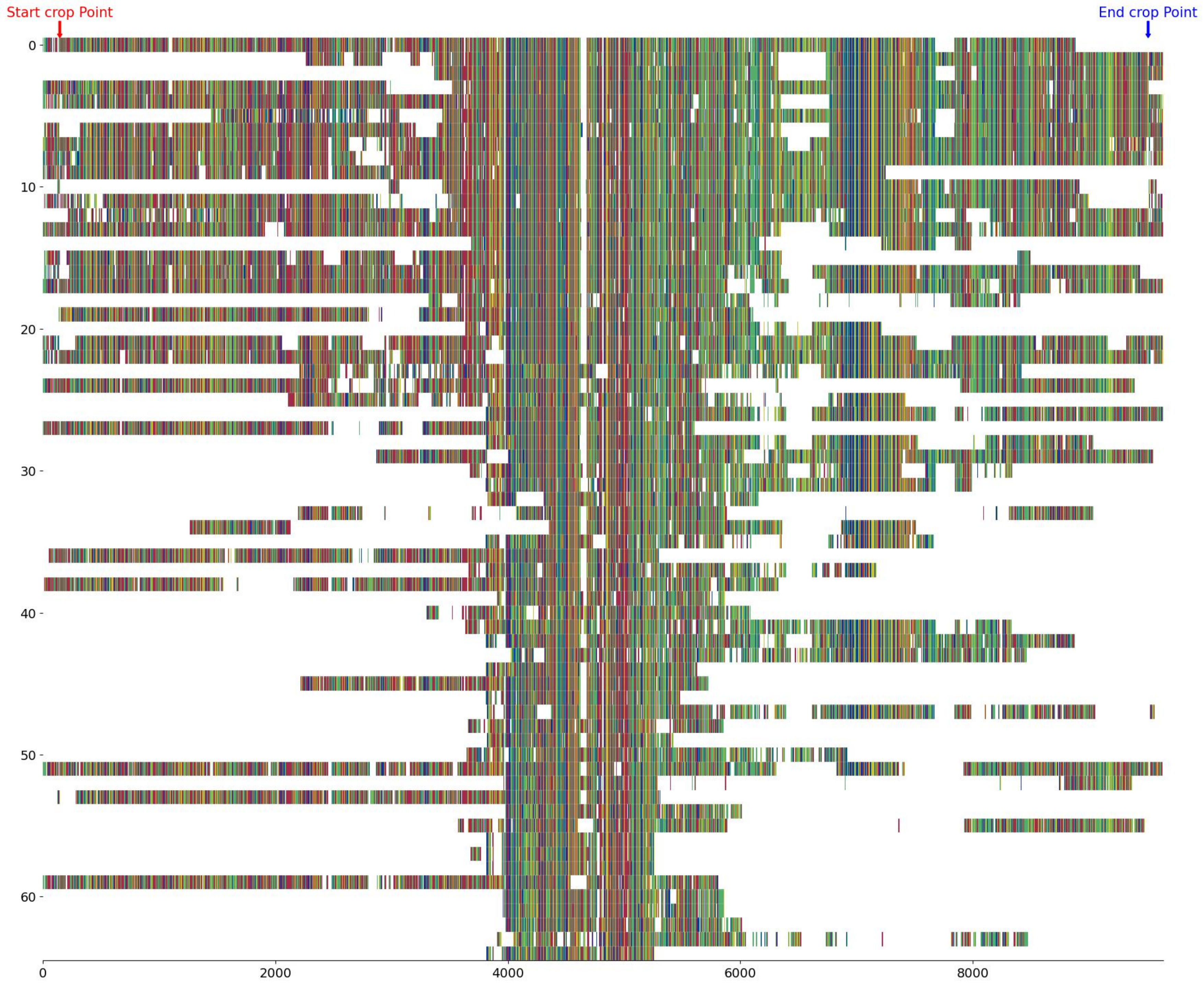
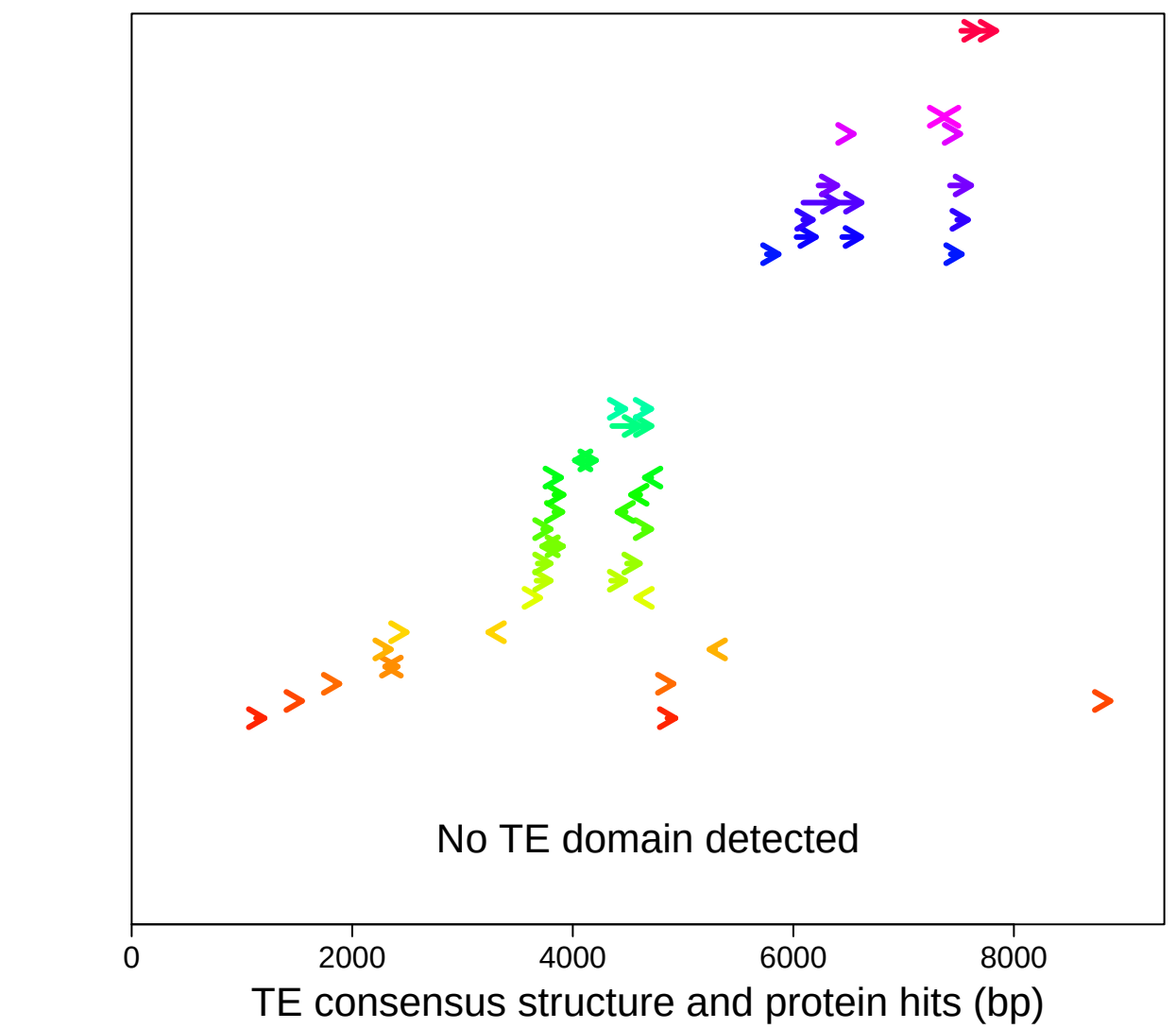
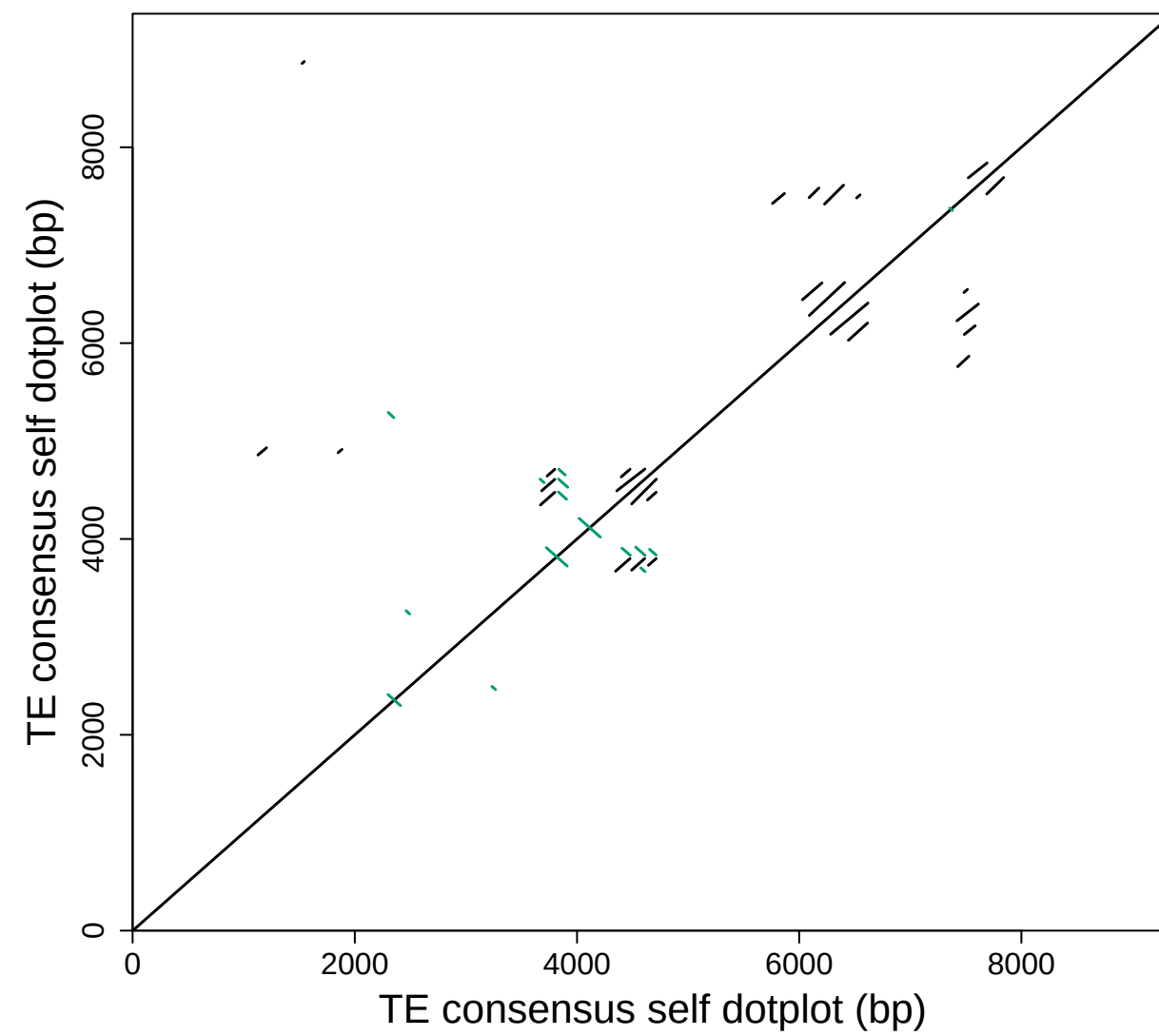
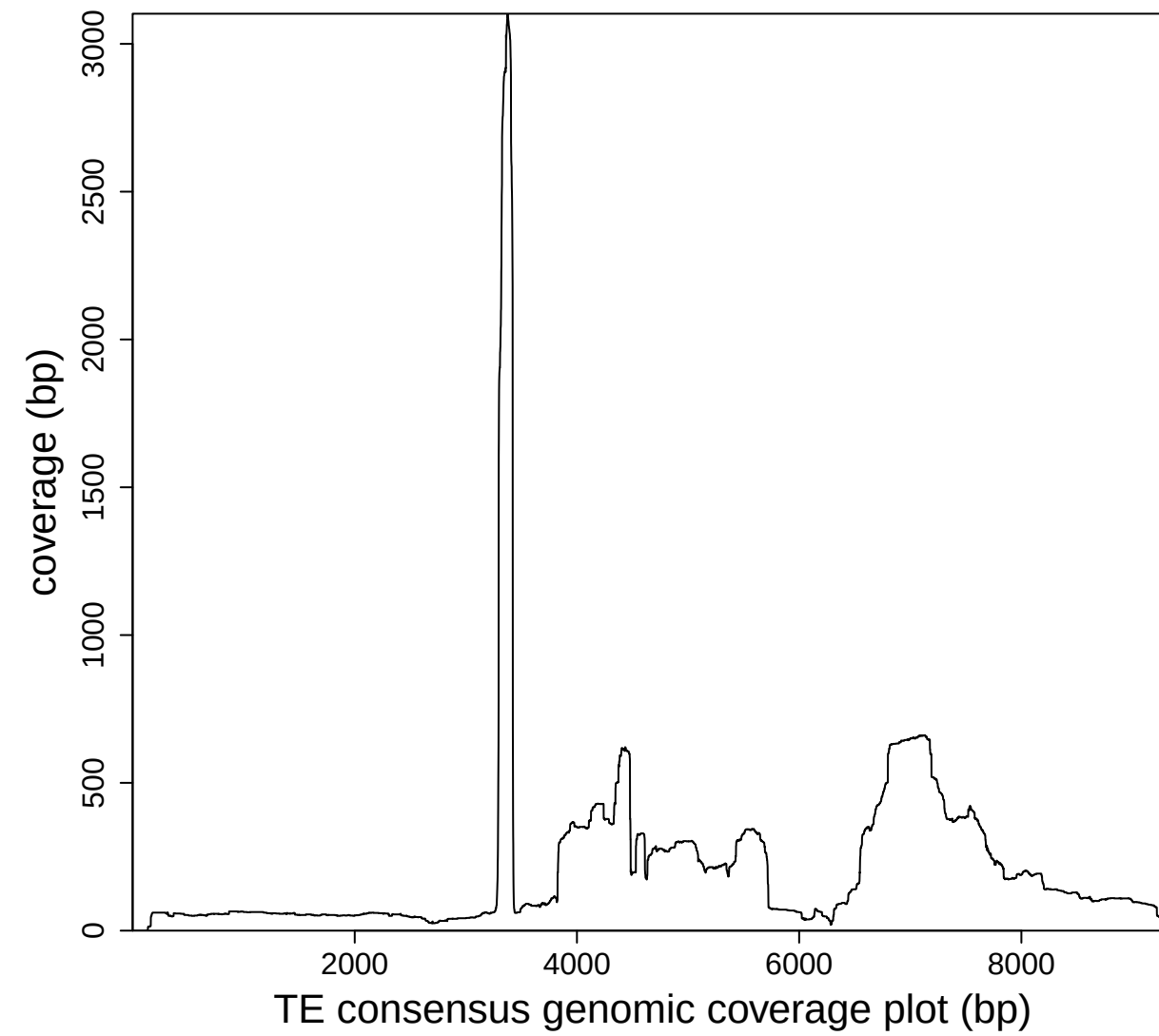
