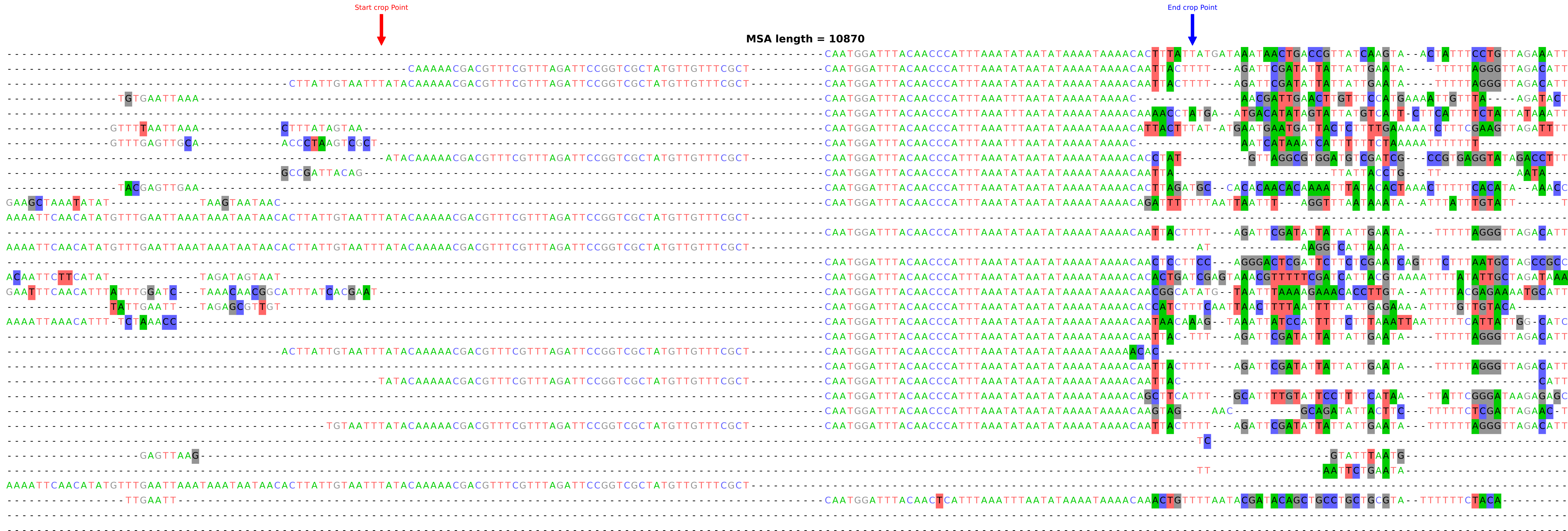
