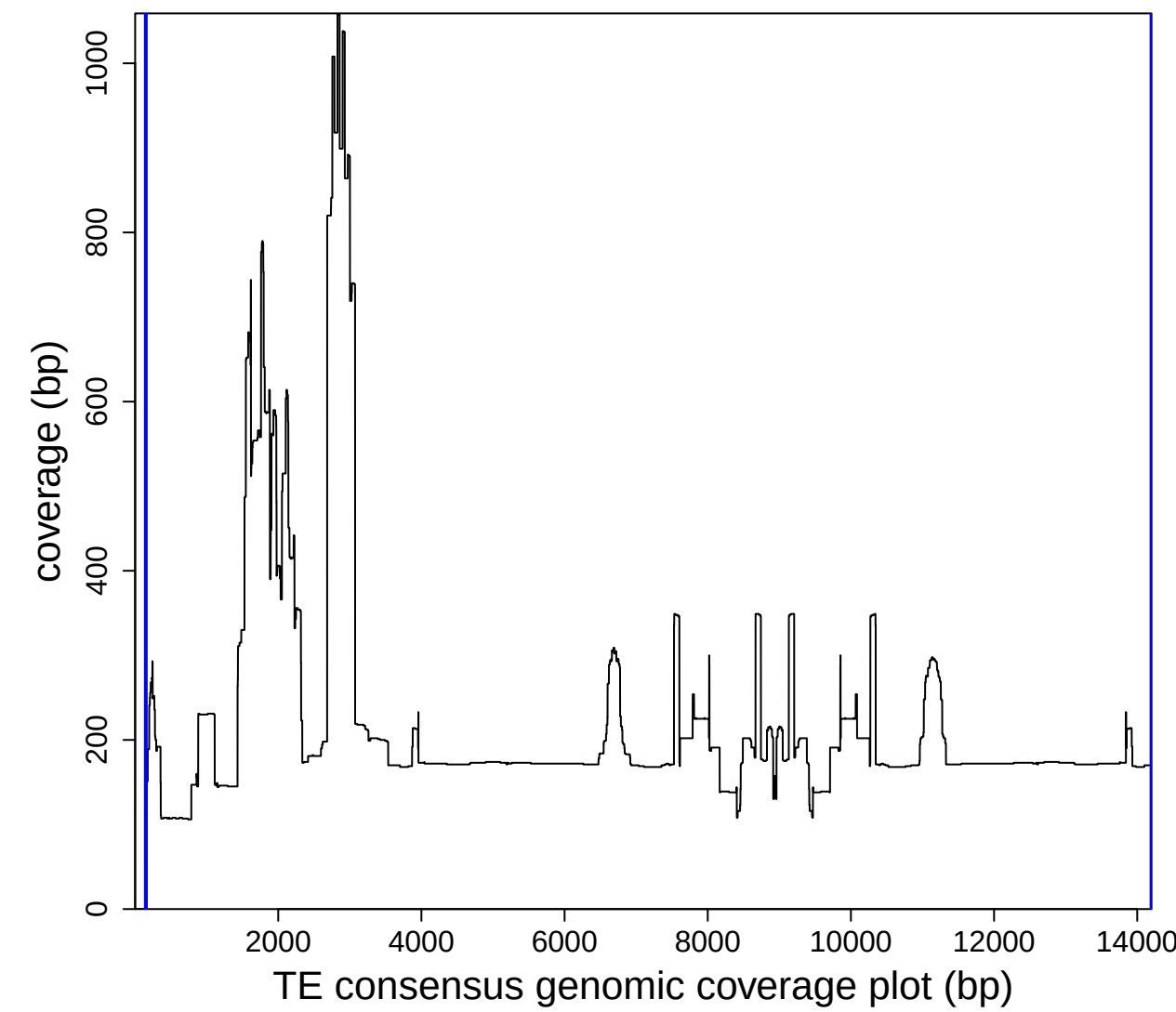
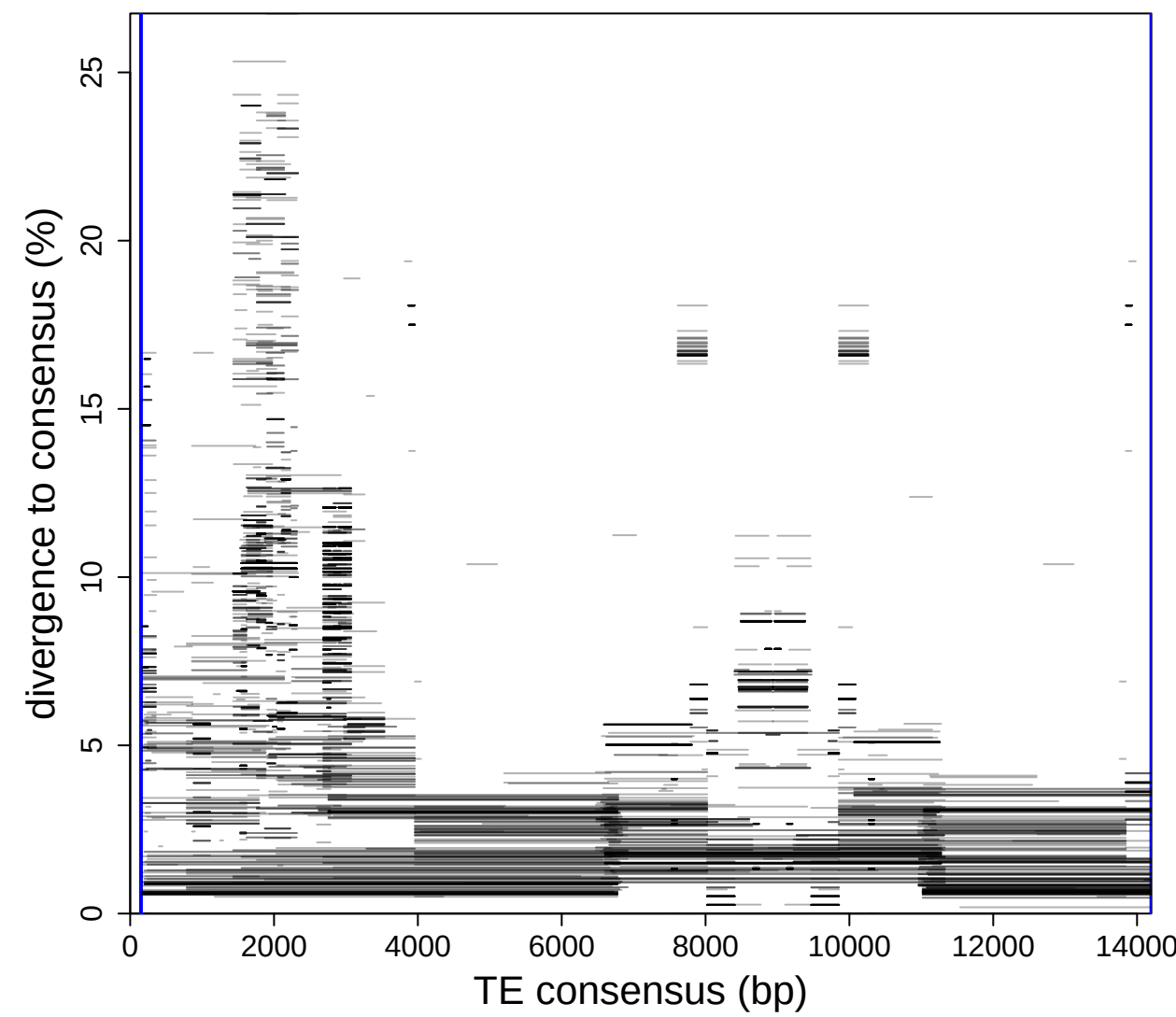
a.b.bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