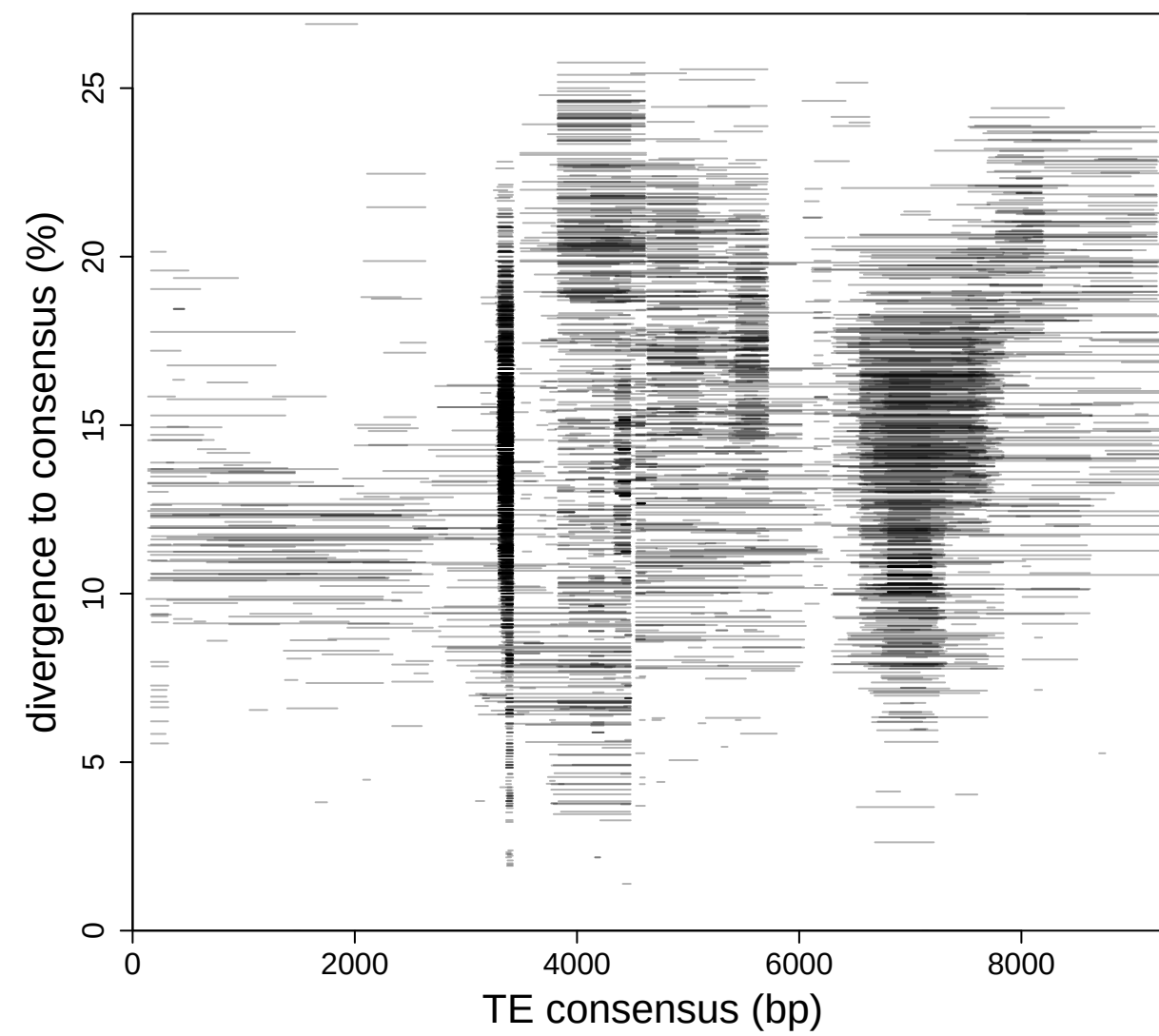






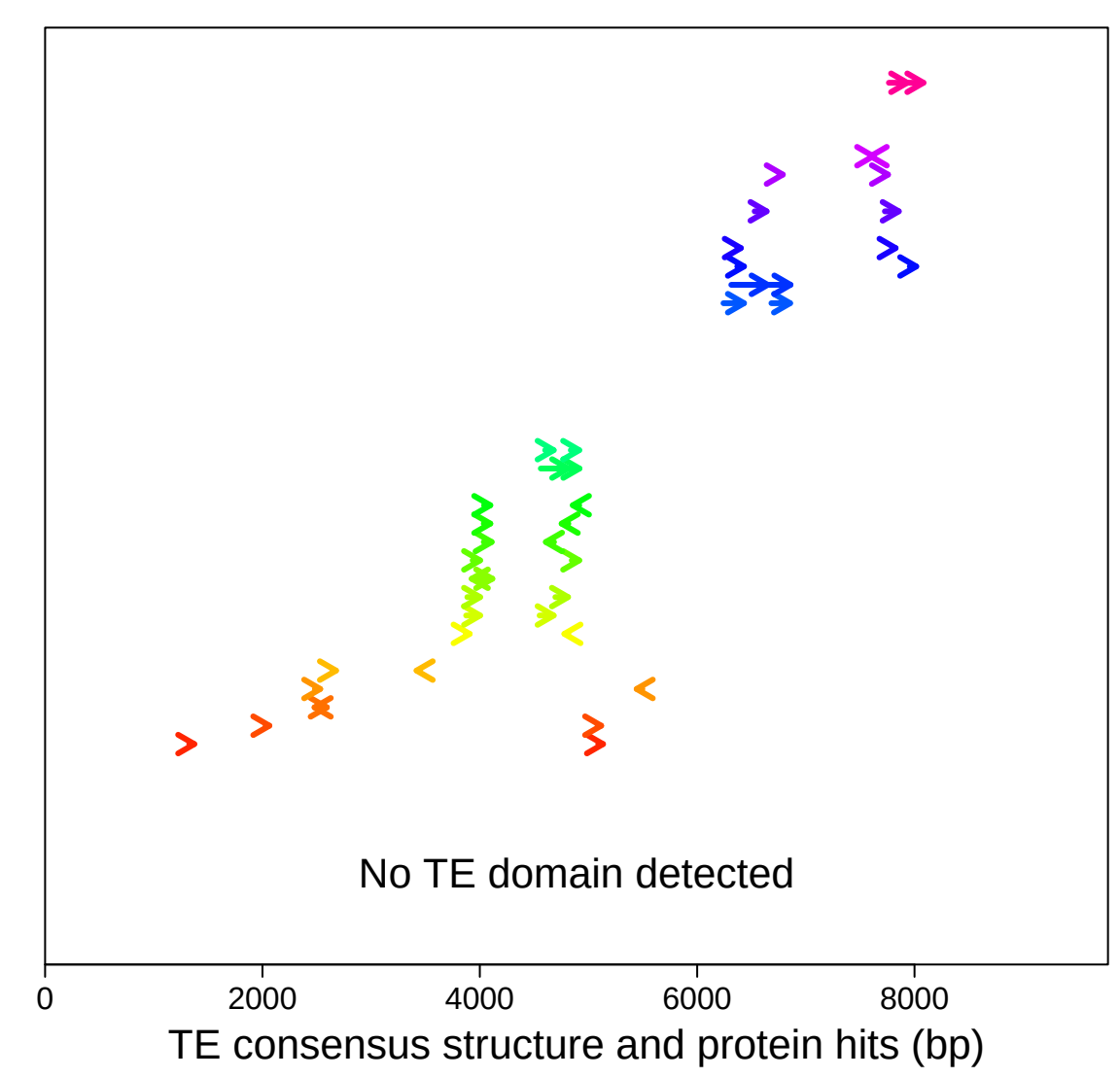
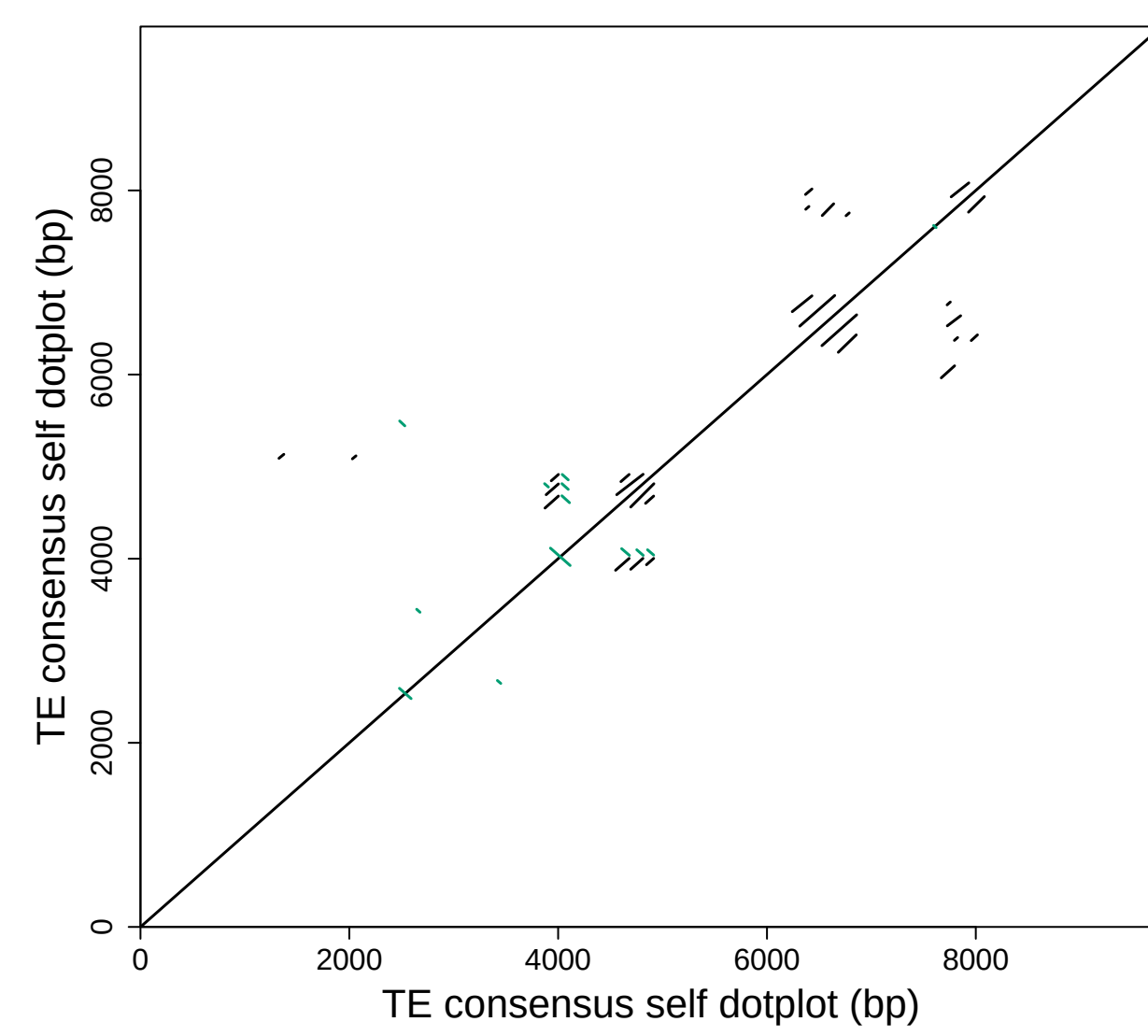
