

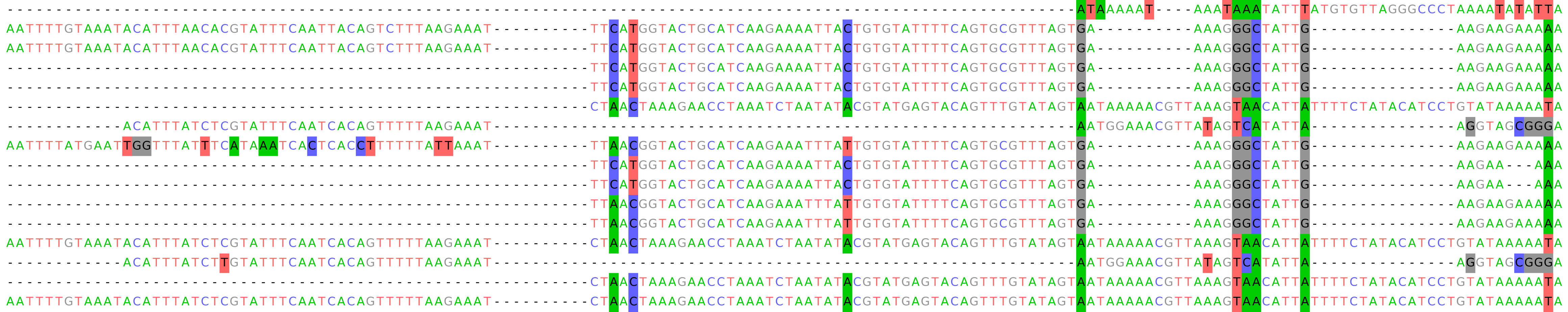
Start crop Point

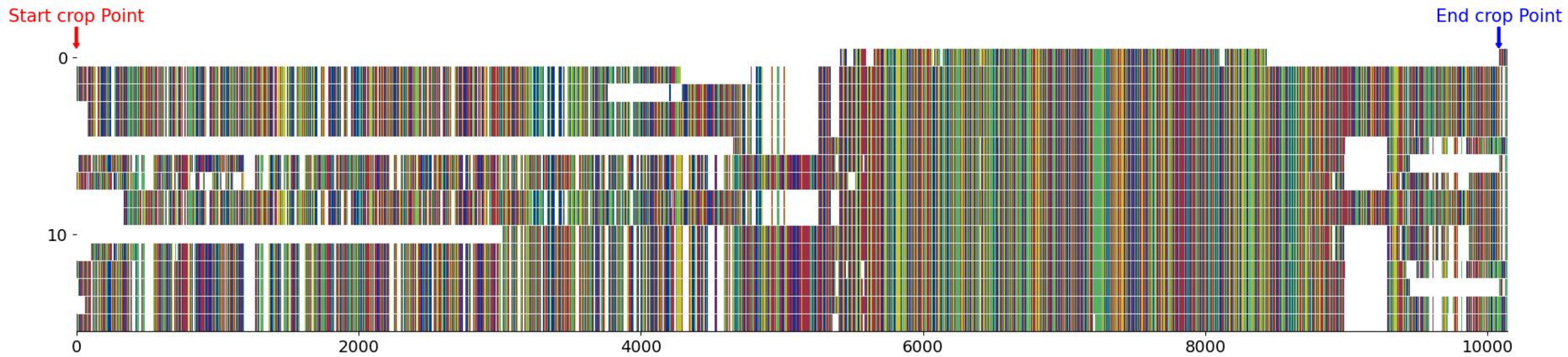


End crop Point



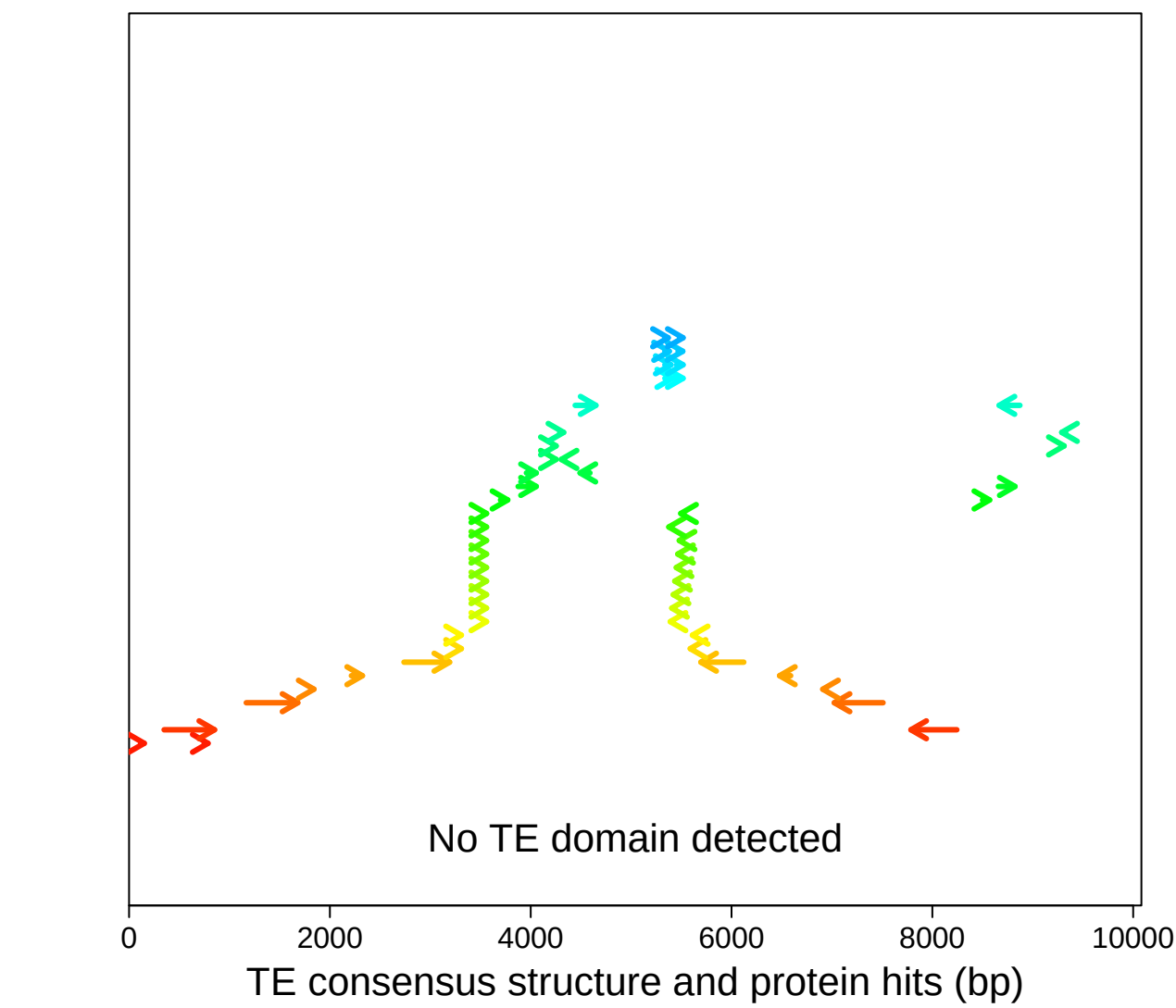
