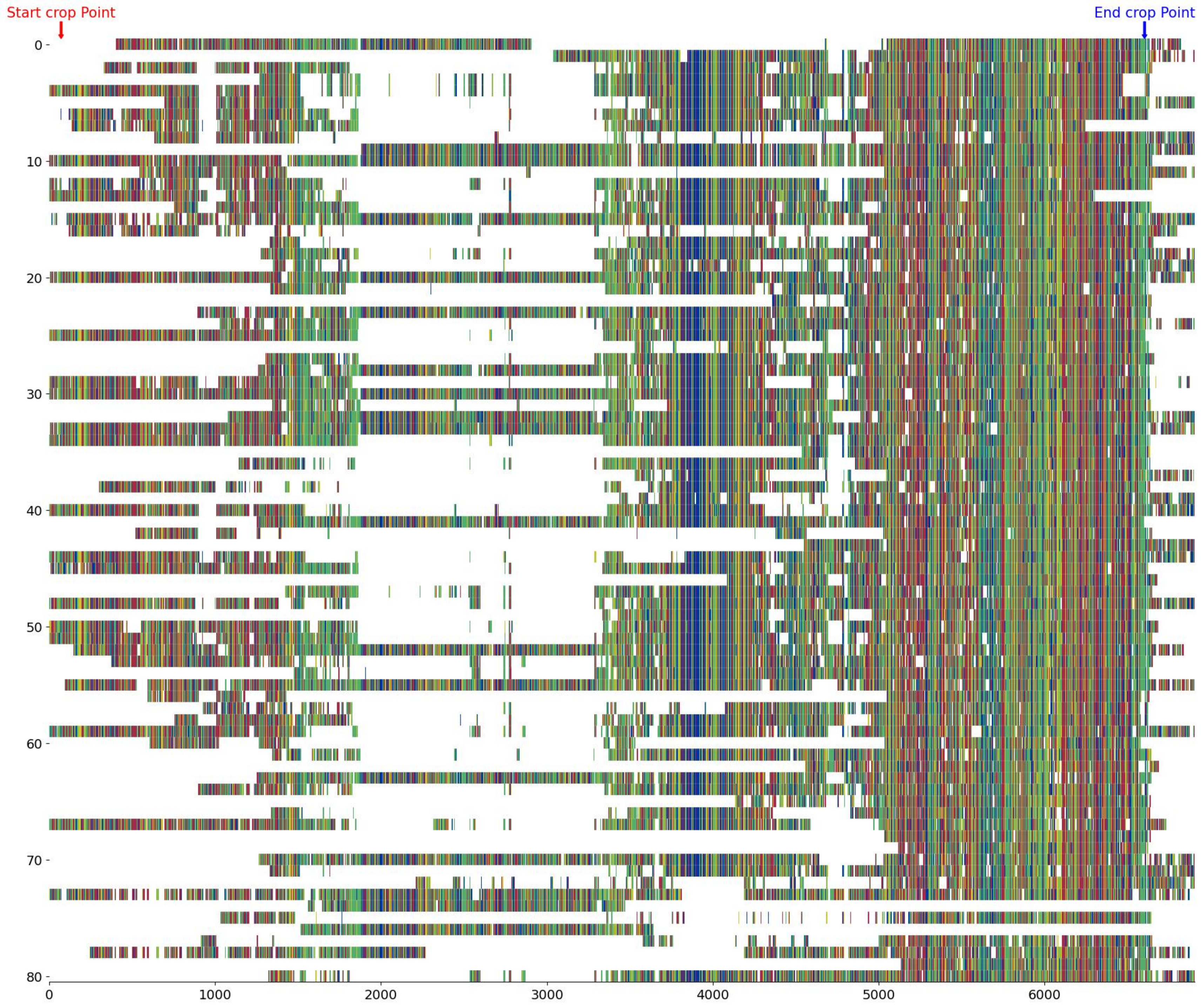
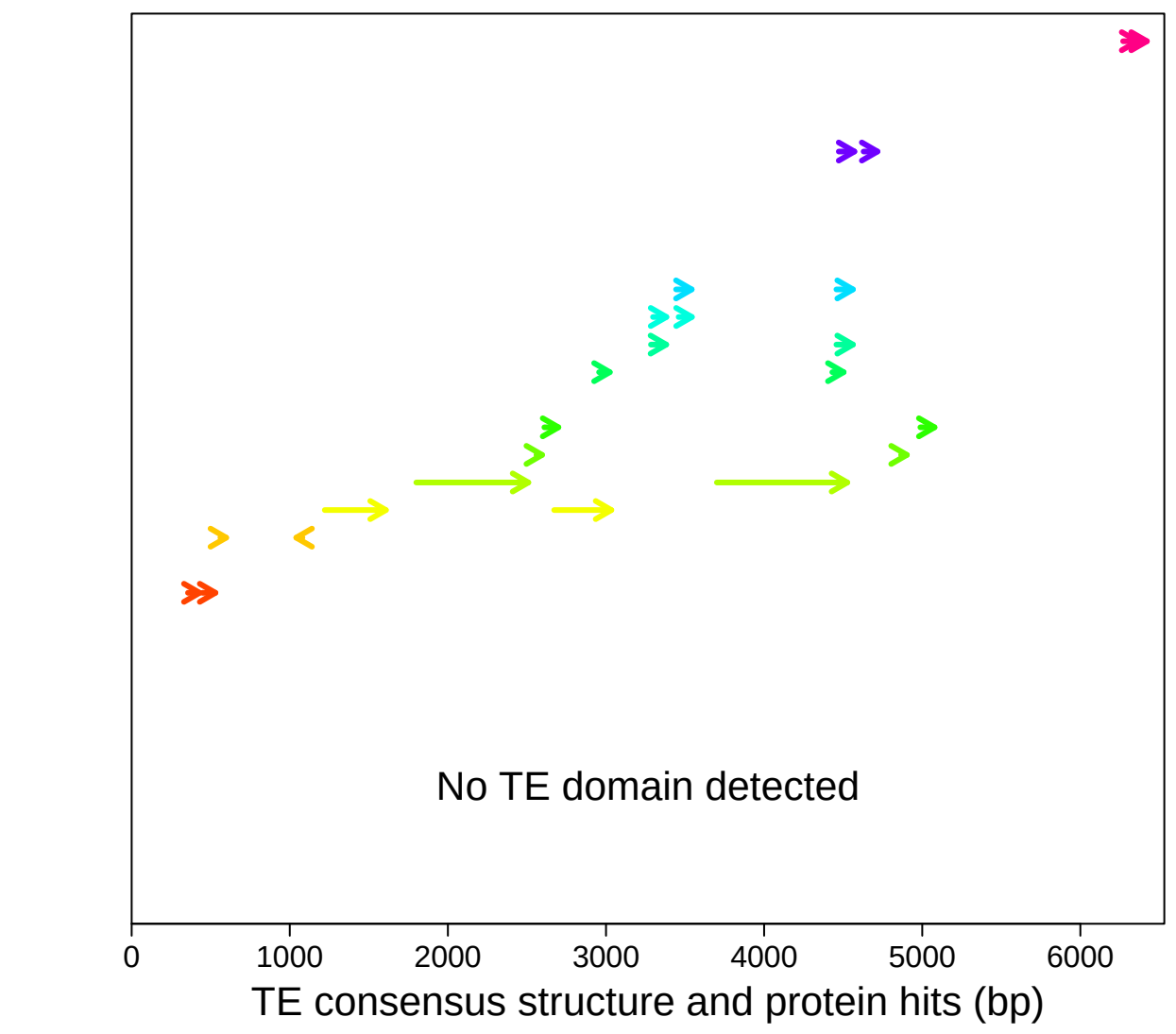
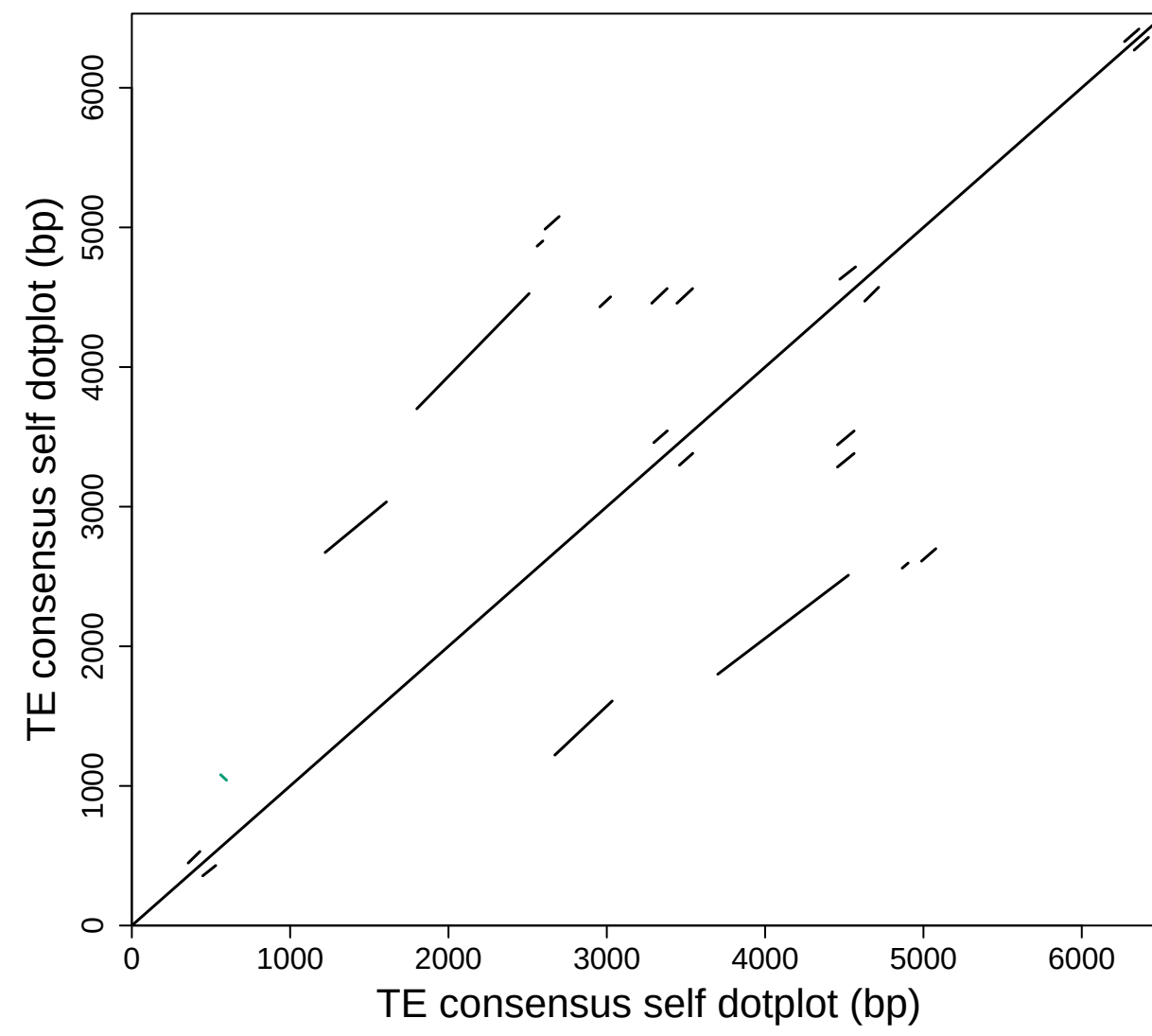
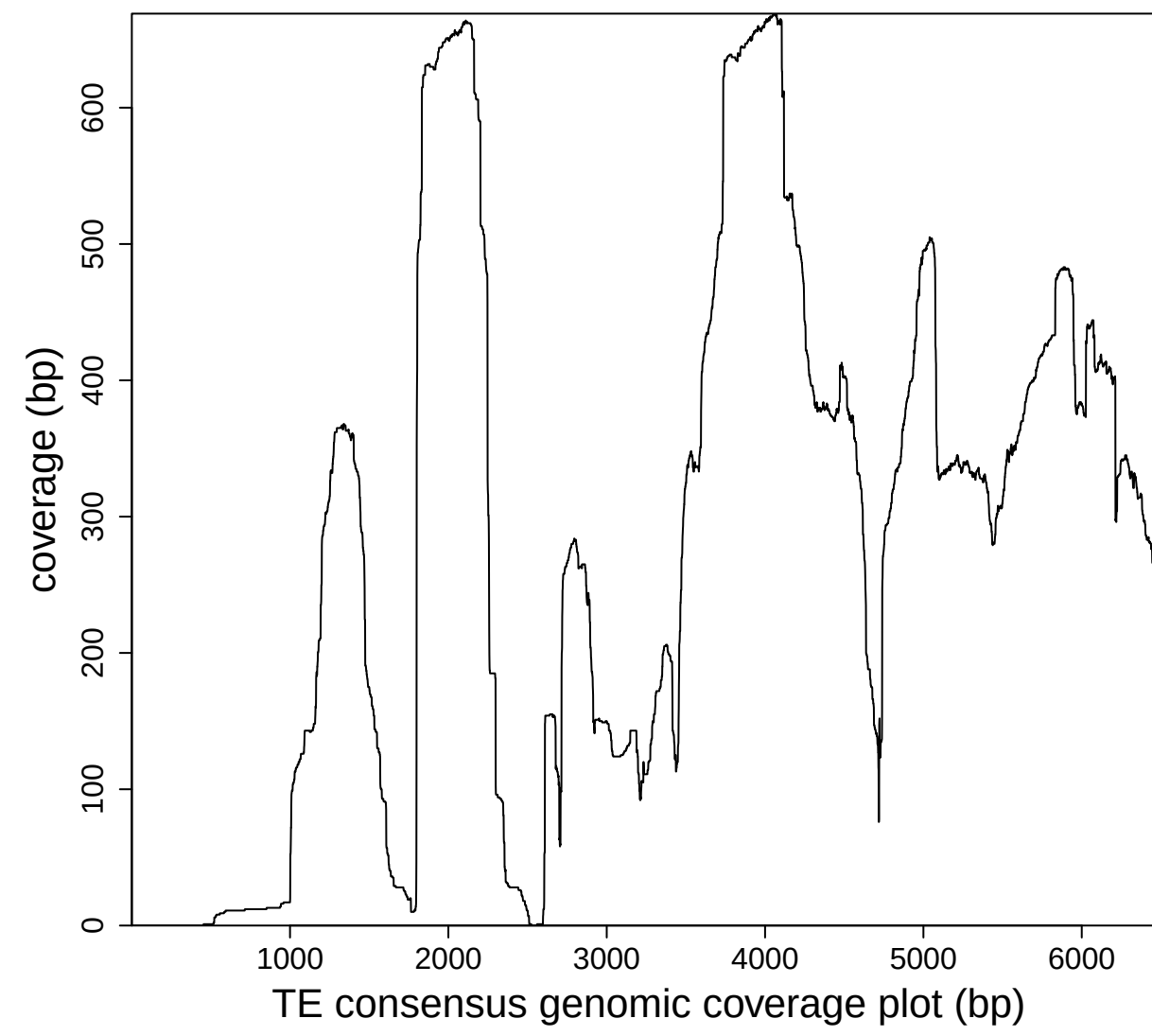