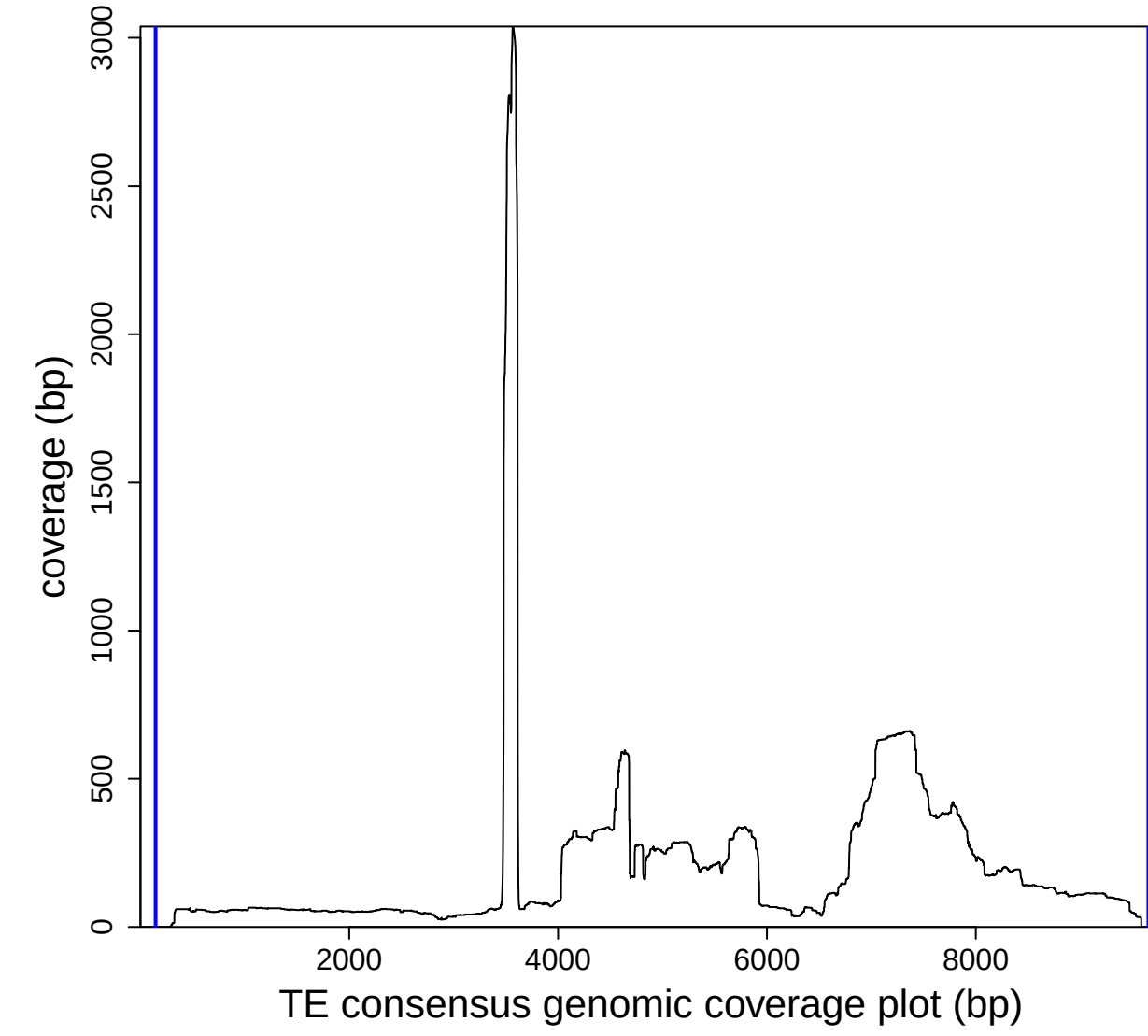
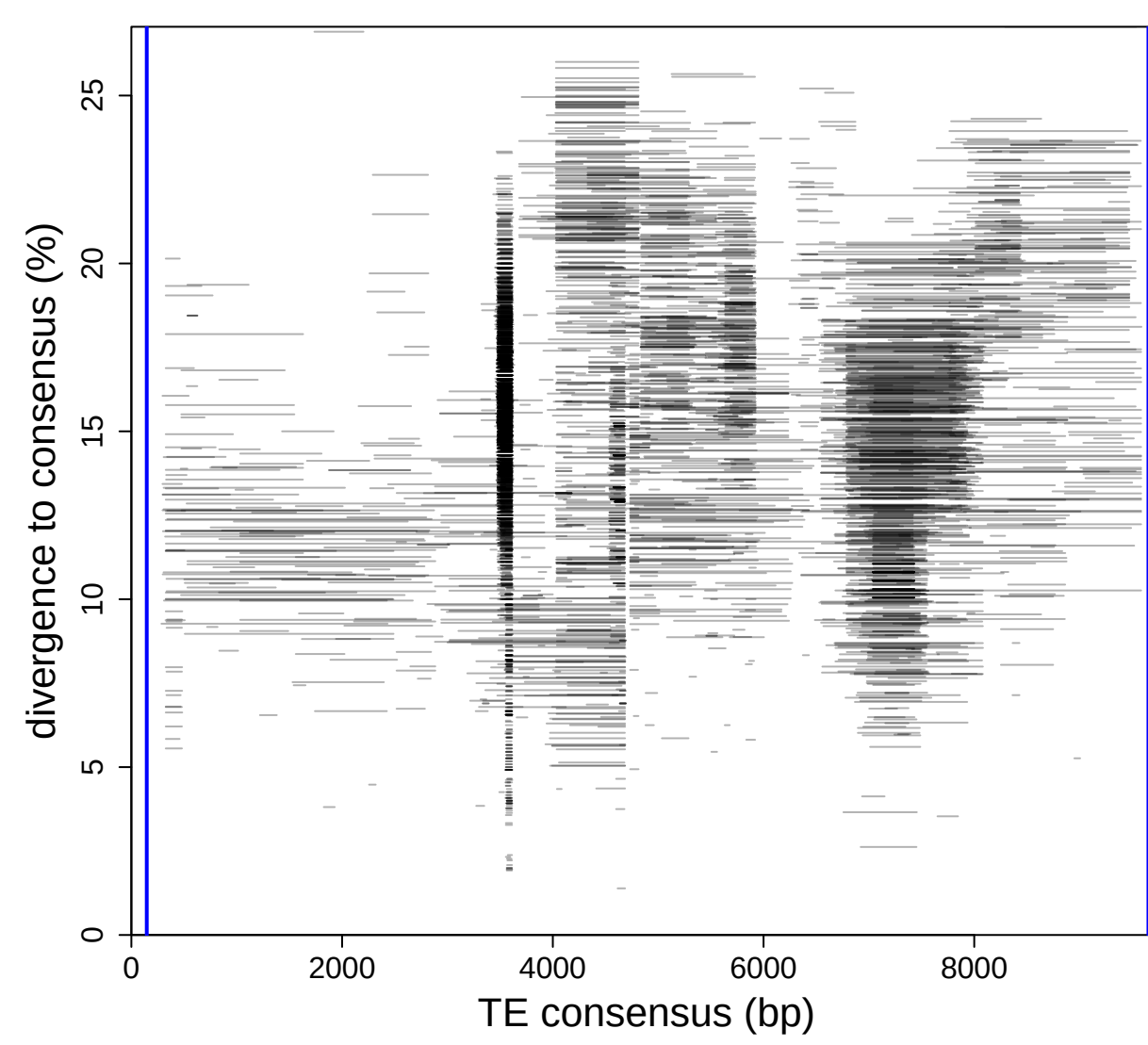
a.b.bed_uf.bed_g_2.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.f