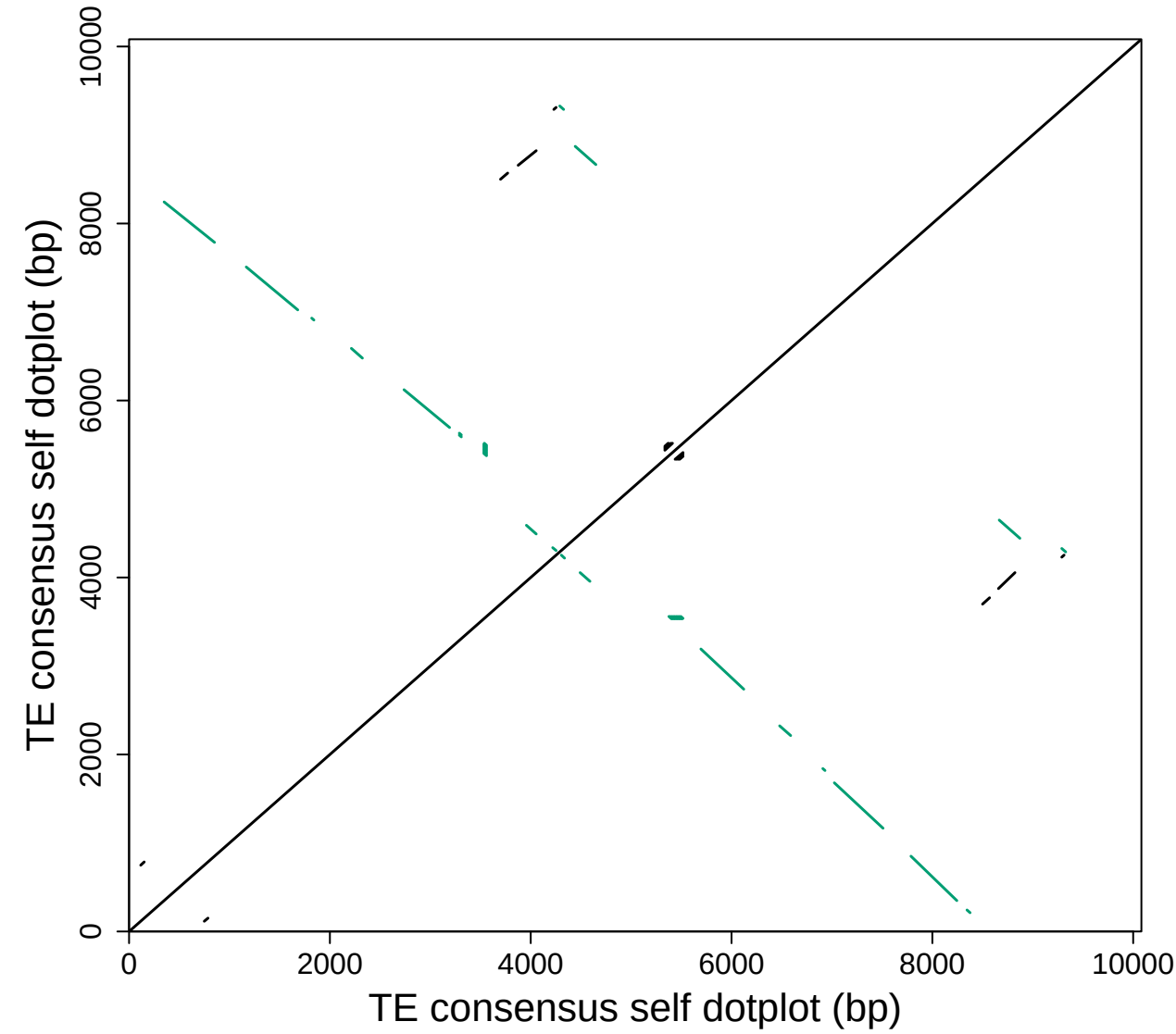
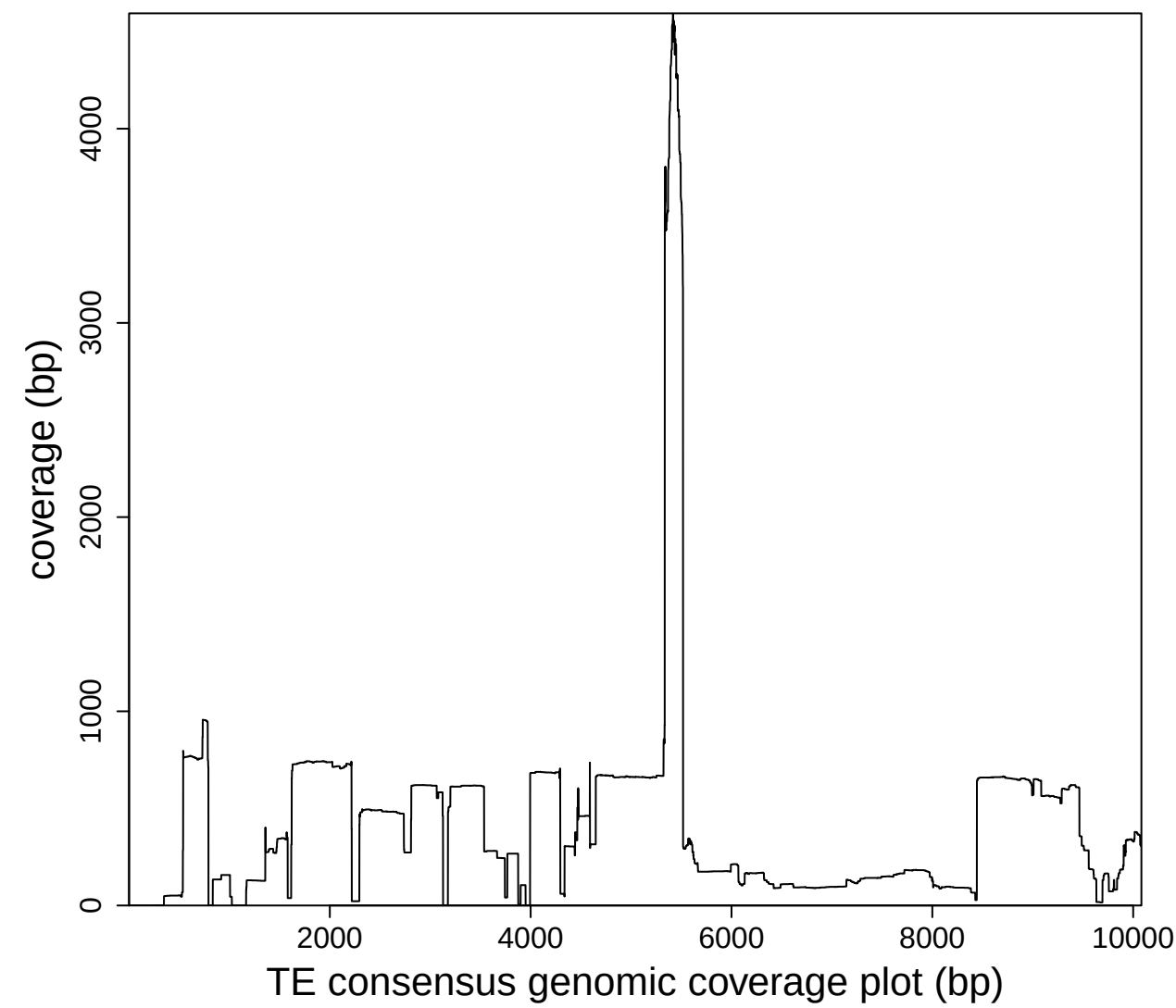
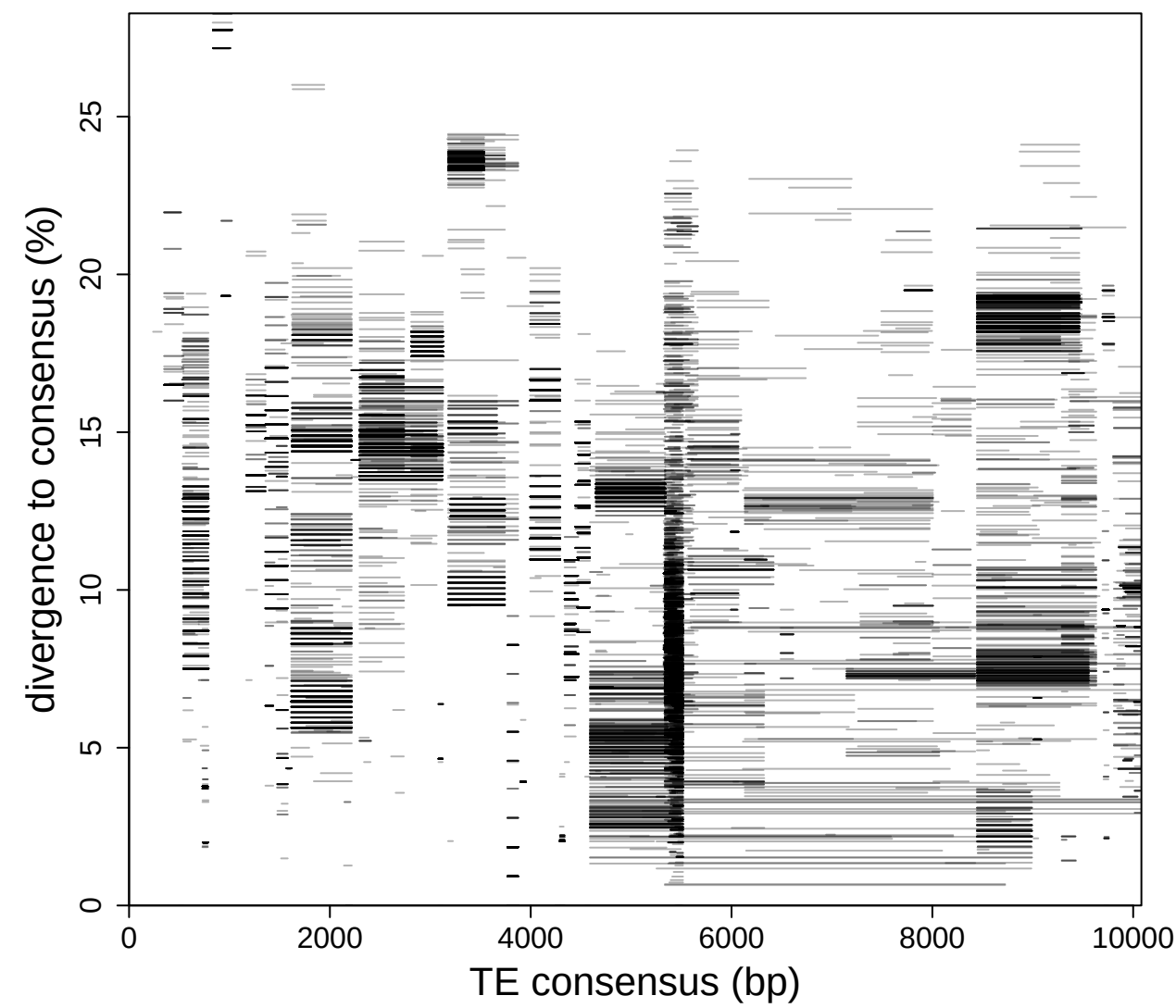
MSA length = 10082





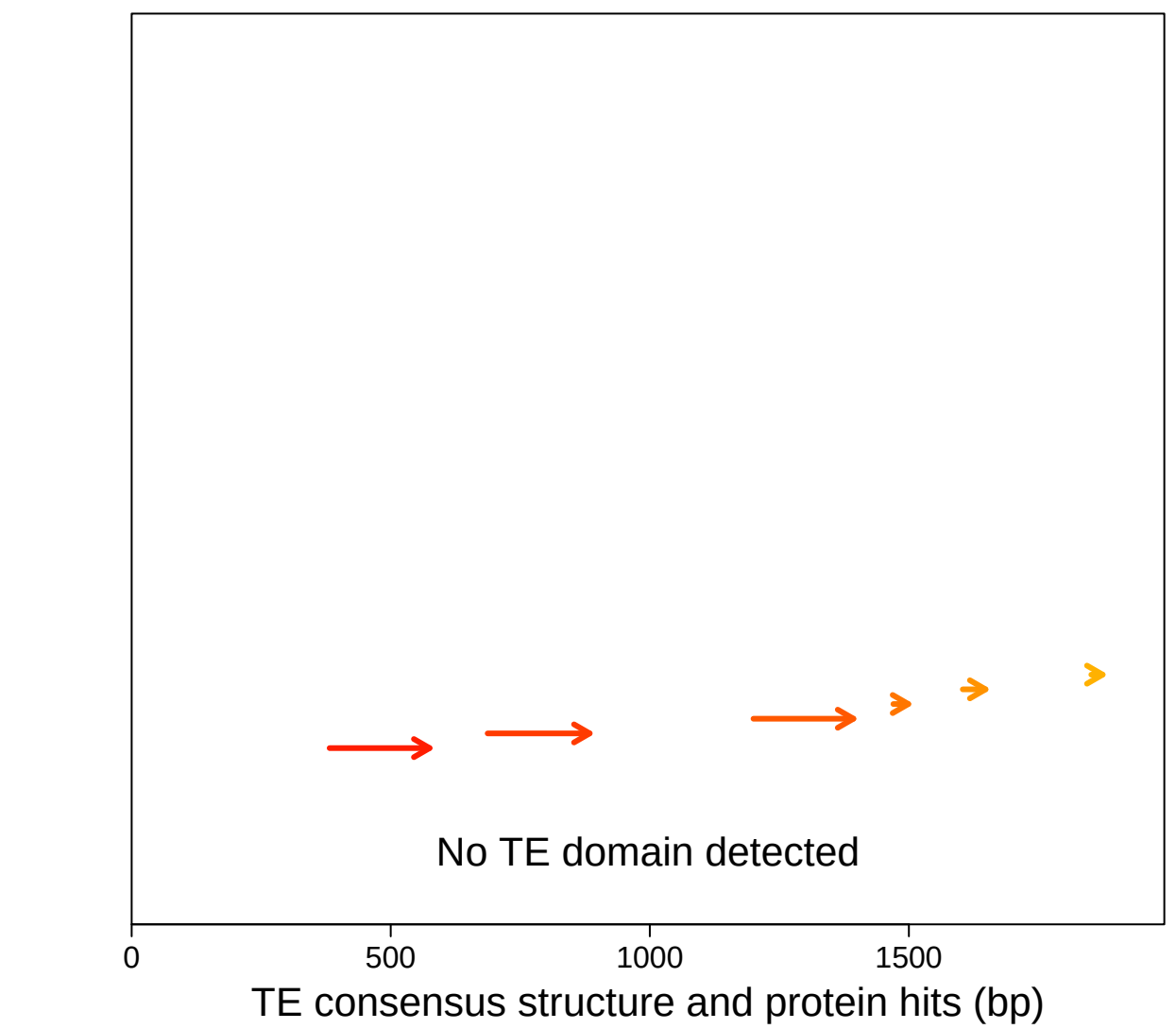