.f
size: 14048bp; fragments: 5100; full length: 0 (>=12643.2bp)

After TEtrimmer 14048 bp



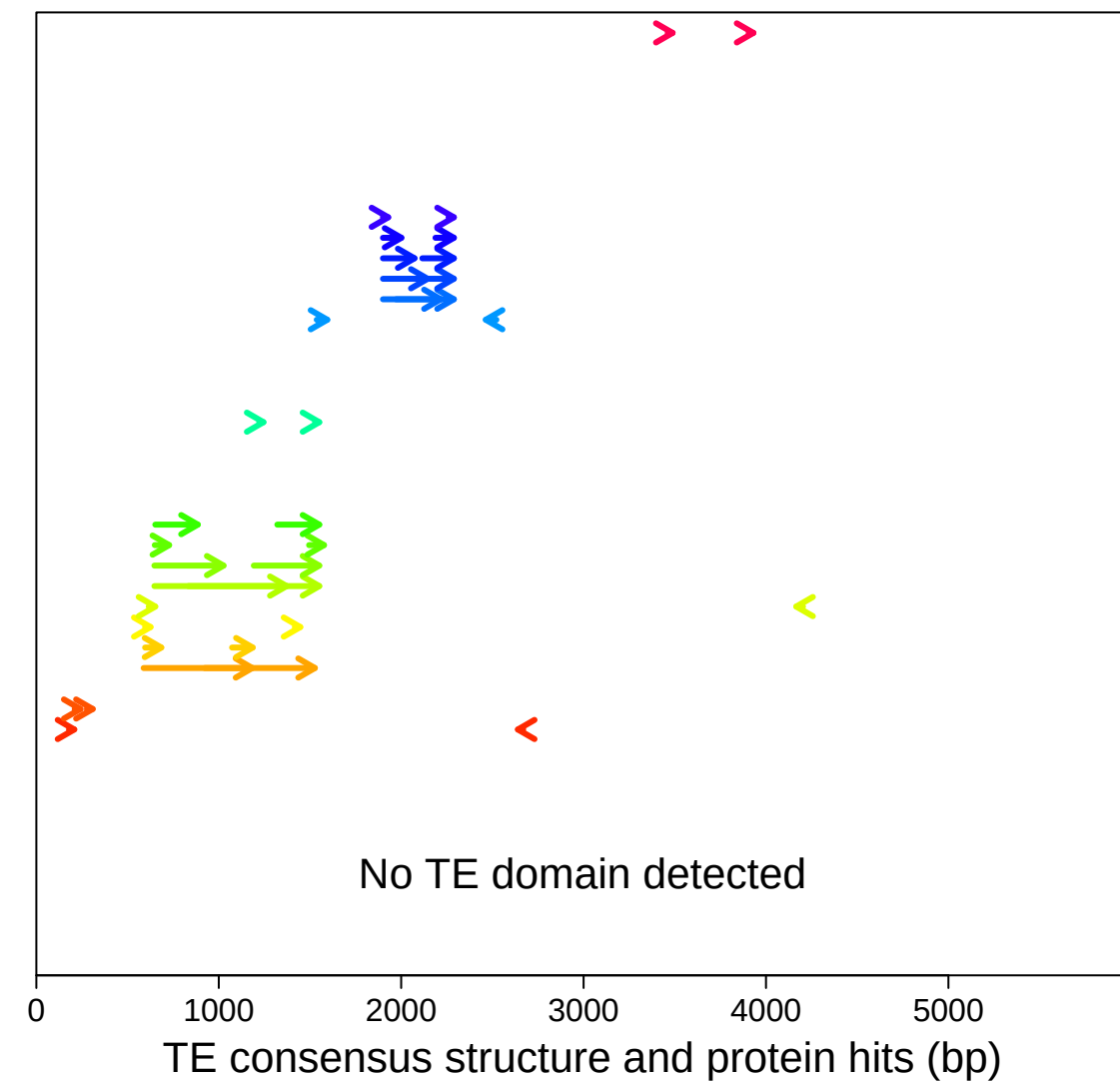
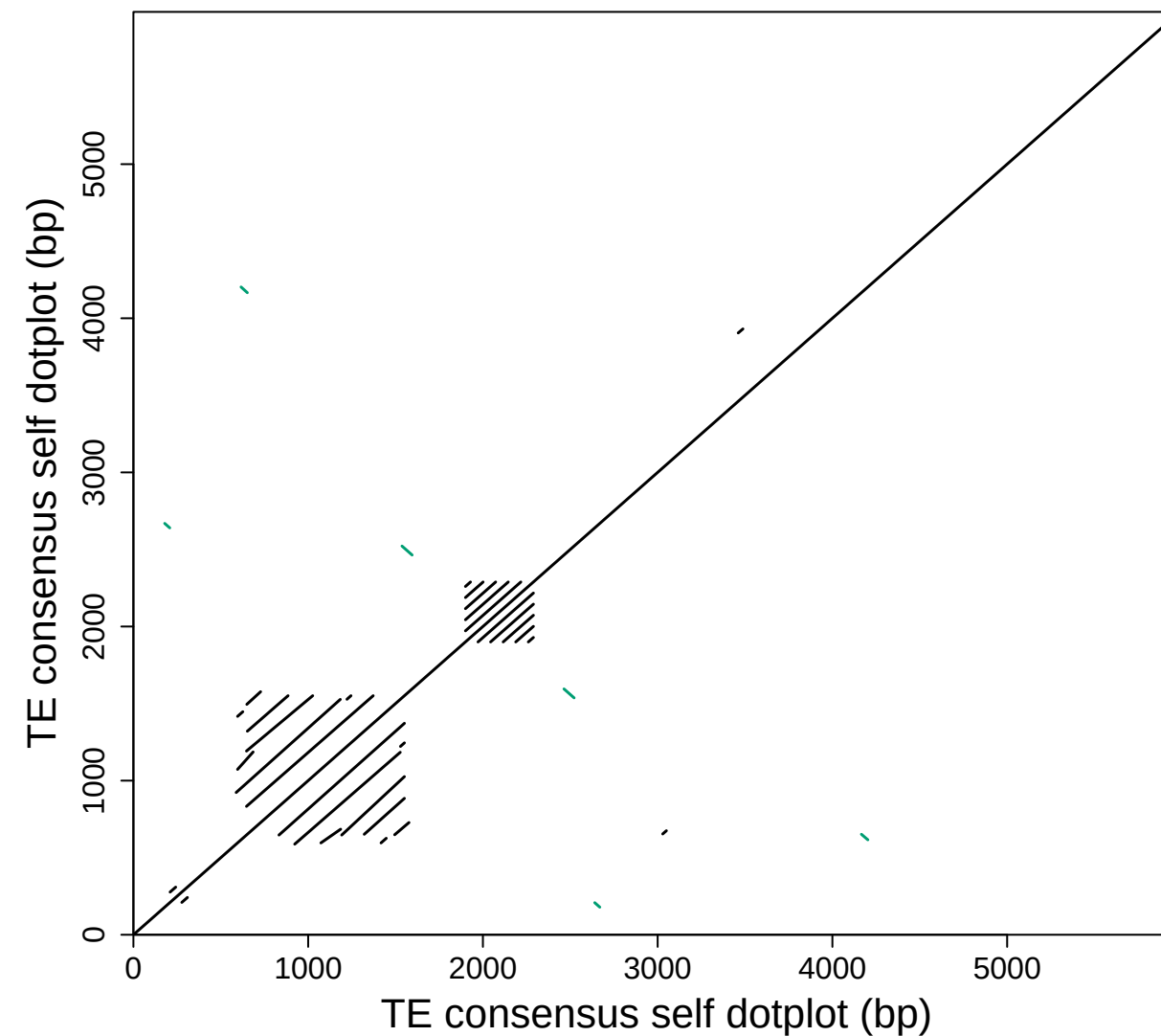
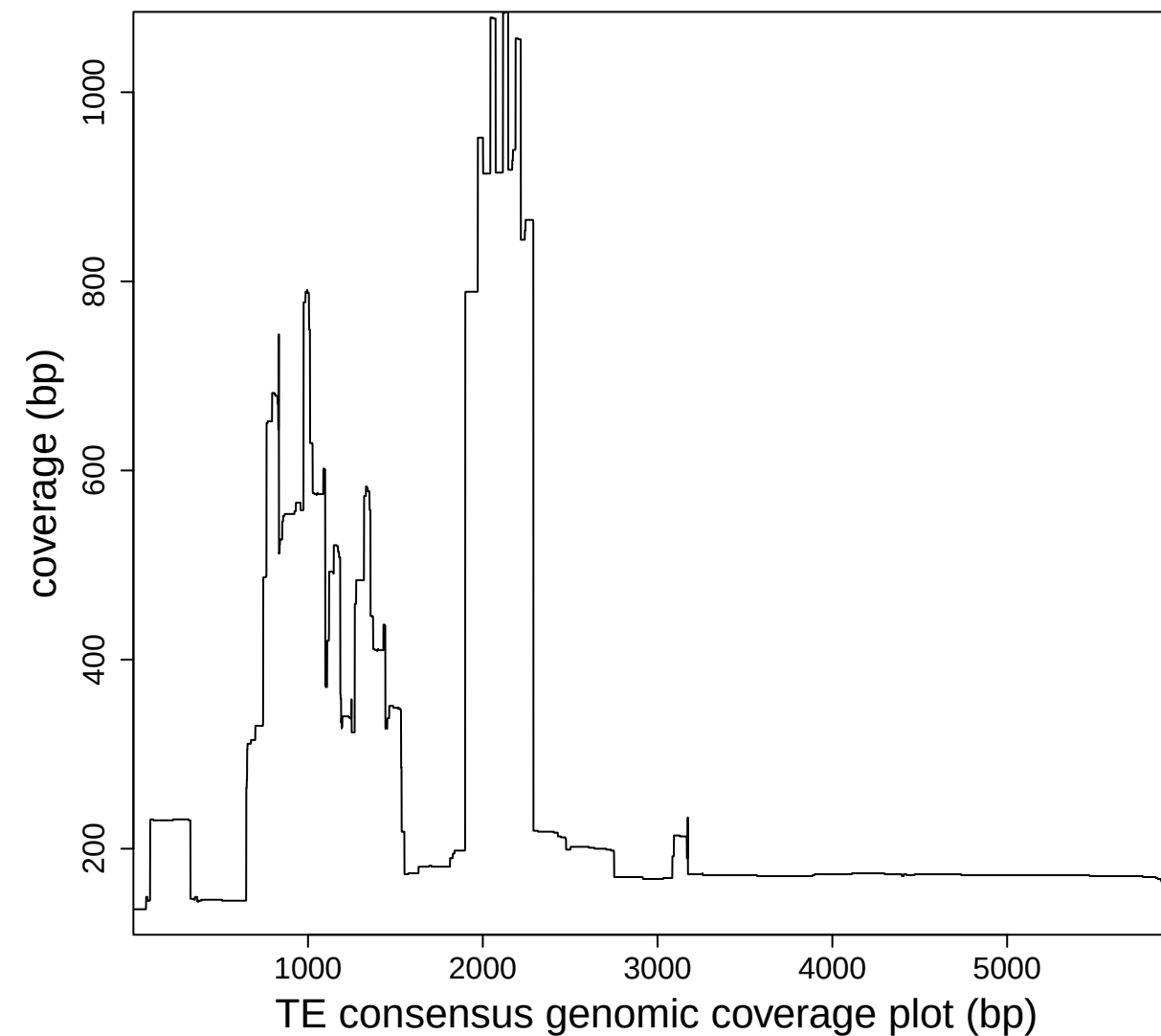