size: 9364bp; fragments: 5895; full length: 0 (>=8427.6bp)

After TEtrimmer 9364 bp



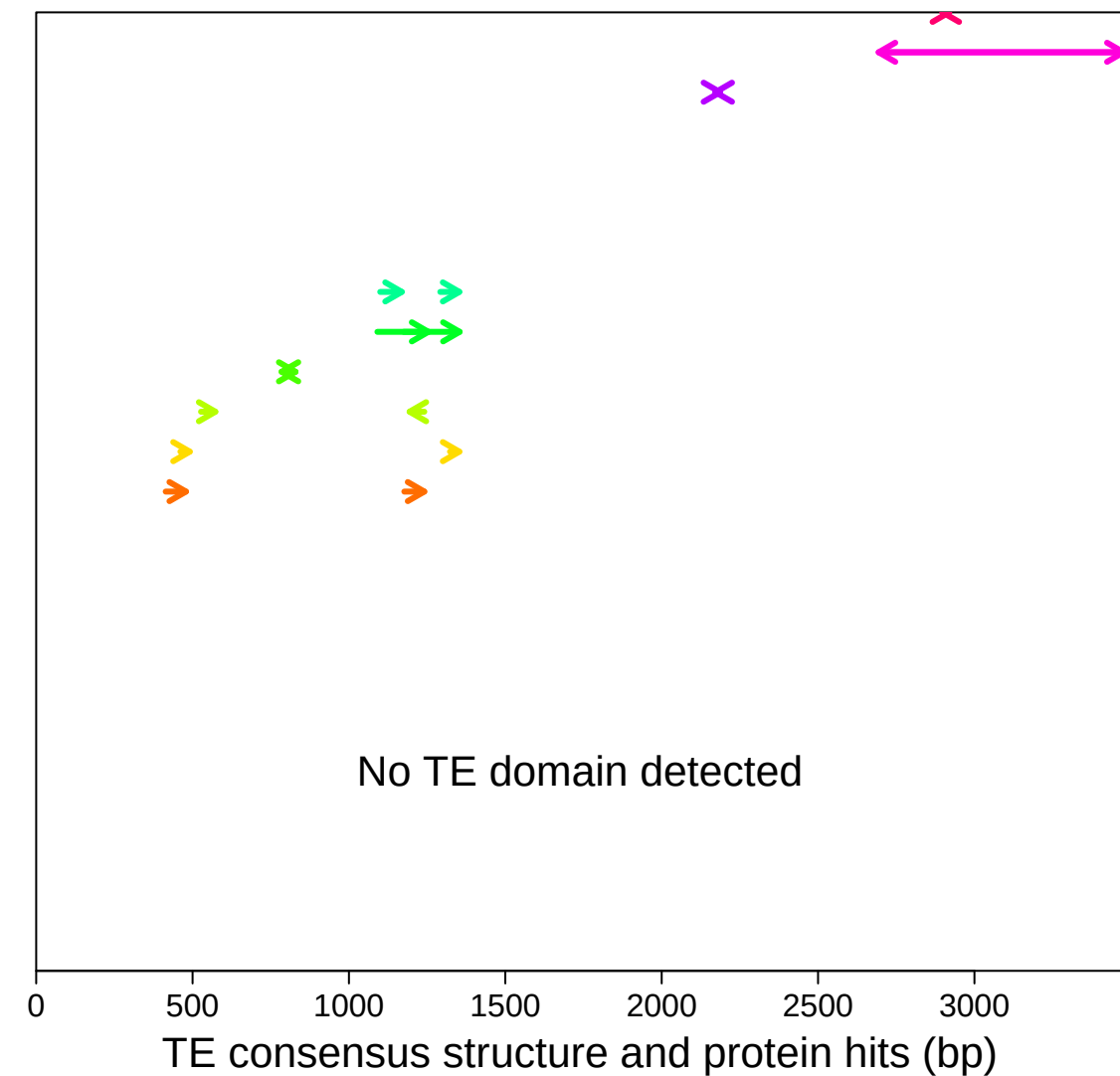
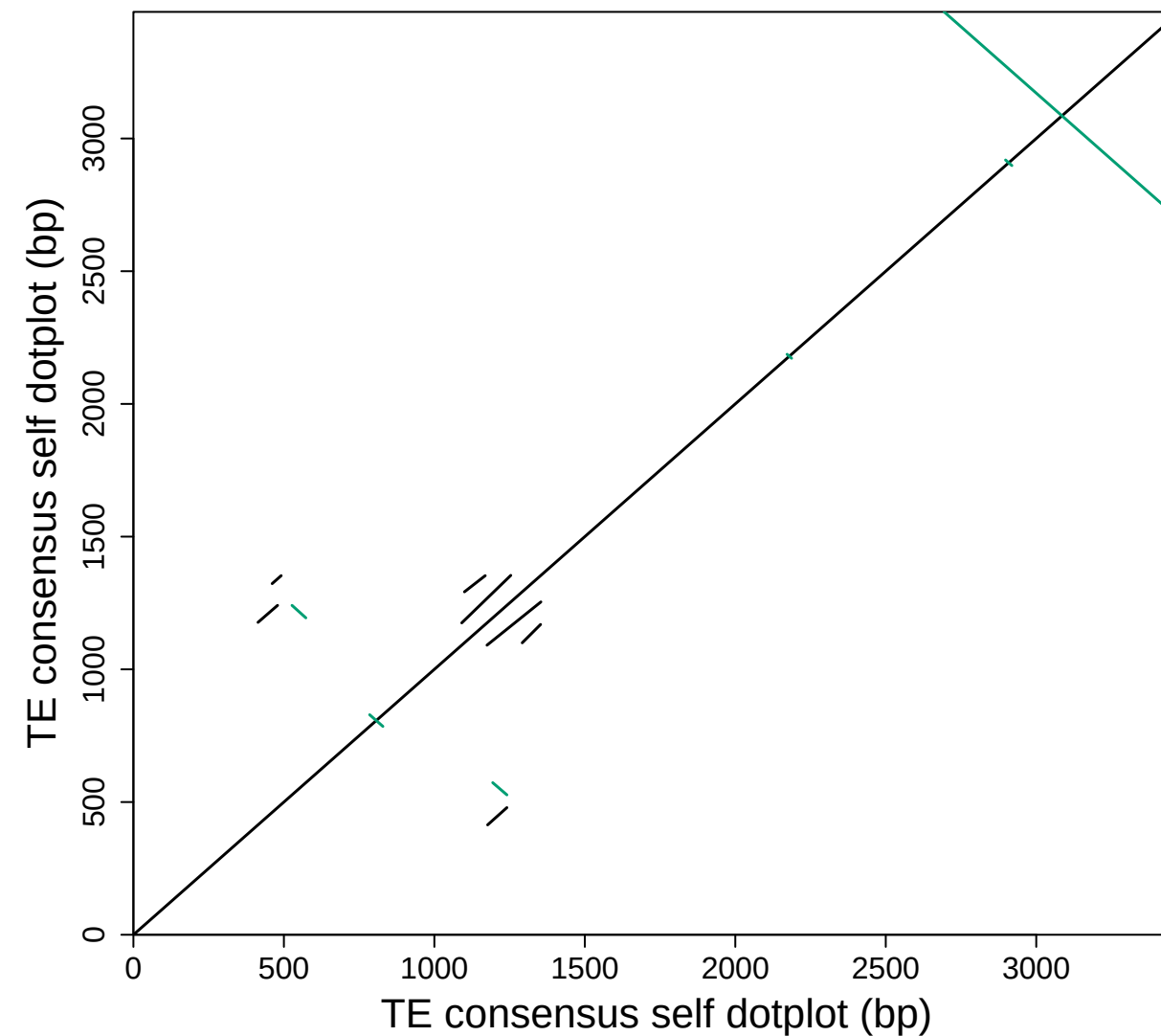