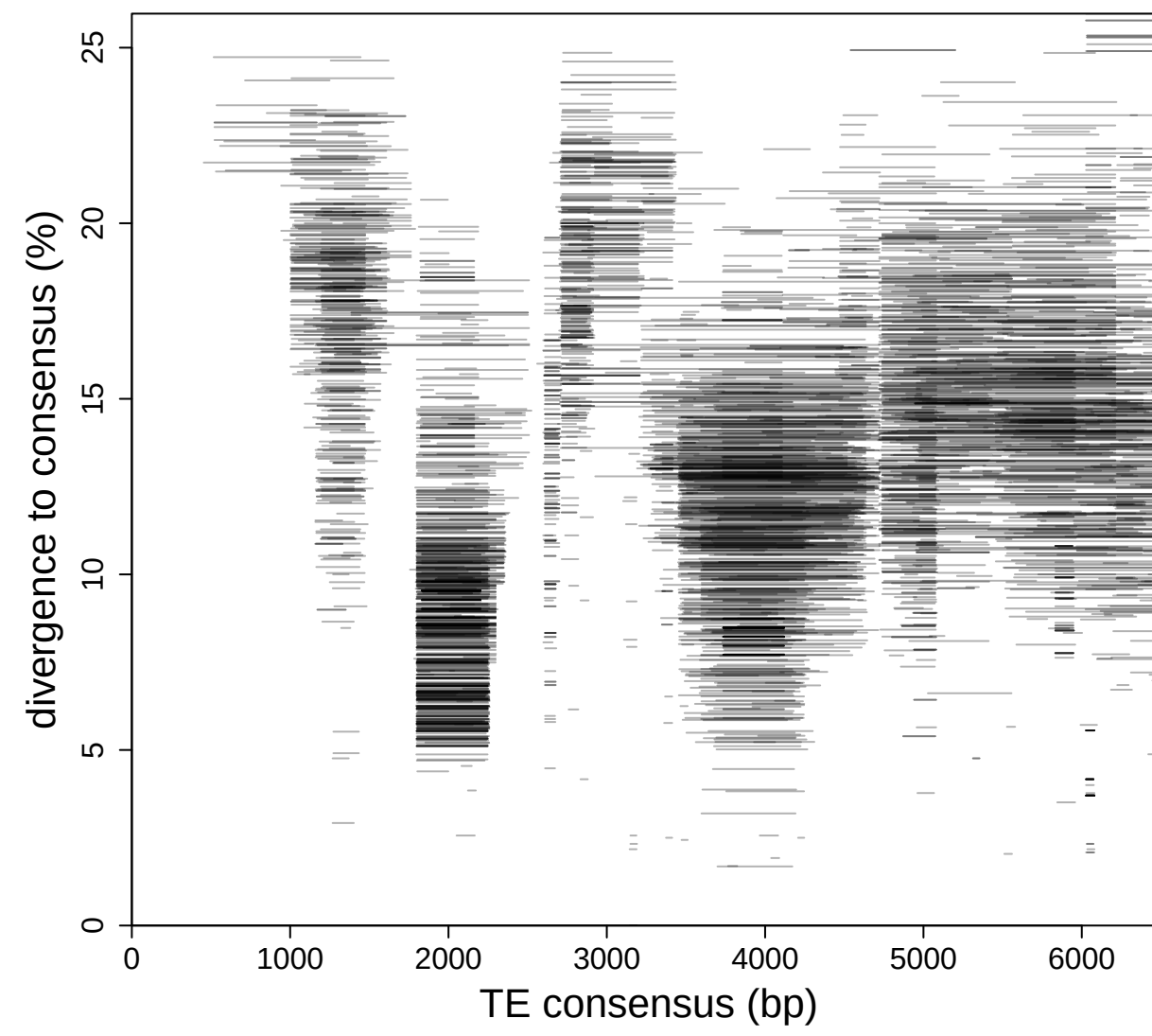






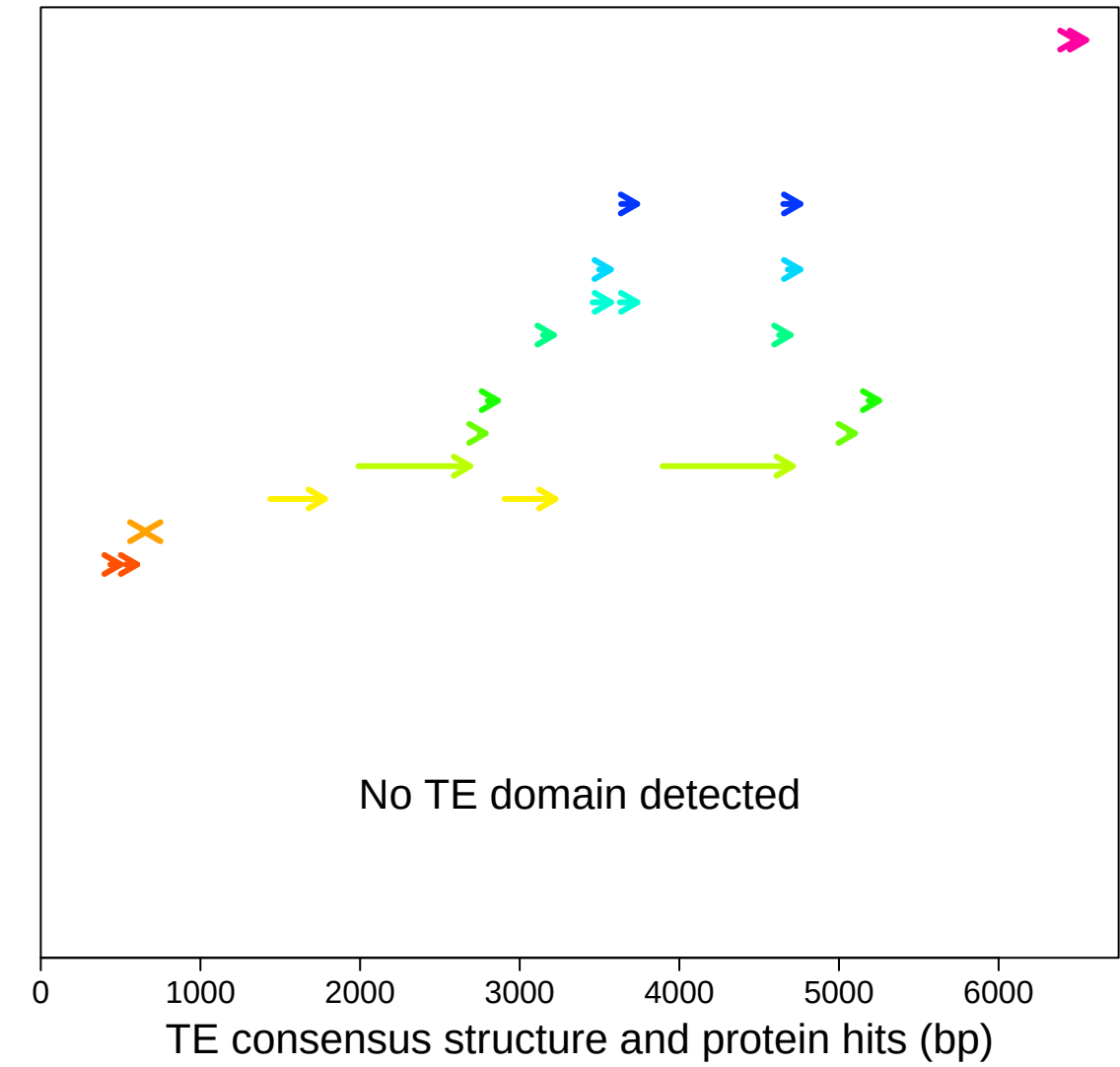
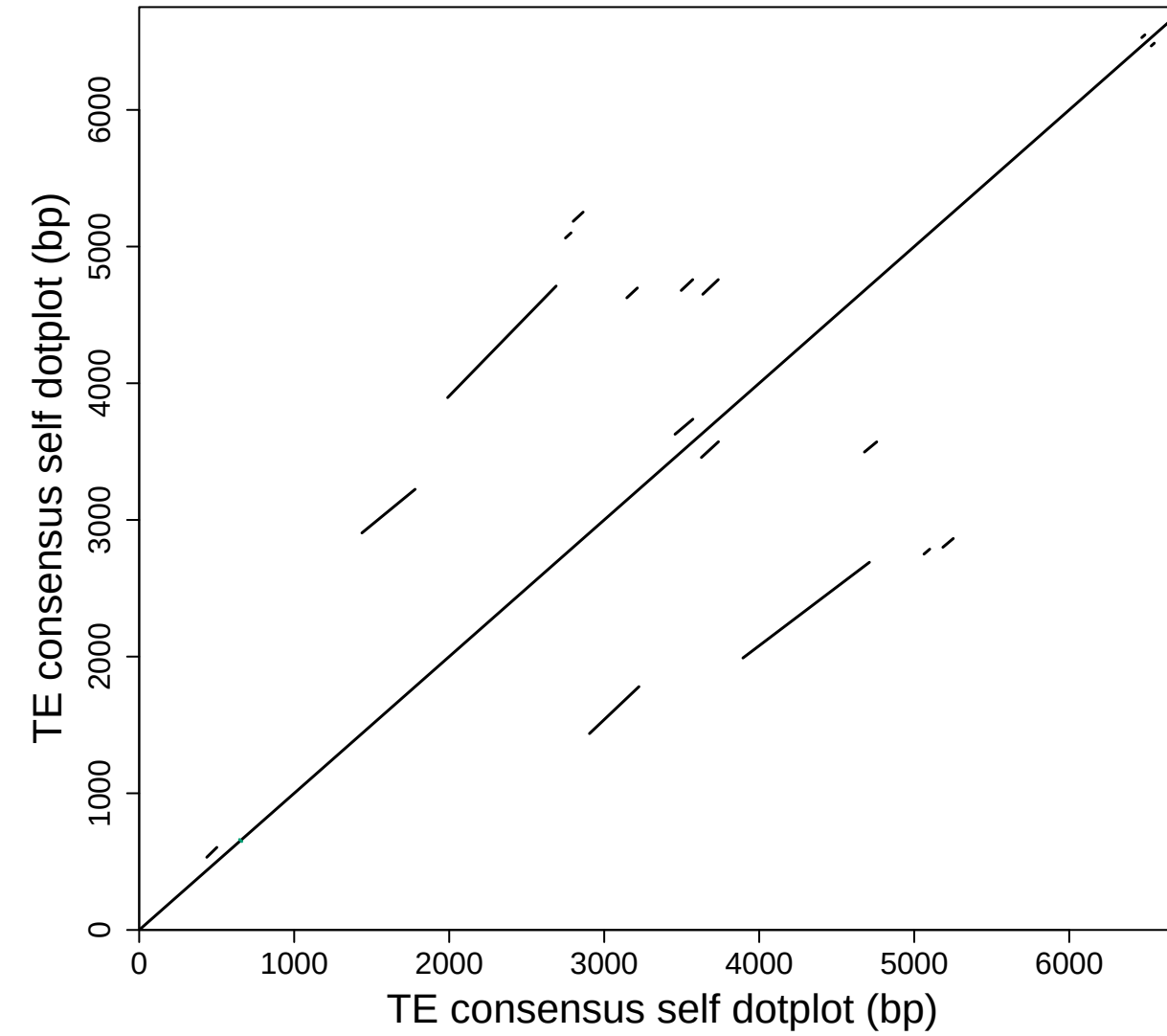