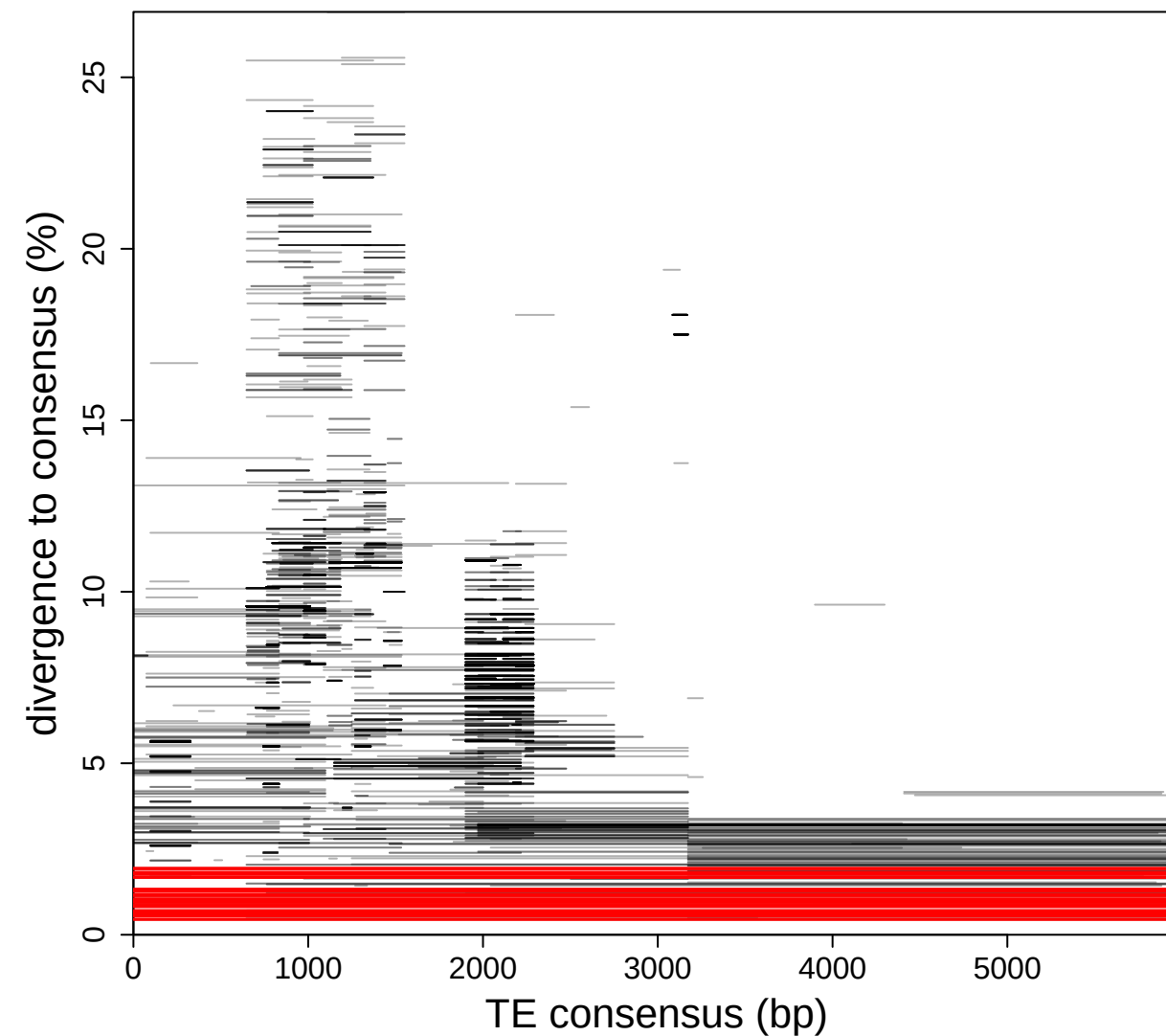
0.fasta.b.bed uf.bed g_1.bed fm_1.bed 0_0_bcln.fa aln.fa cl.f