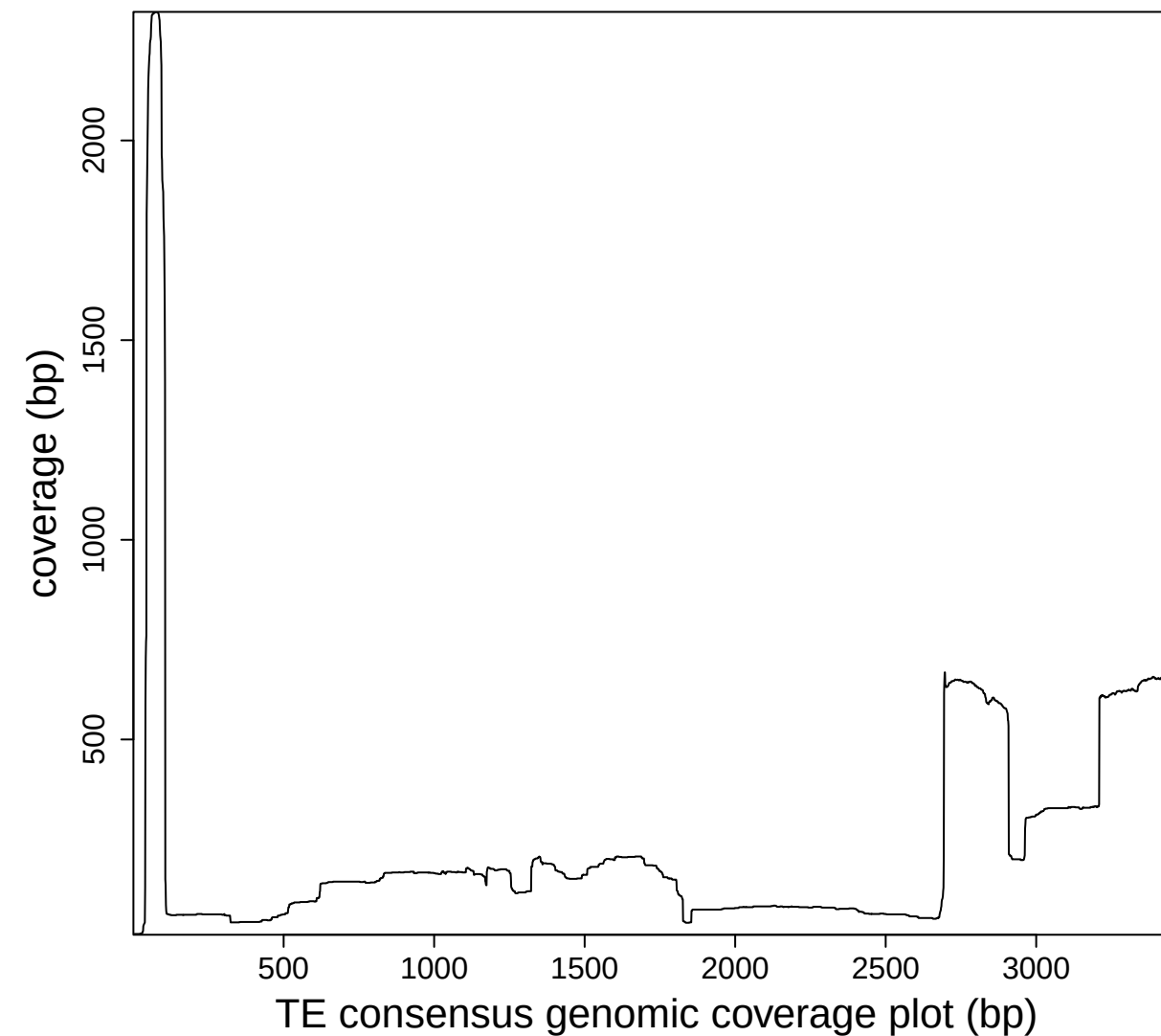
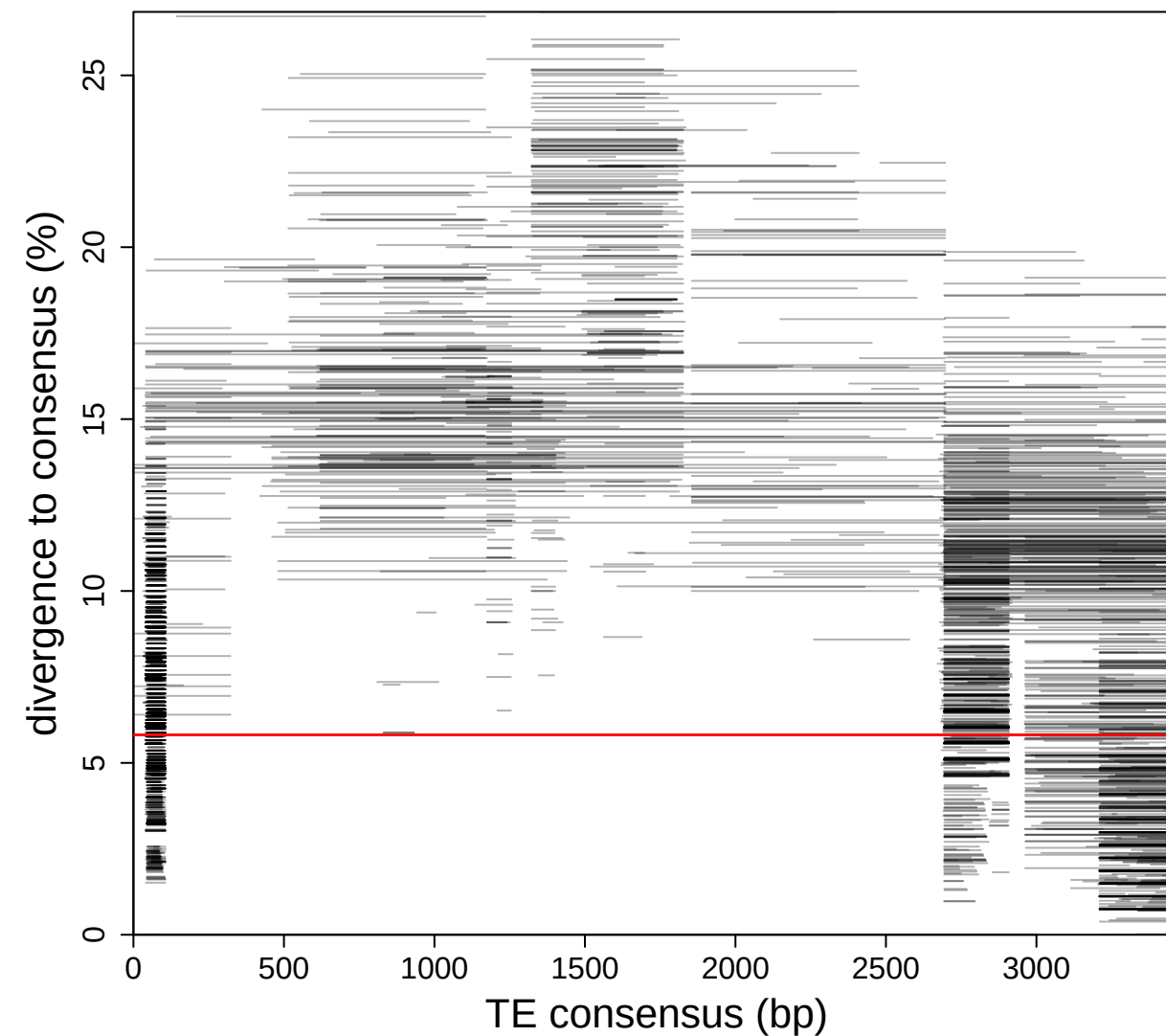
9.fasta.b.bed uf.bed g_2.bed fm_1.bed 0 0 bcln.fa aln.fa cl.f