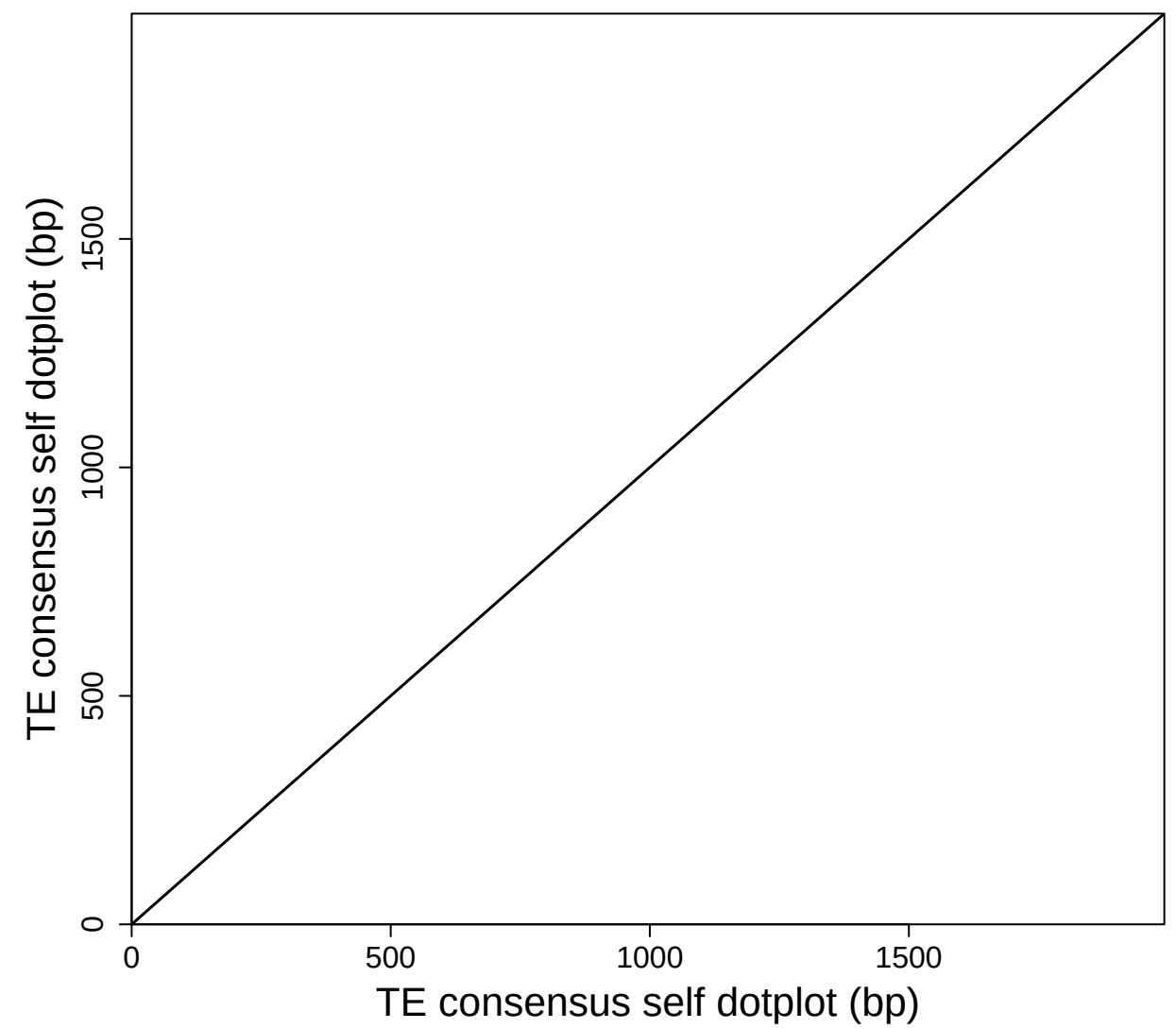
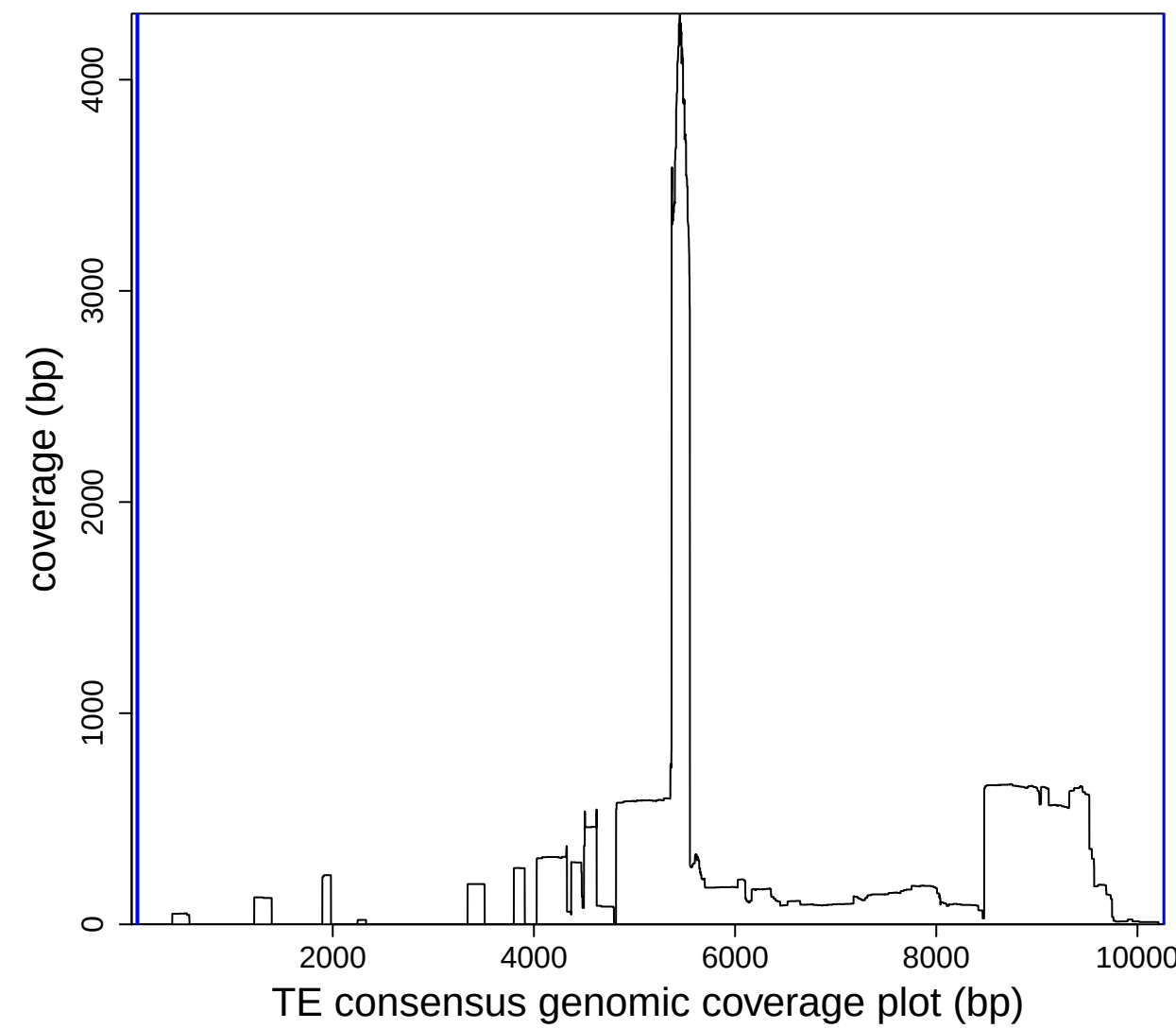
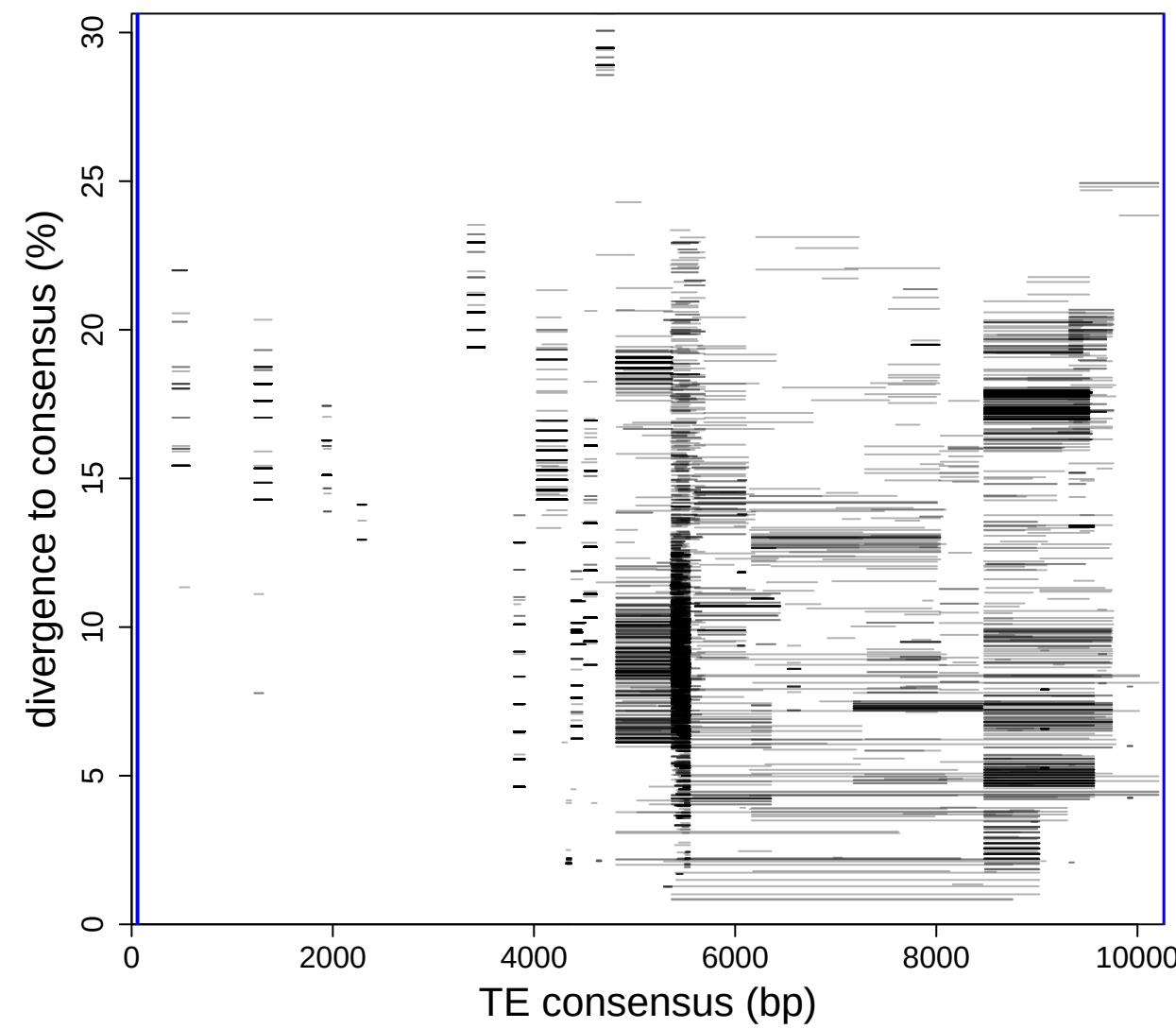
a.b.bed_uf.bed_g_3.