size: 14199bp; fragments: 5058; full length: 0 (>=12779.1bp)

After TEtrimmer Extended plot Blue lines are boundaries

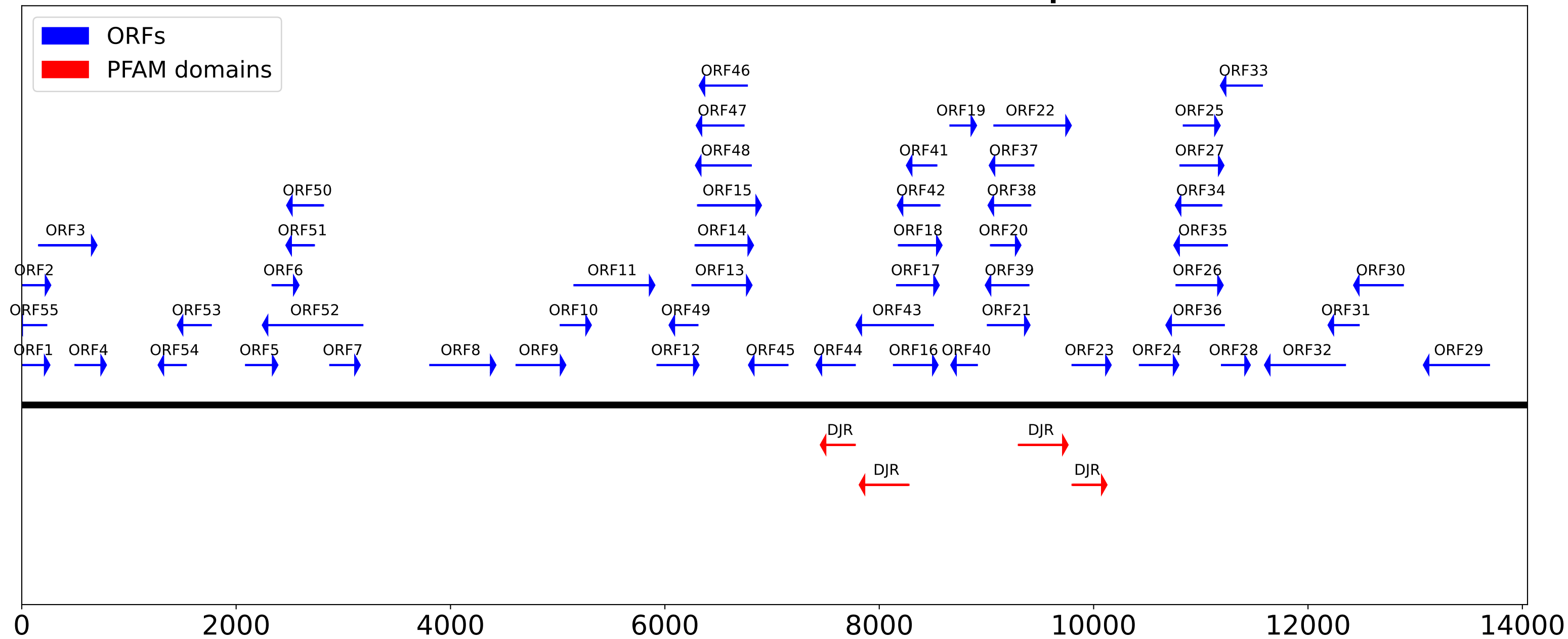


TE: rnd_5_family_1590
size: 5988bp; fragments: 3104; full length: 