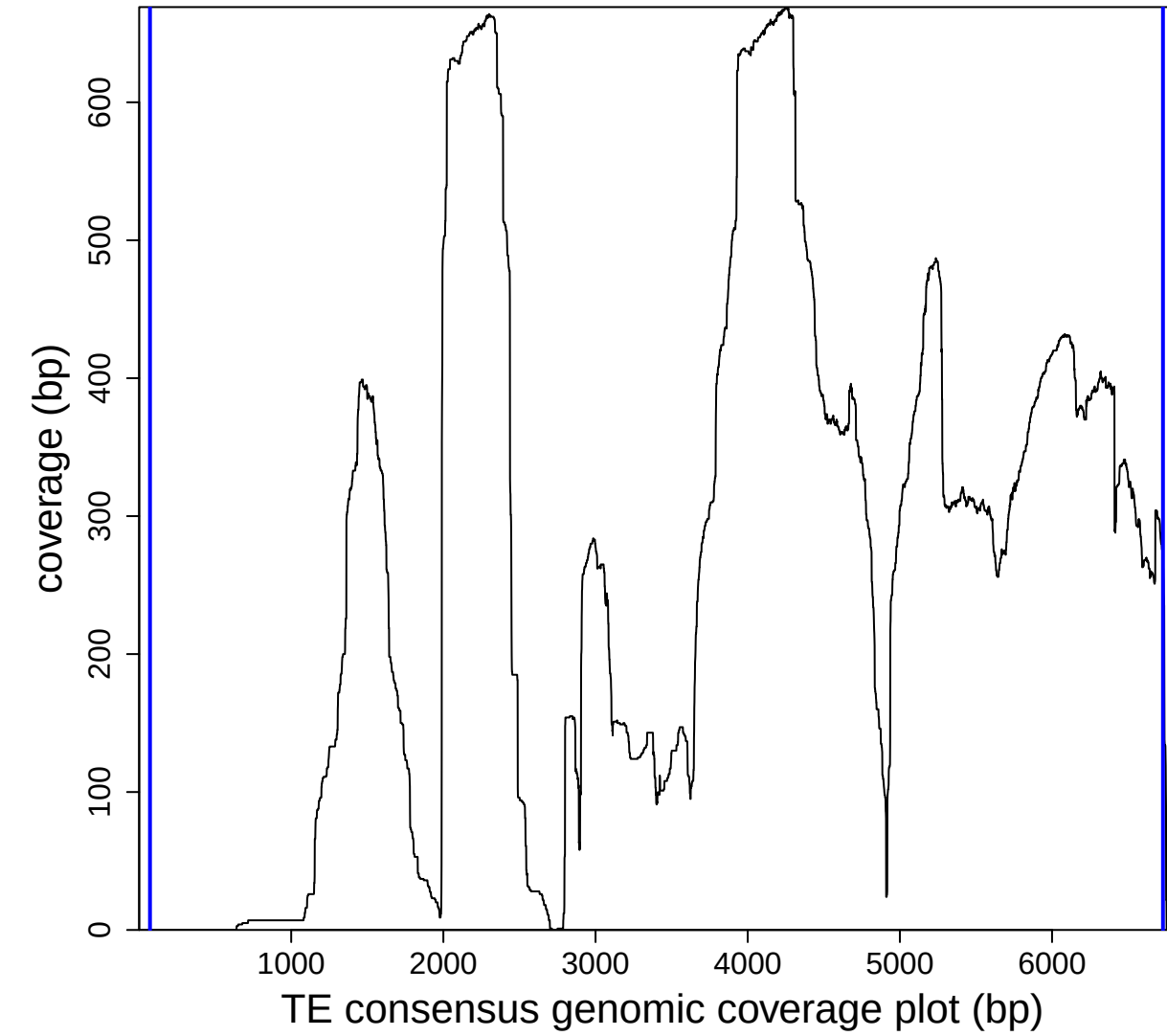
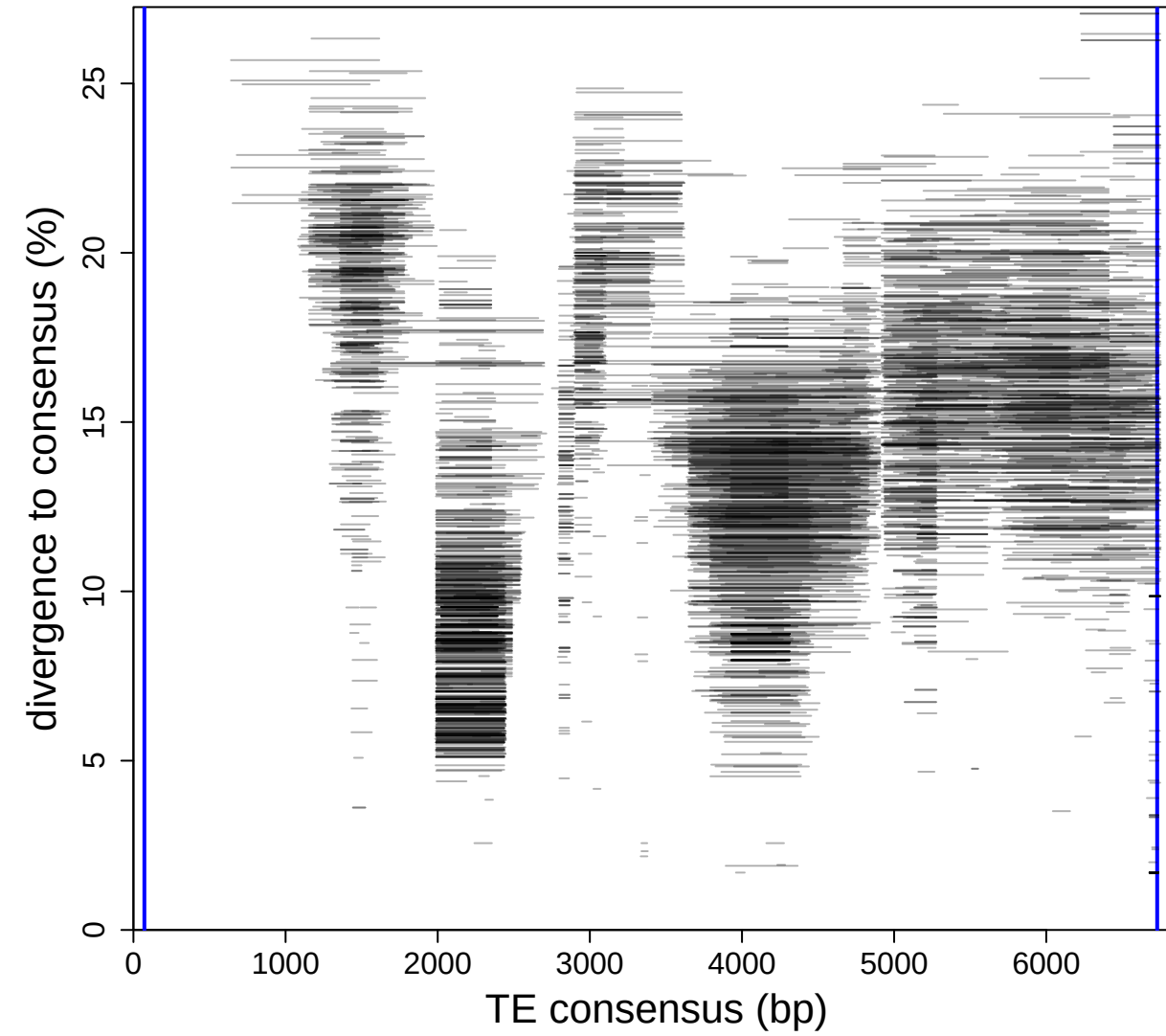
a.b.bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.f