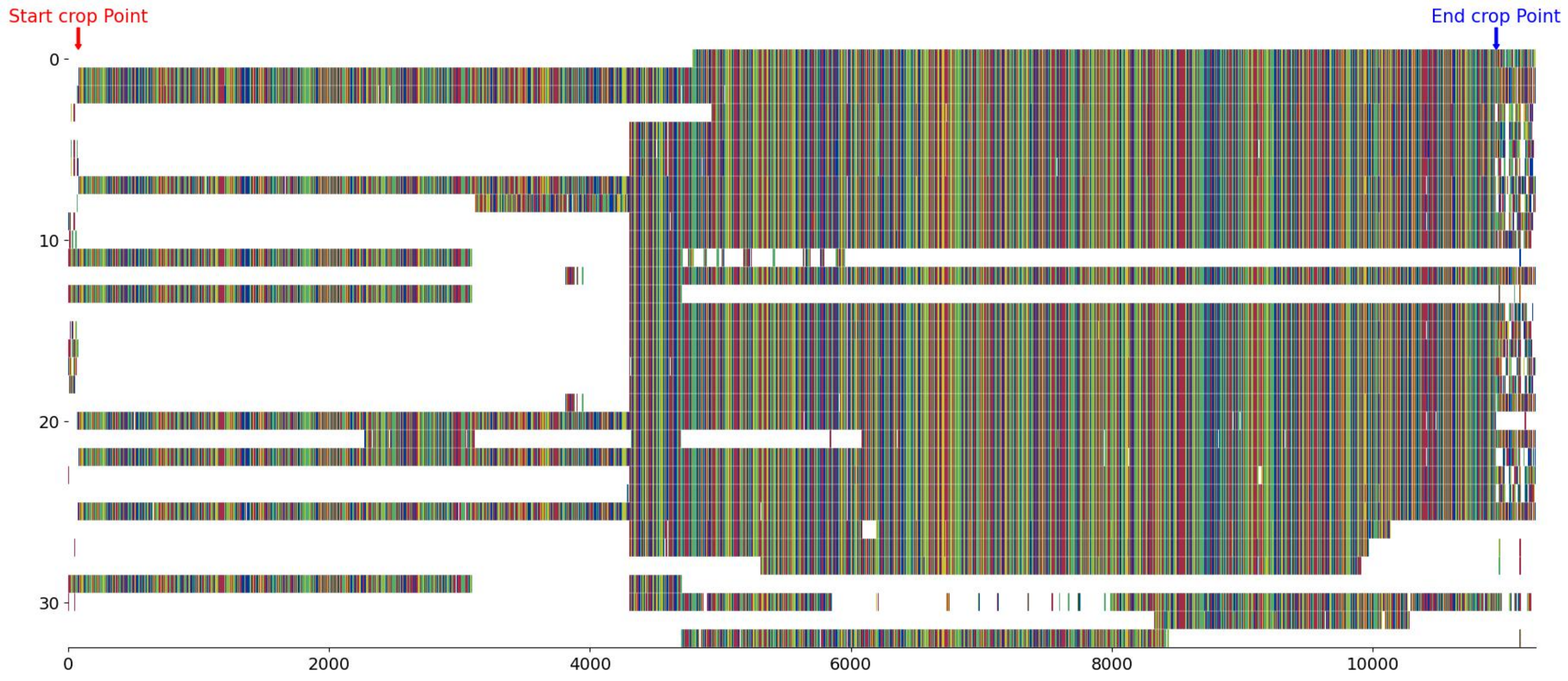