70 (≥ 5389.2 bp)

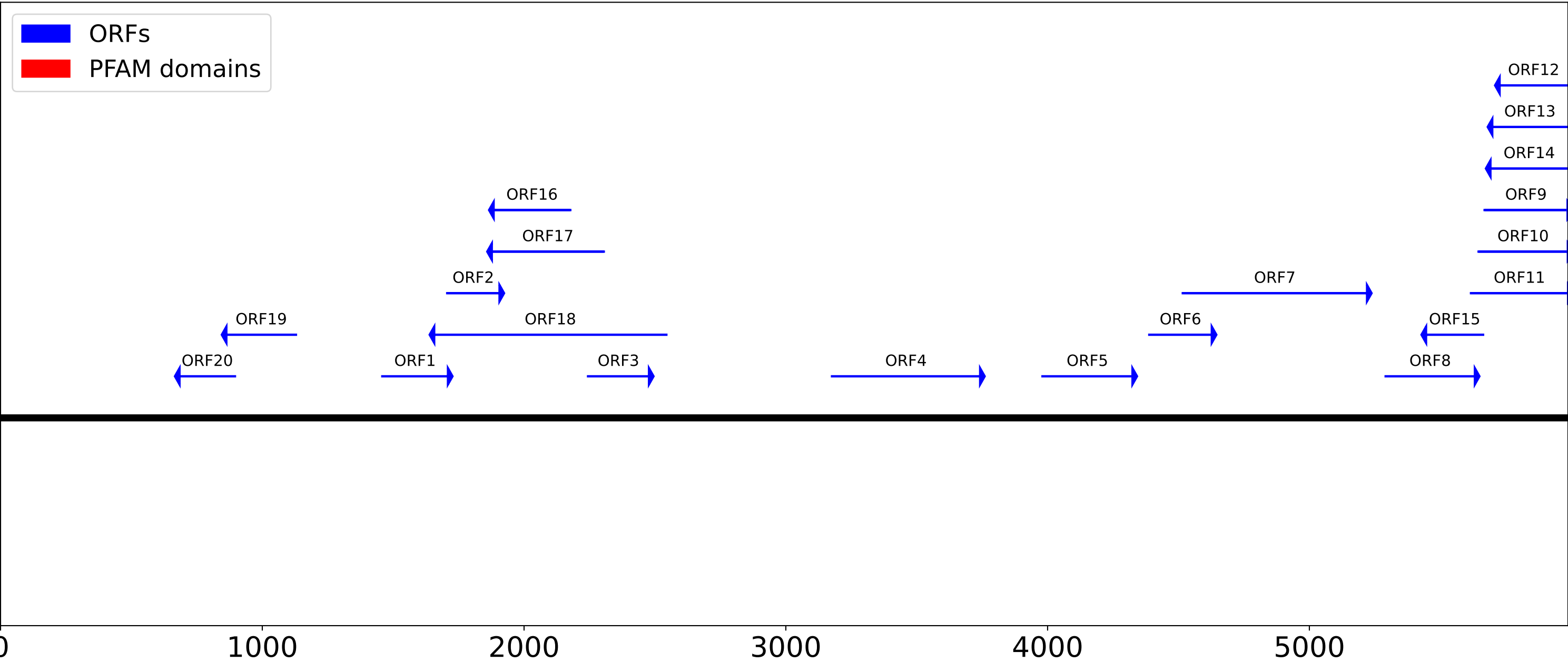
Before TEtrimmer 5988 bp



After TEtrimmer ORF and