size: 6531bp; fragments: 3635; full length: 0 (>=5877.9bp)

After TEtrimmer 6531 bp



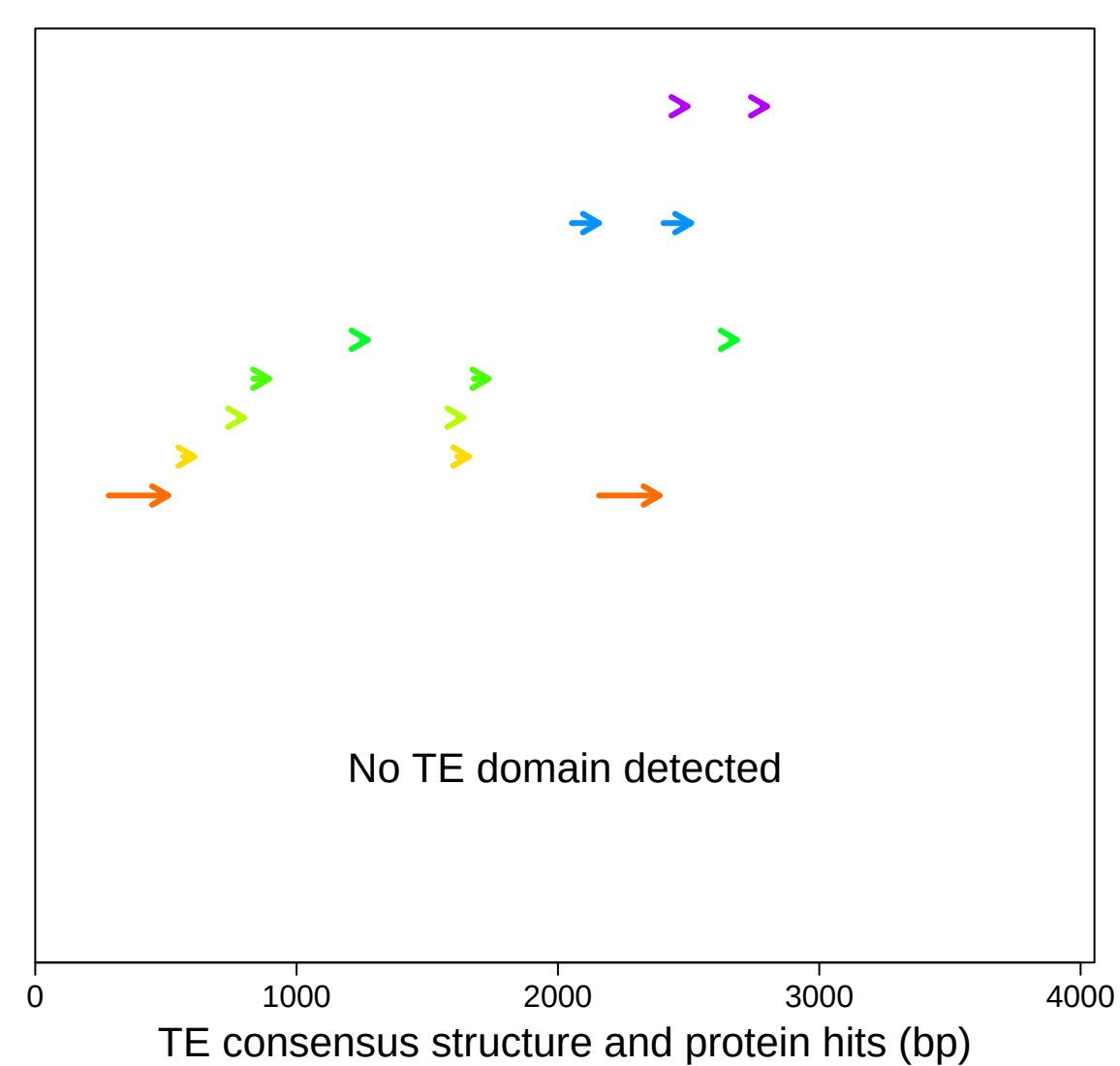
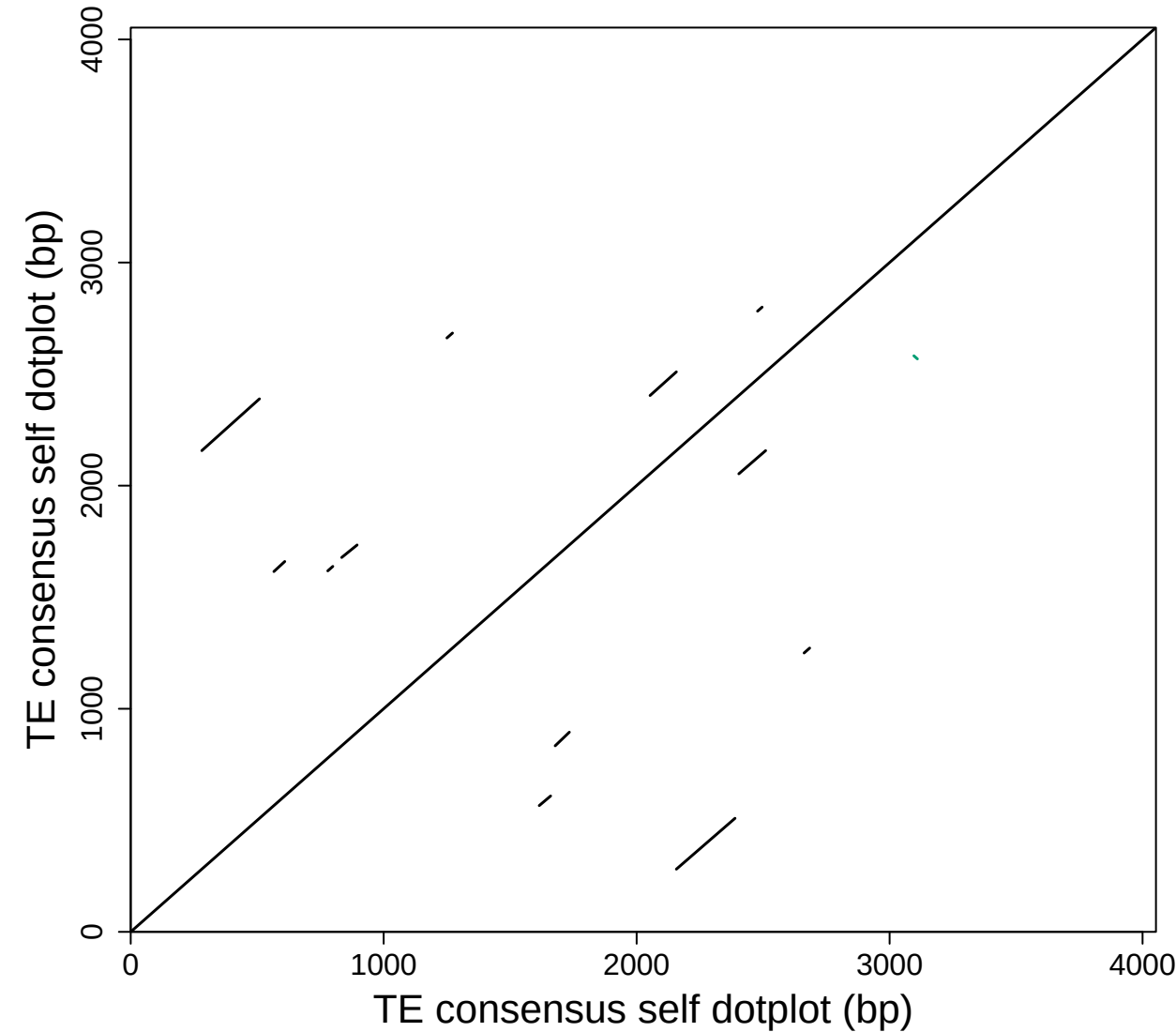