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.f
size: 10082bp; fragments: 13193; full length: 0 (>=9073.8bp)

After TEtrimmer 10082 bp



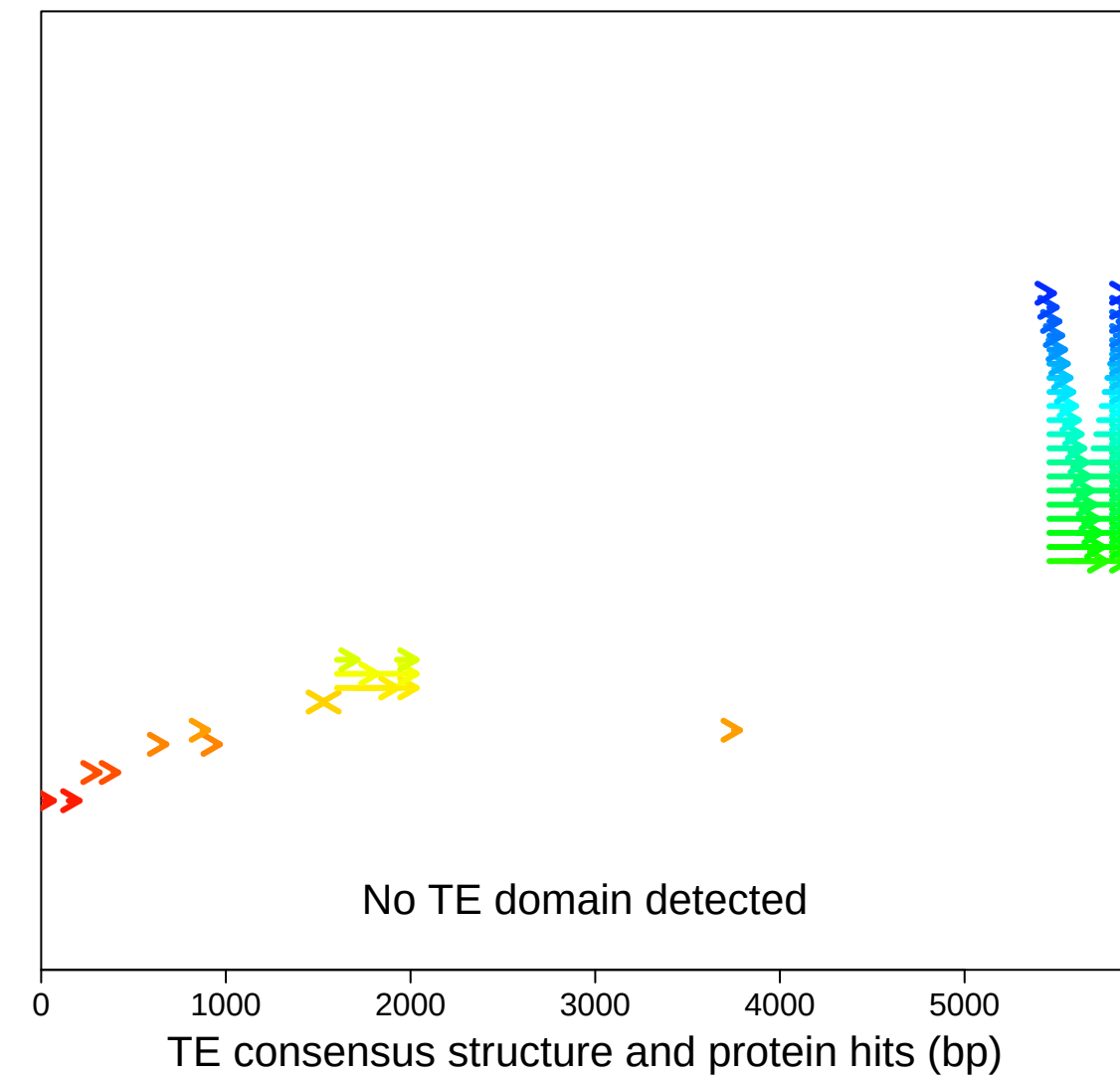