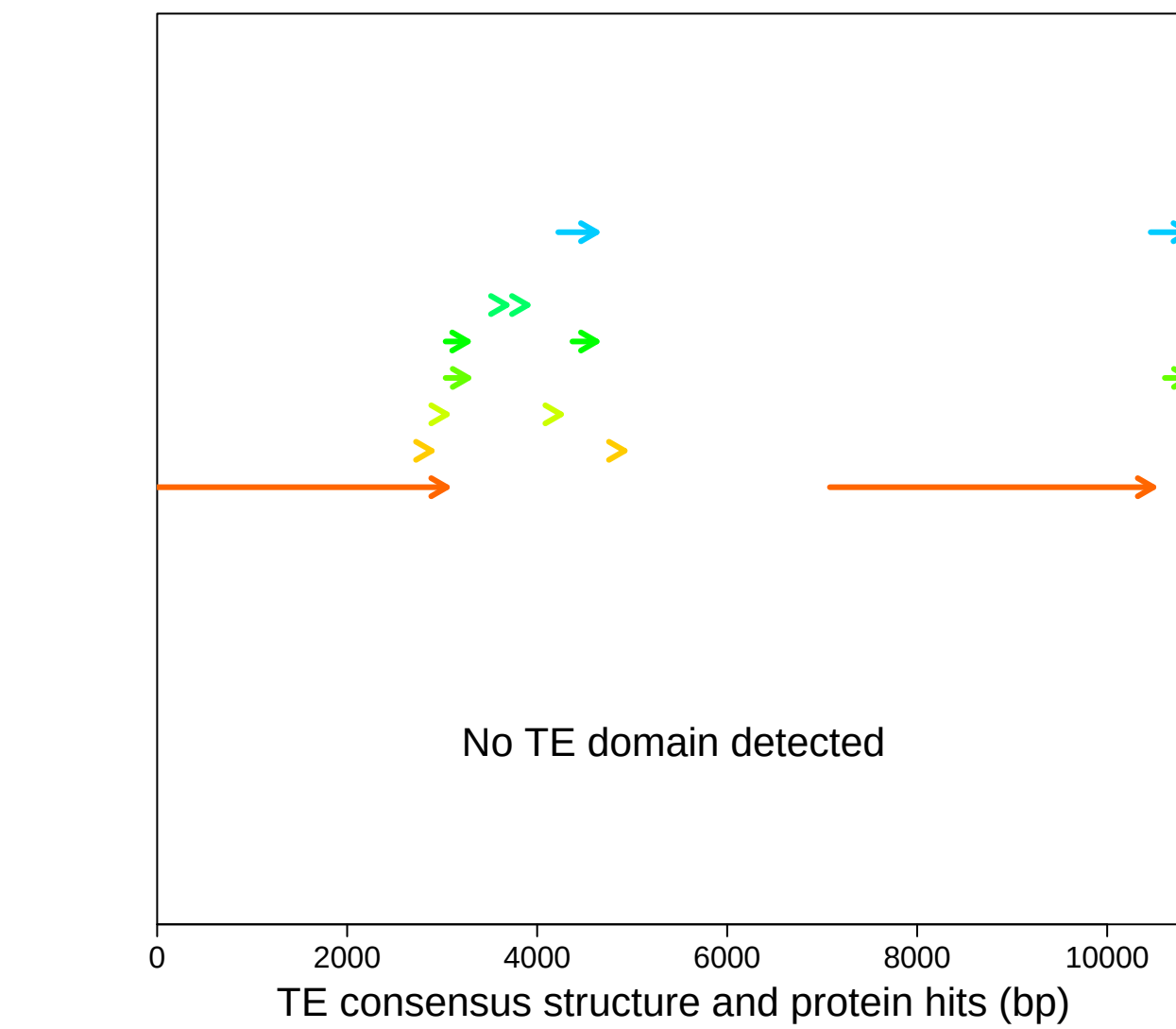
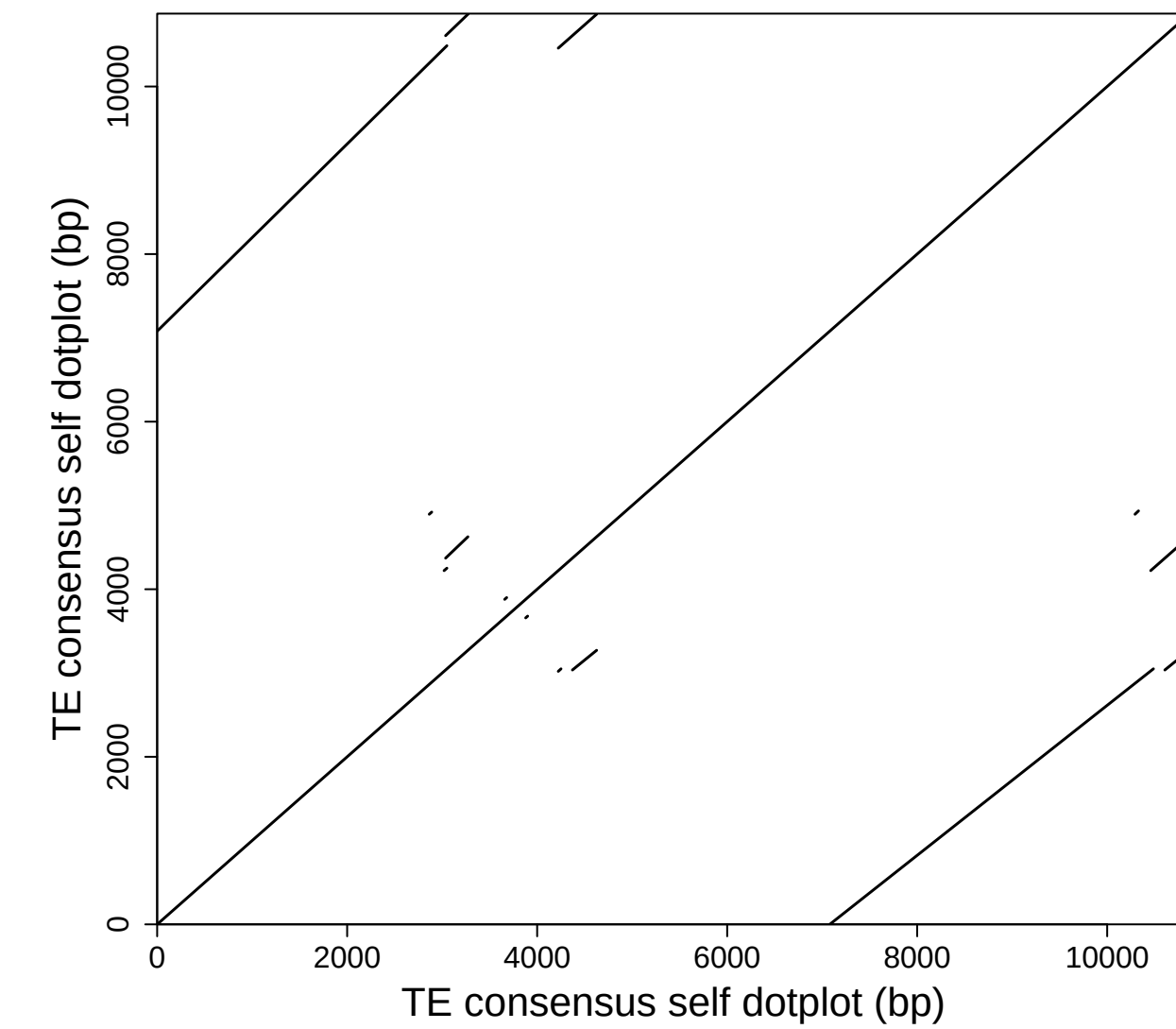