size: 9780bp; fragments: 5669; full length: 0 (>=8802bp)

After TEtrimmer Extended plot Blue lines are boundaries

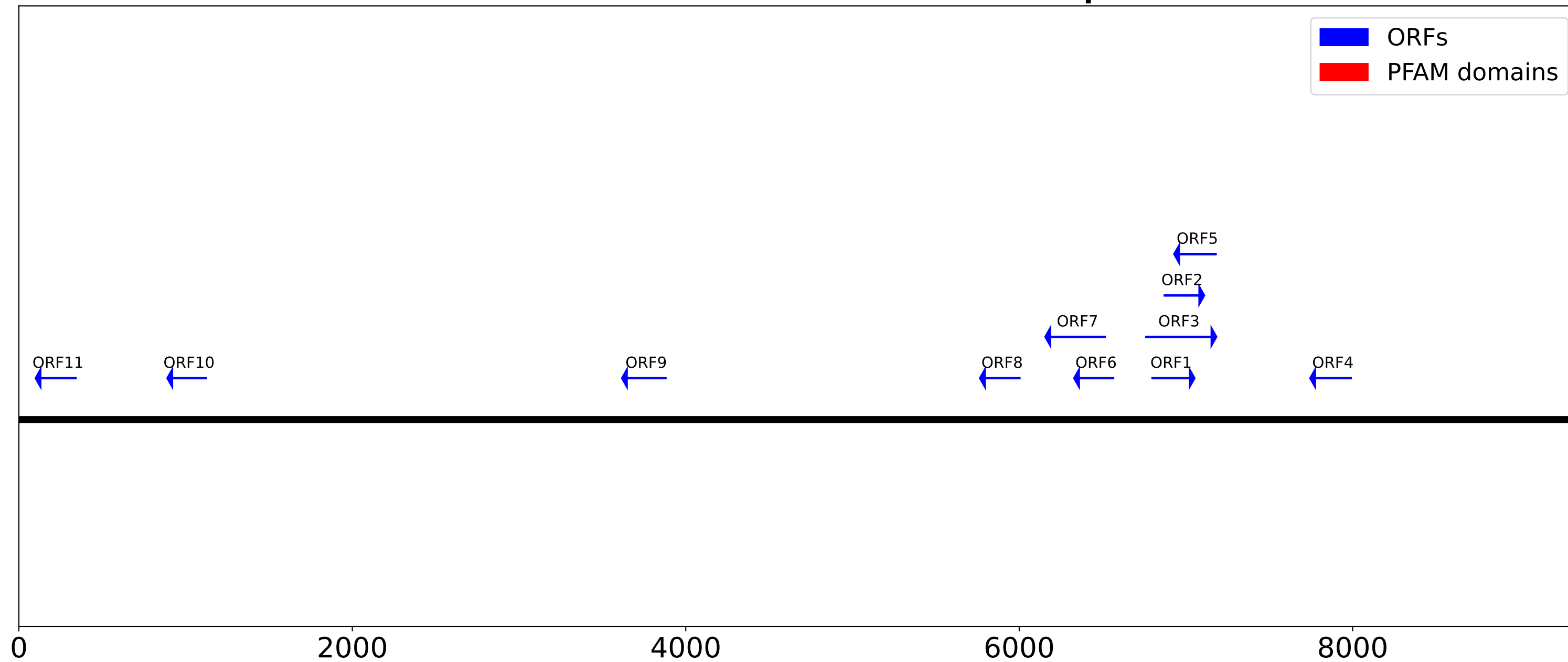