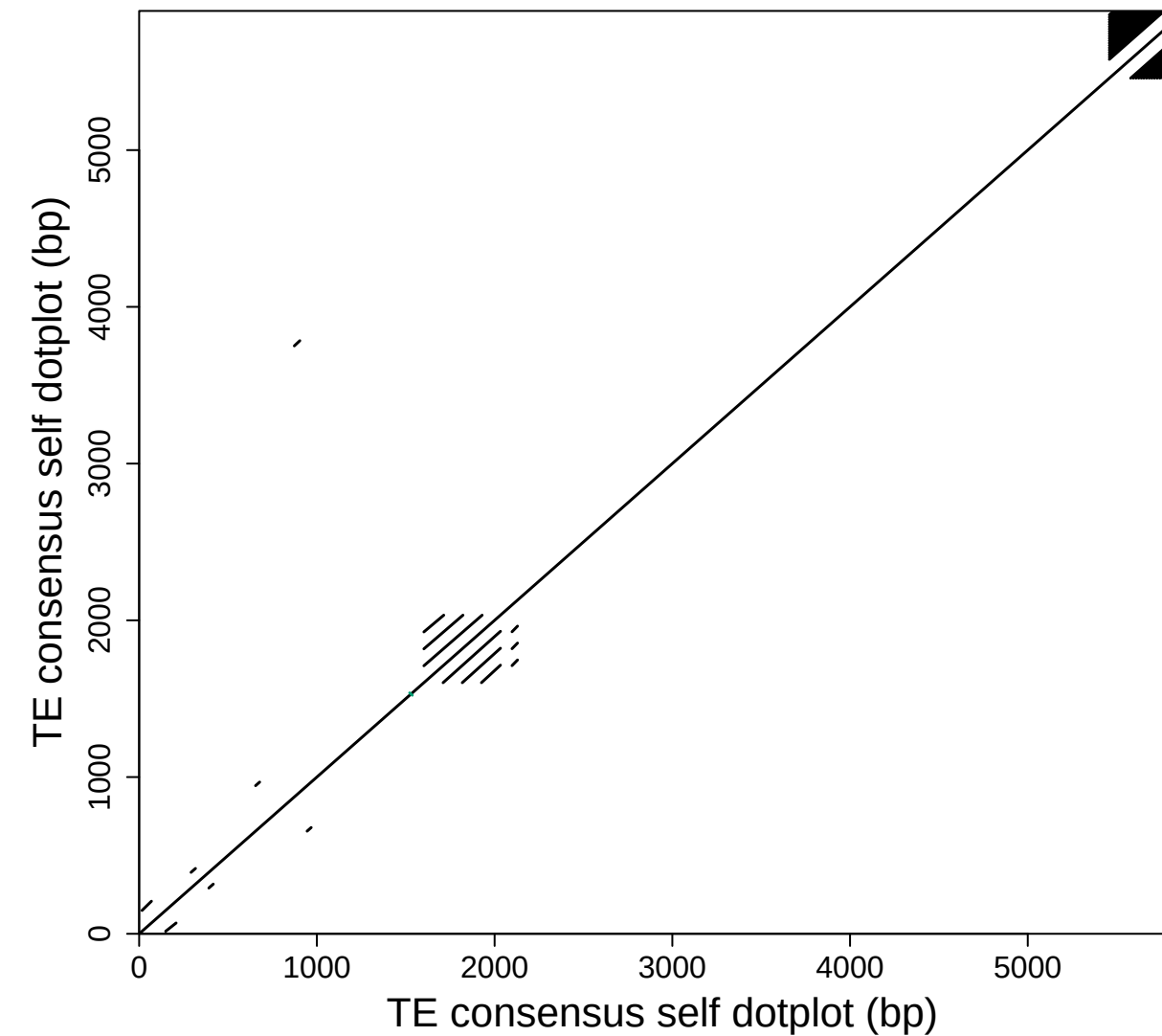
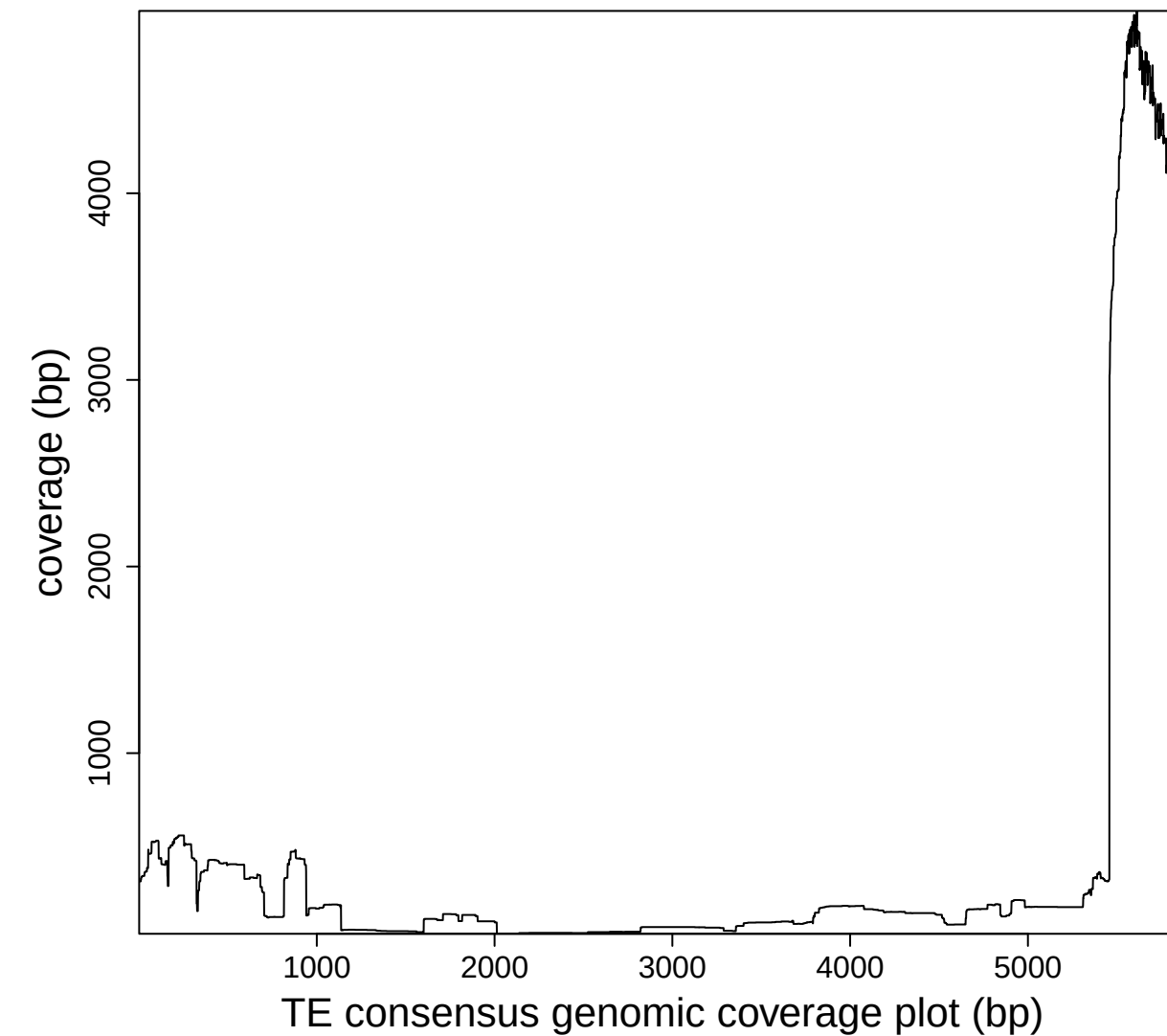
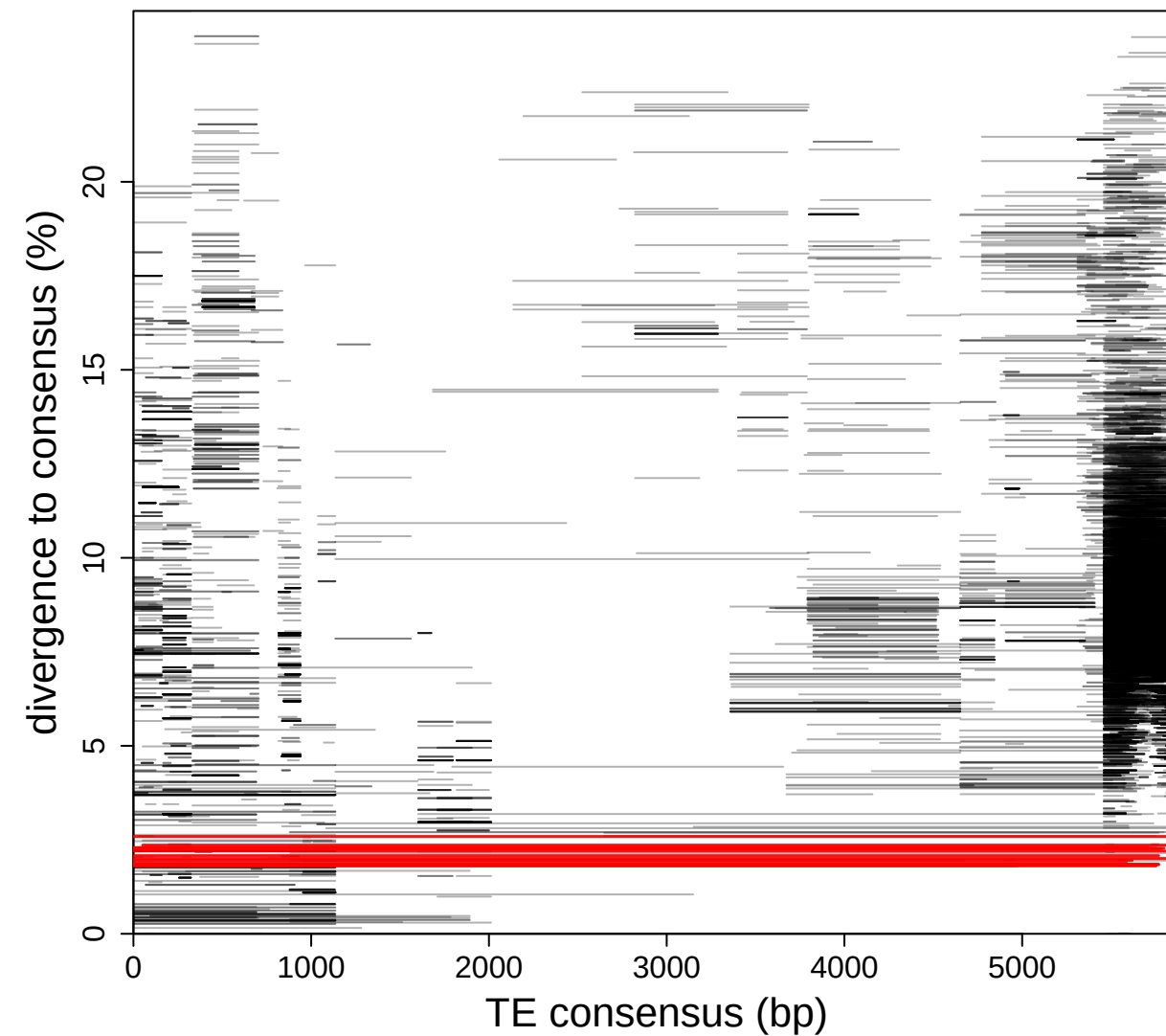
asta.b.bed_uf.bed_g_3.