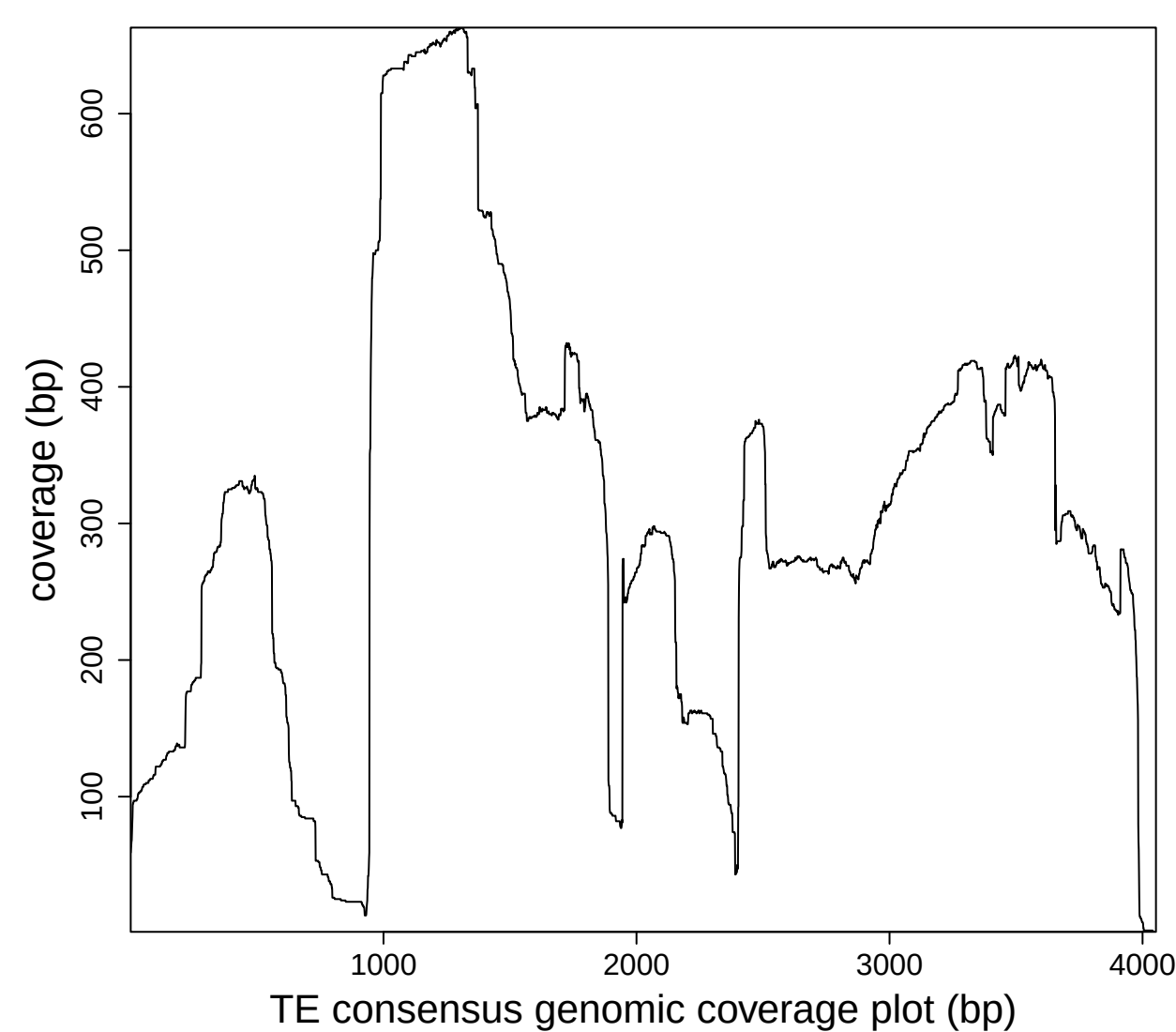
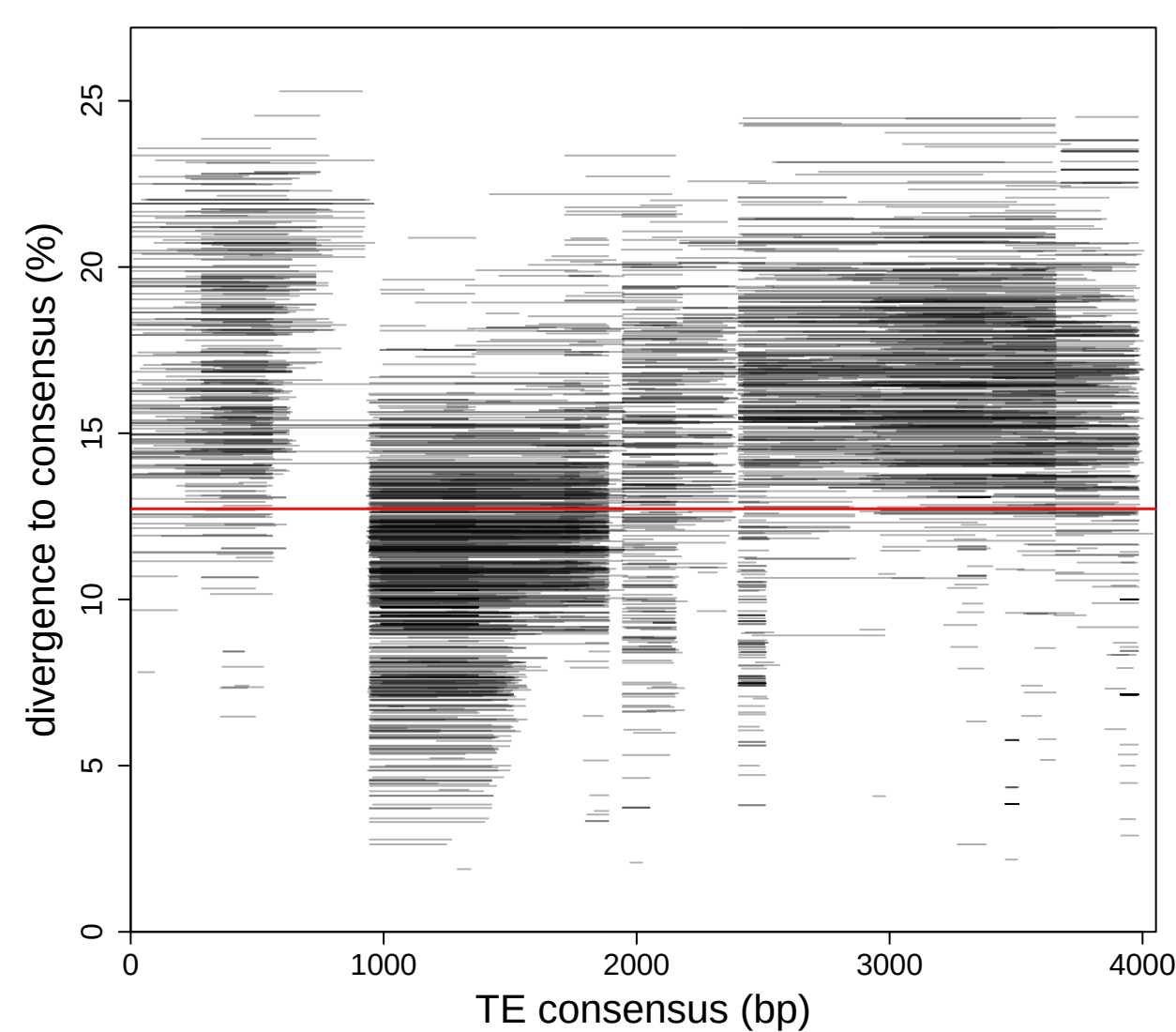
1.fasta.b.bed uf.bed g_1.bed fm_1.bed 0 0 bcln.fa aln.fa cl.f