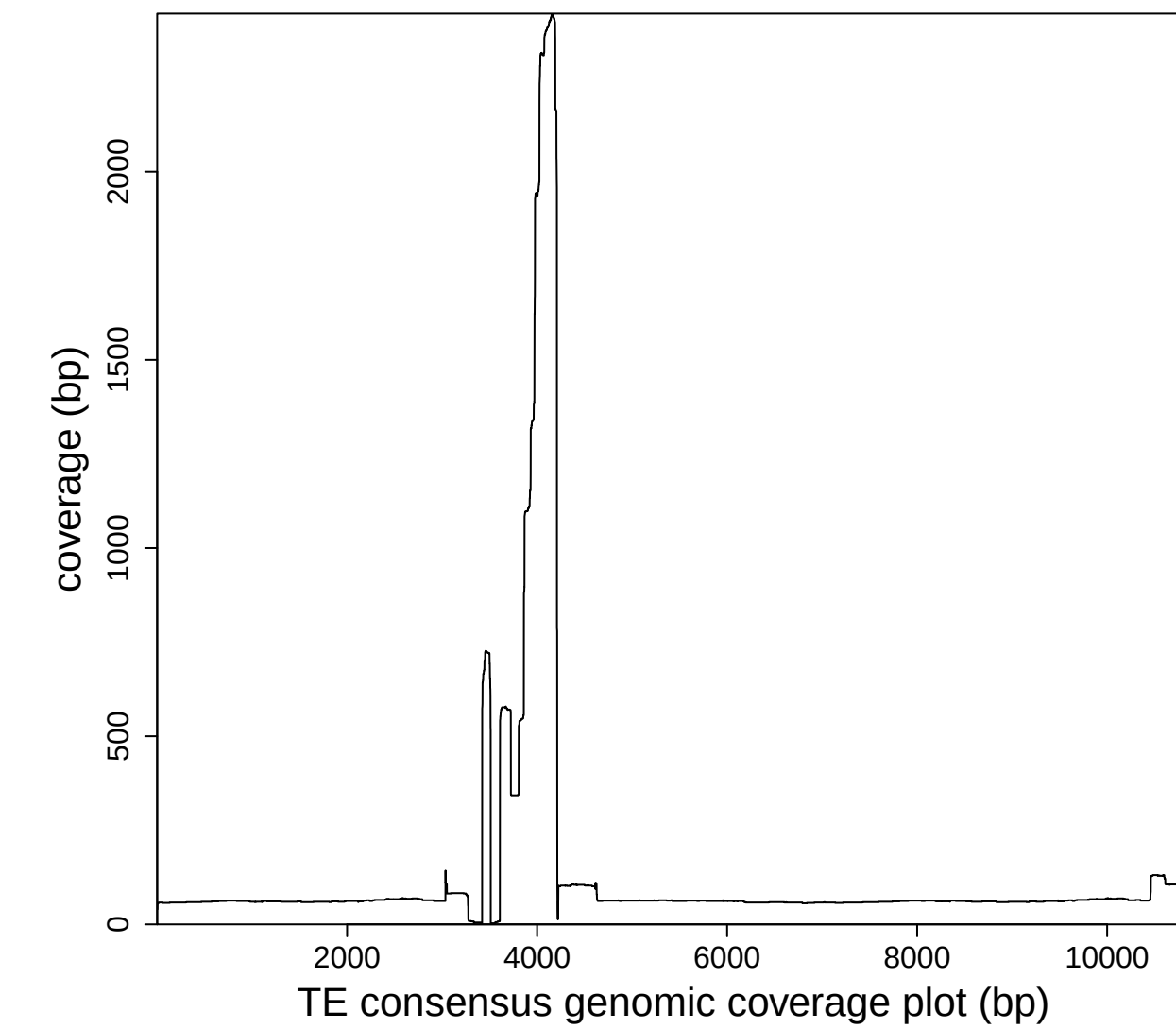