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_c
size: 10267bp; fragments: 8410; full length: 0 (>=9240.3bp)

After TEtrimmer Extended plot Blue lines are boundaries

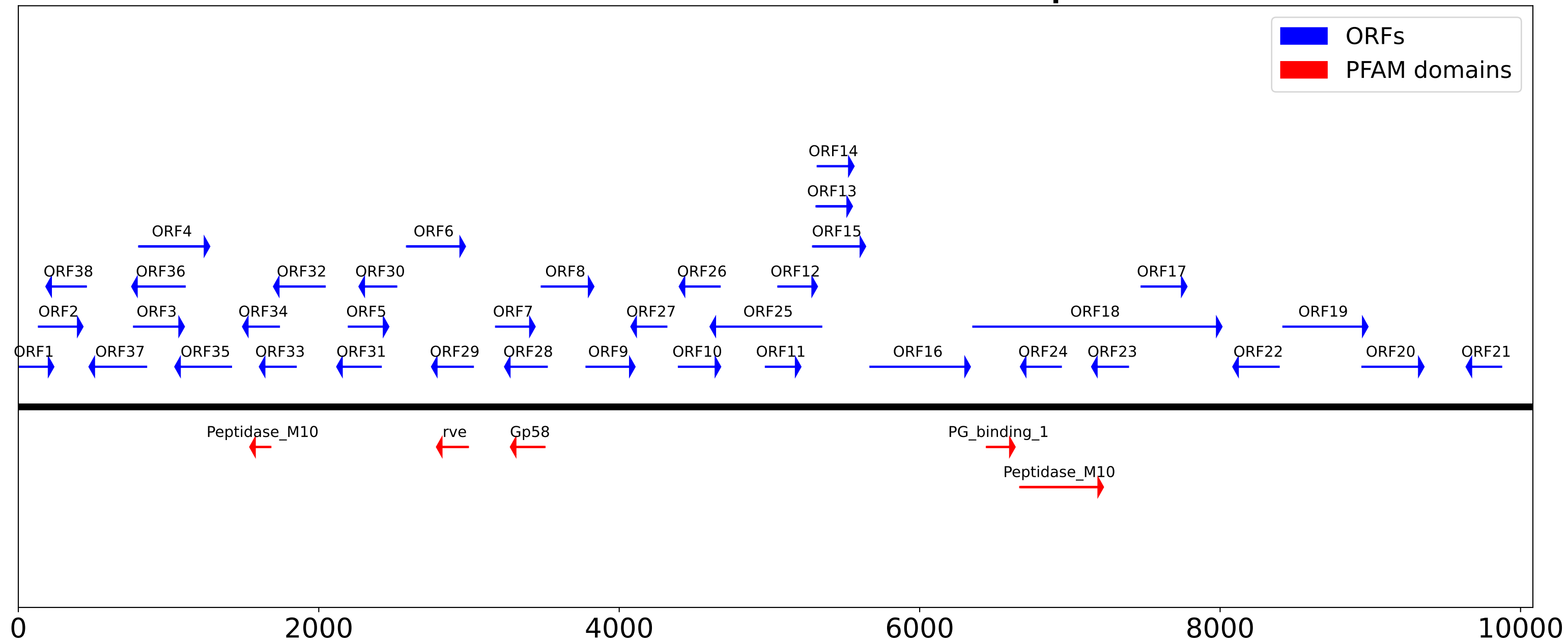


TE: rnd_5_family_7375
size: 5888bp; fragments: 10720; full length: 15 (≥ 5299.2 bp)

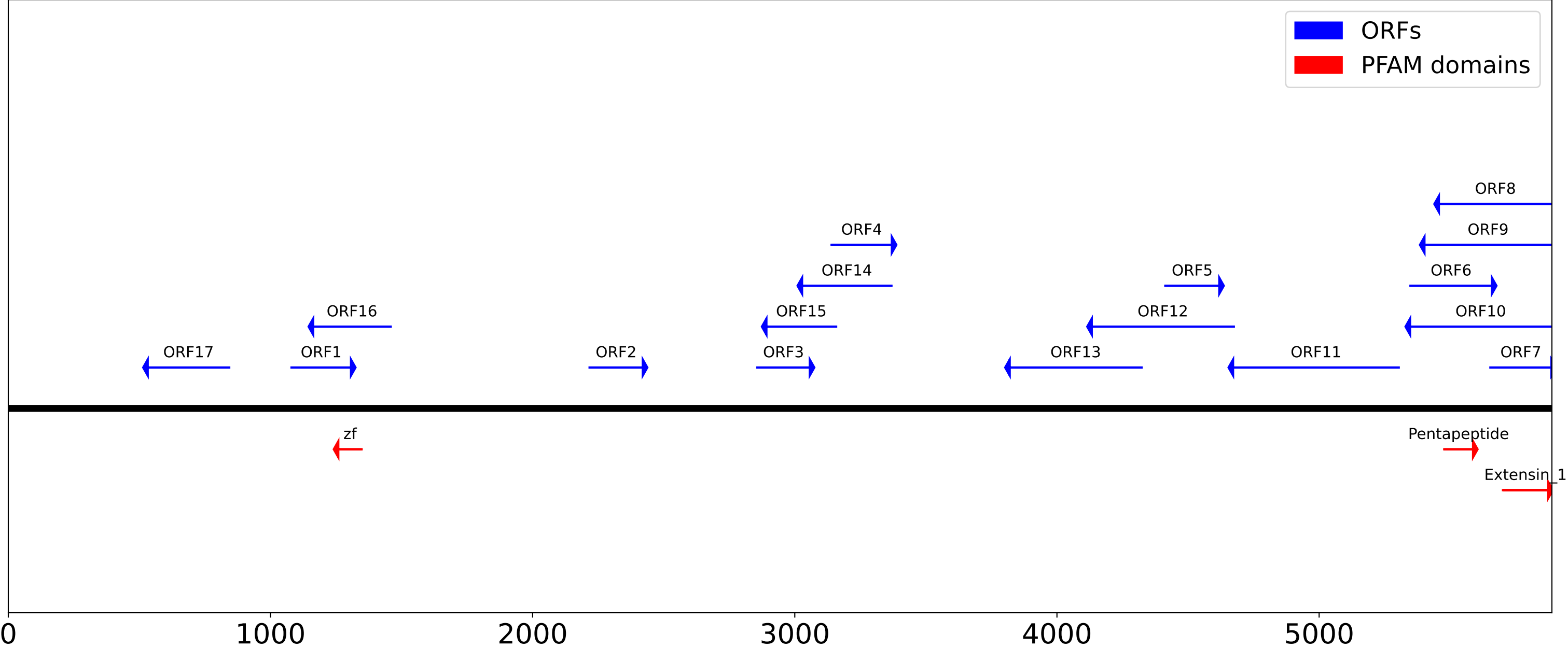