size: 6879bp; fragments: 3444; full length: 0 (>=6191.1bp)

After TEtrimmer Extended plot Blue lines are boundaries

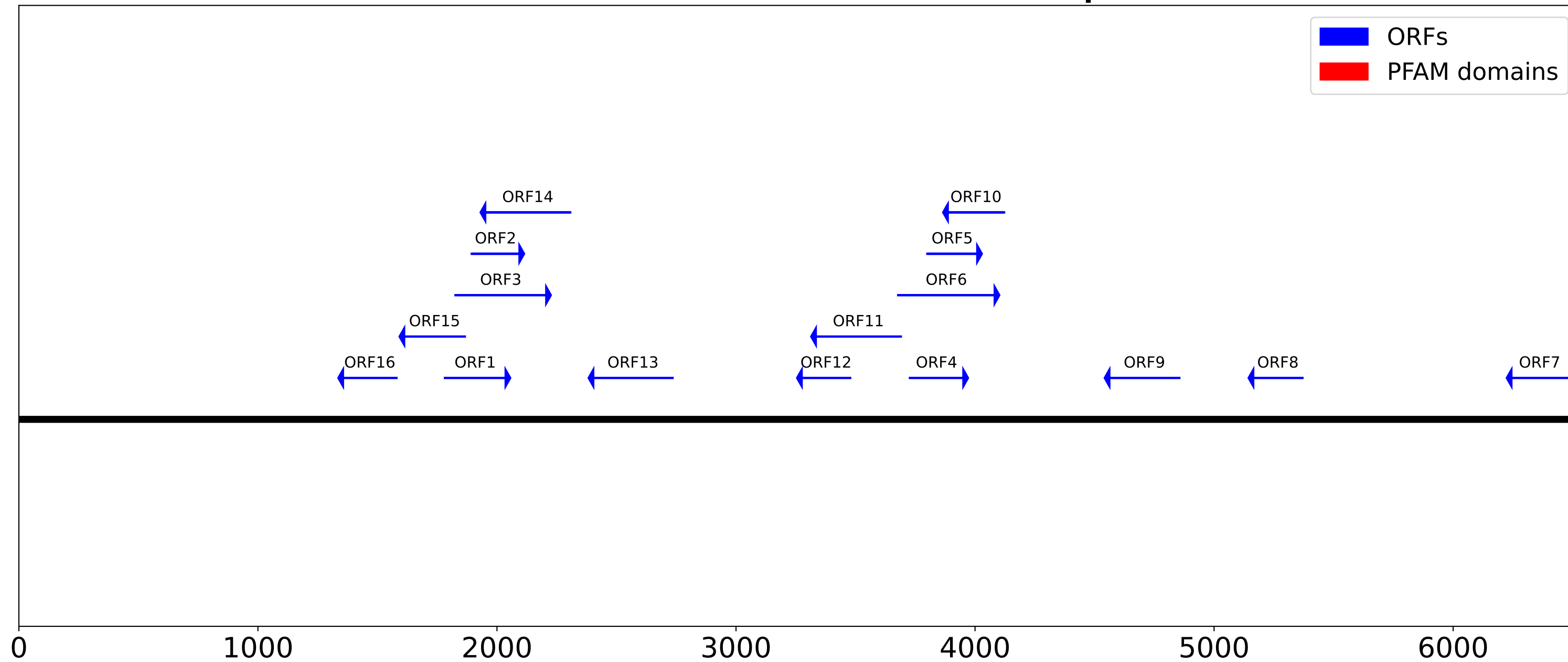