PFAM domain plot



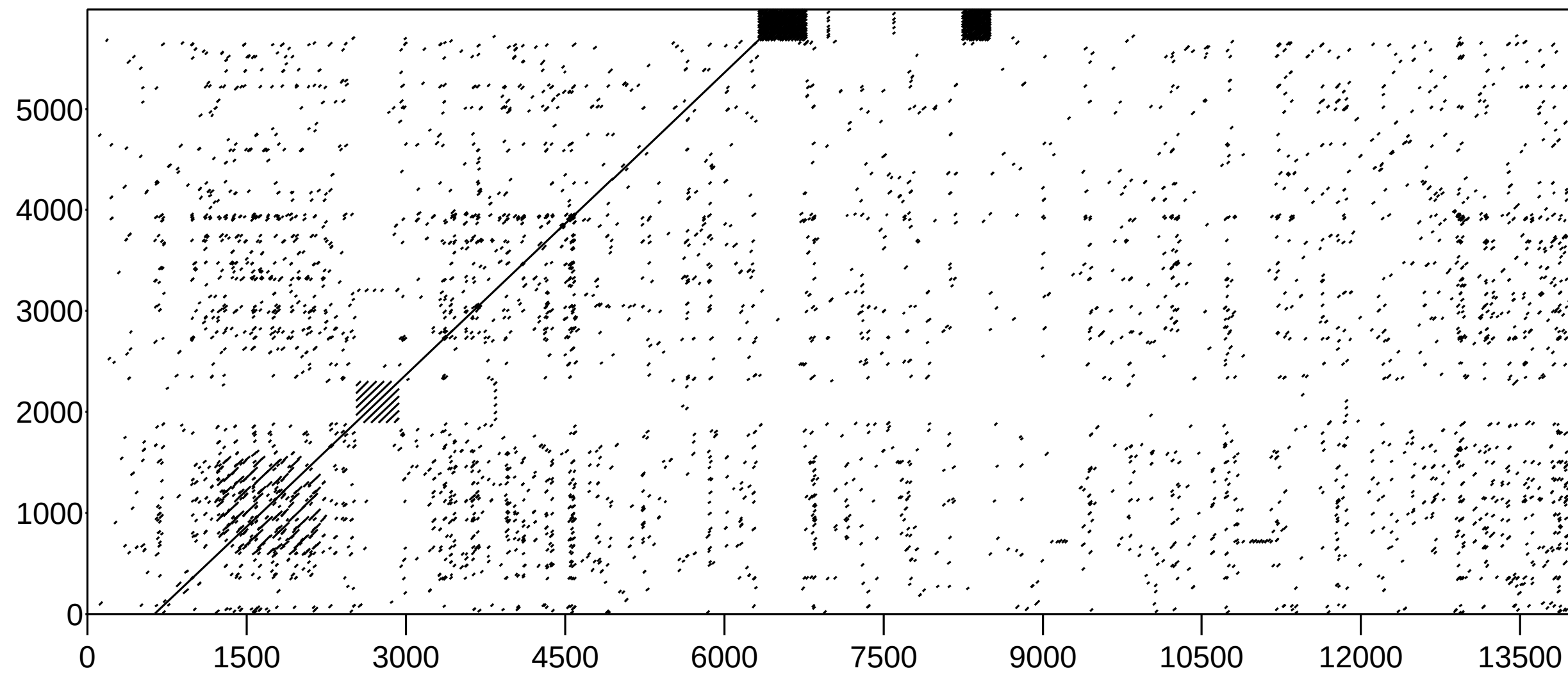
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)