Before TEtrimmer 5888 bp



After TEtrimmer ORF and PFAM domain plot



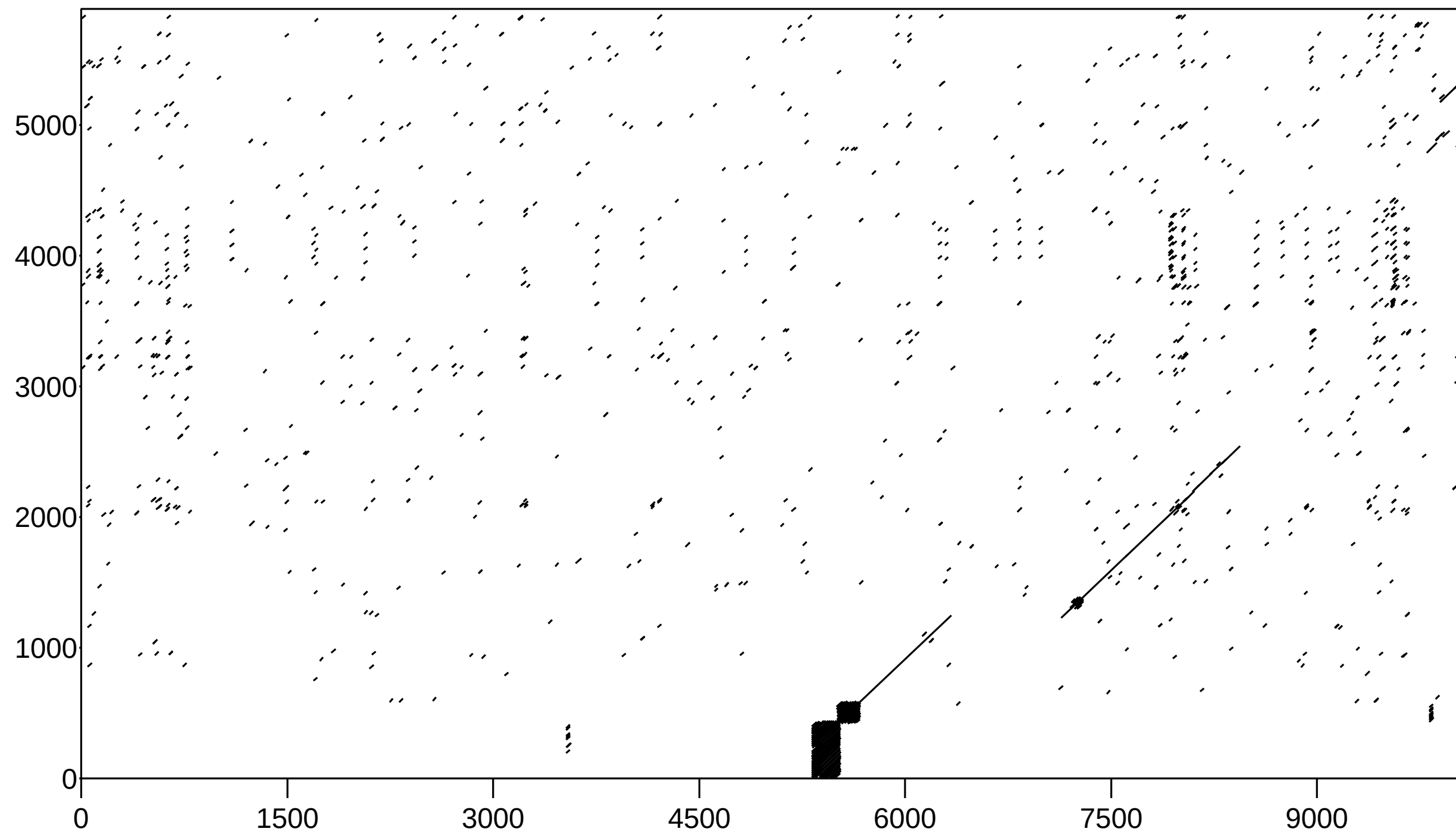
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)