TE: rnd_5_family_8361
size: 4053bp; fragments: 2440; full length: 1 (≥ 3647.7 bp)

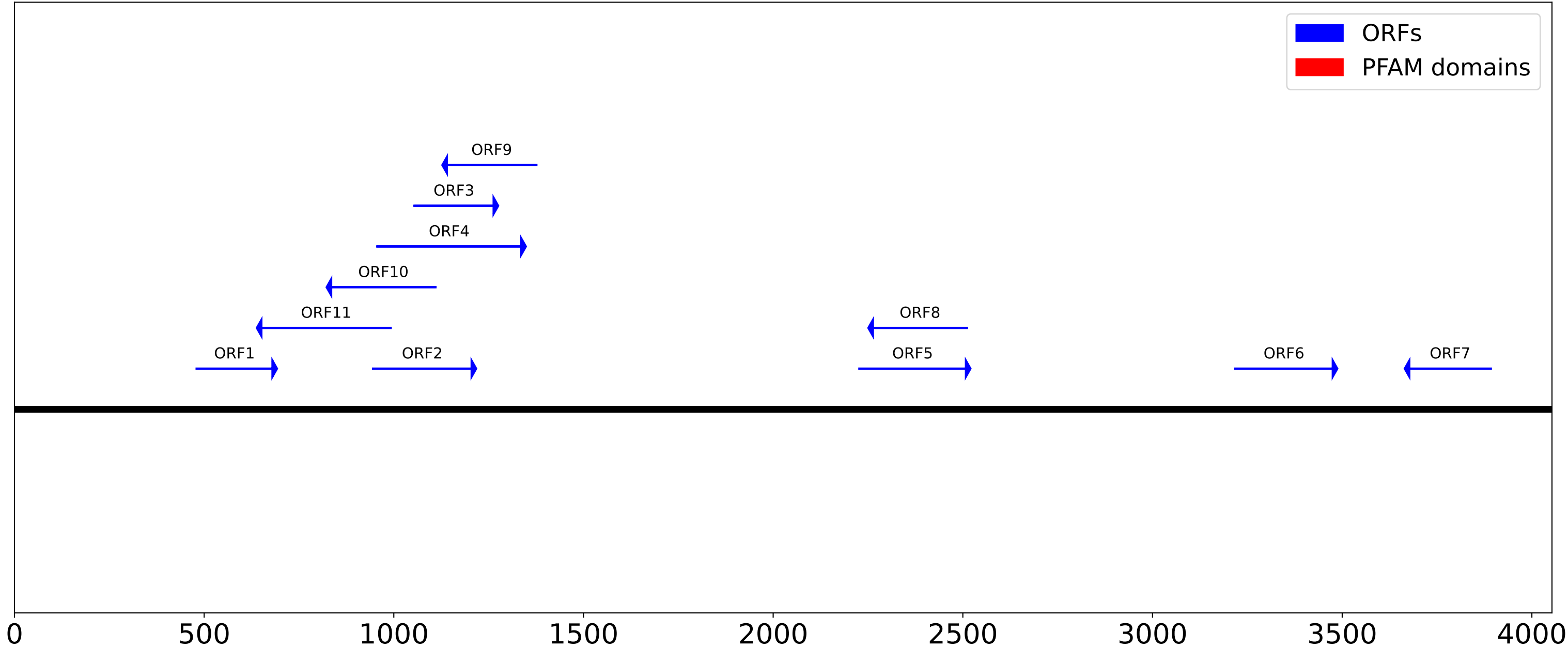
Before TEtrimmer 4053 bp