TE: rnd_5_family_2589
size: 3477bp; fragments: 4018; full length: 1 (≥ 3129.3 bp)

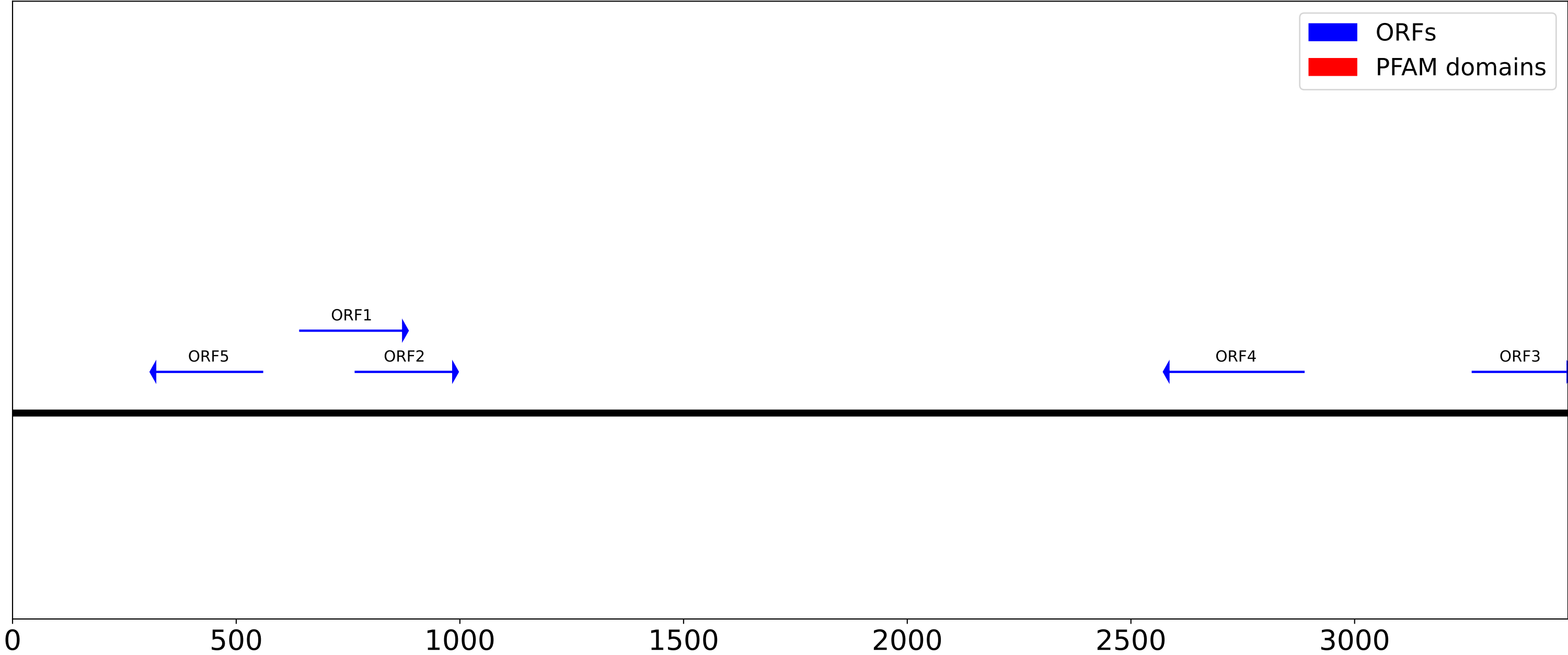
Before TEtrimmer 3477 bp