After TEtrimmer ORF and PFAM domain plot



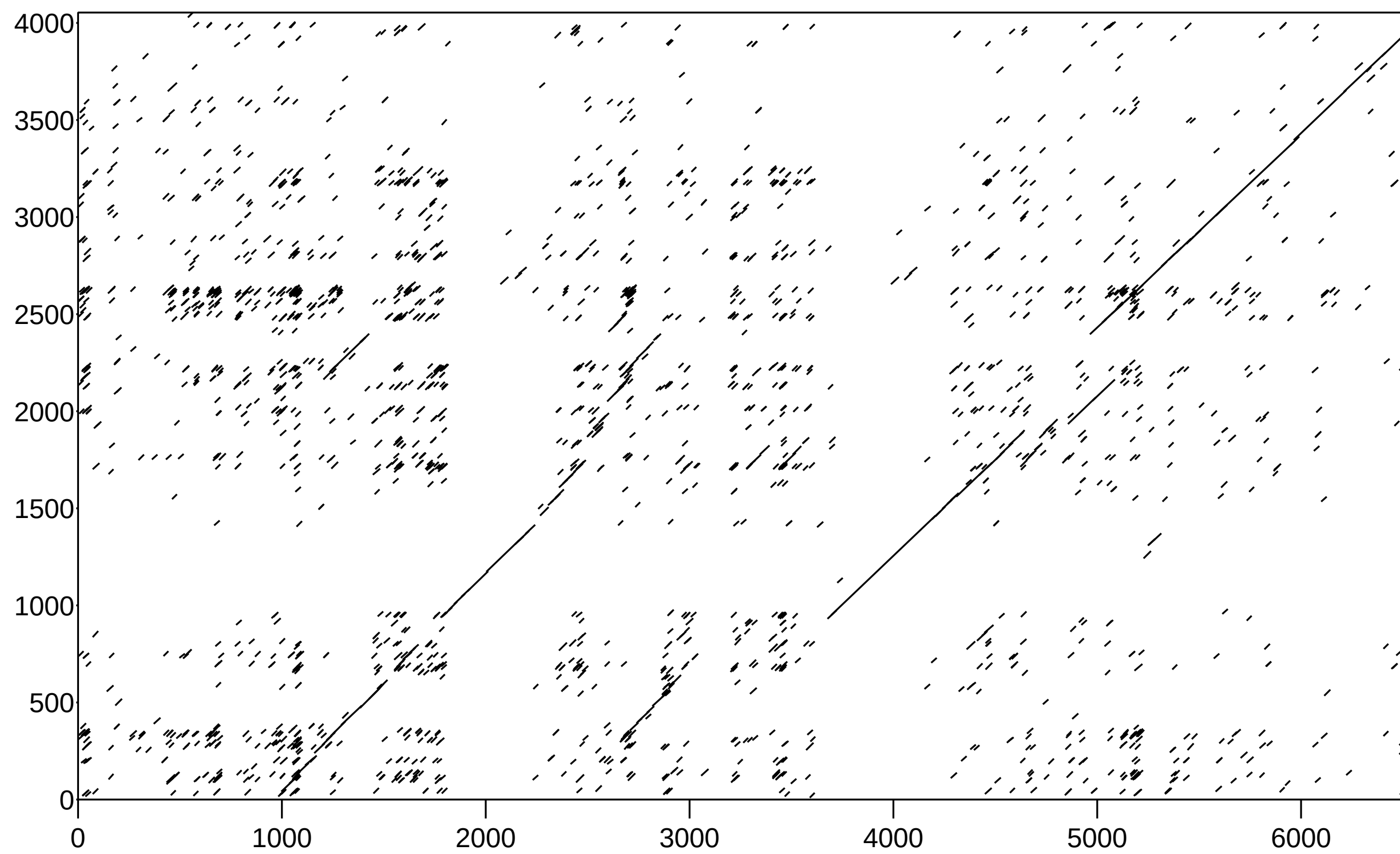
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)