After TEtrimmer ORF and PFAM domain plot



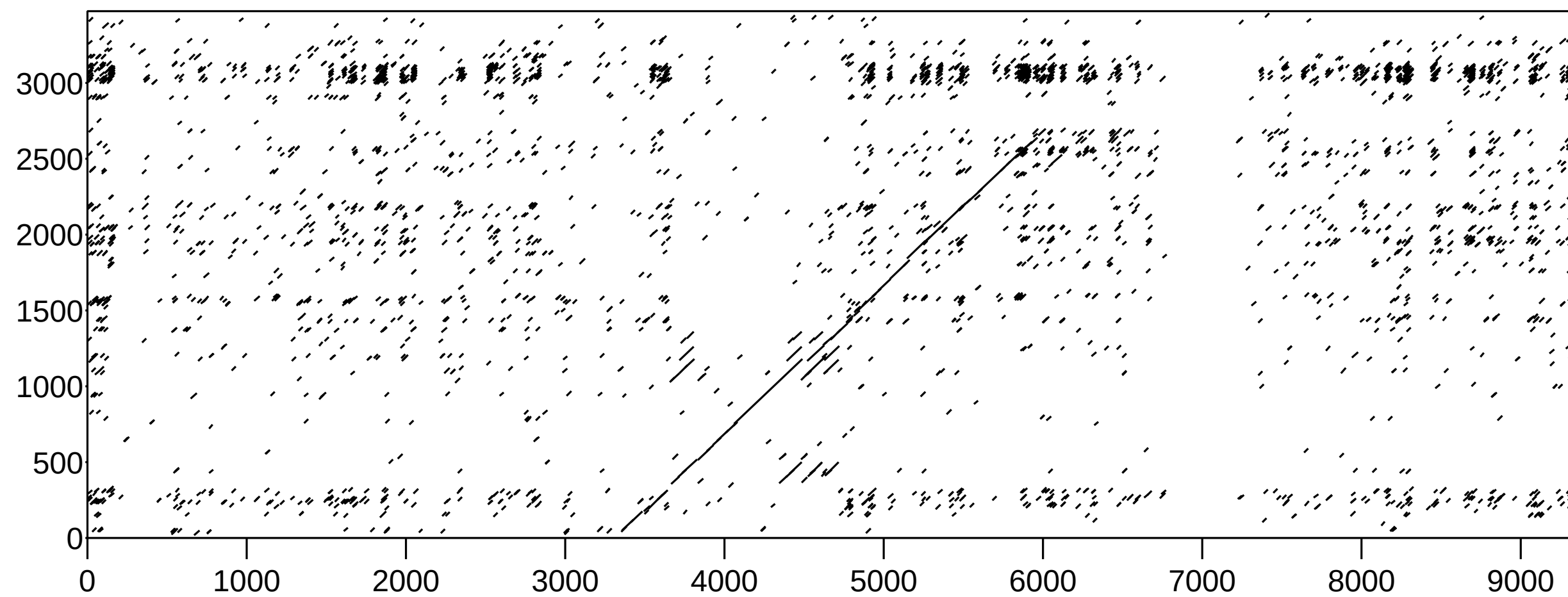
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)