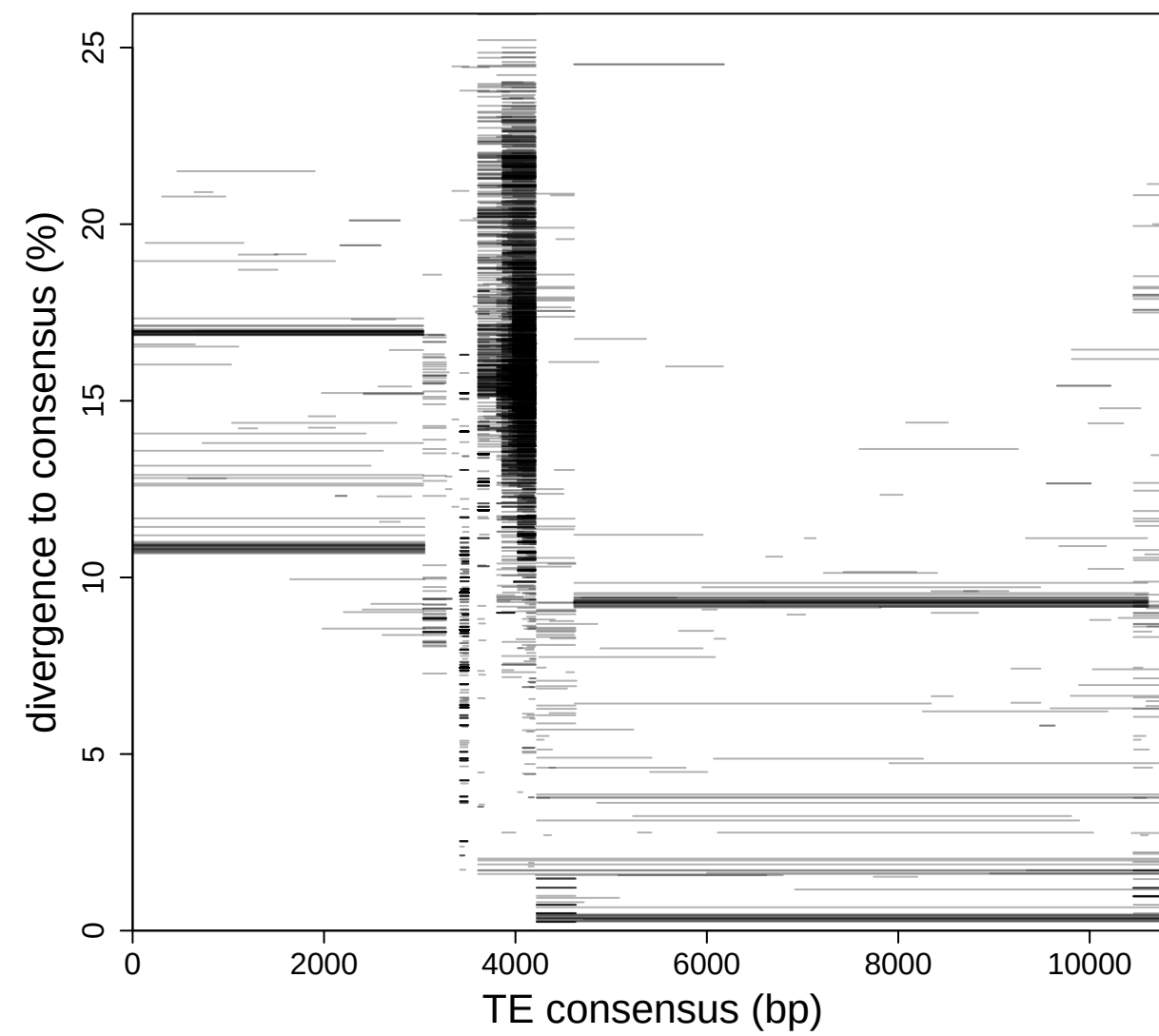






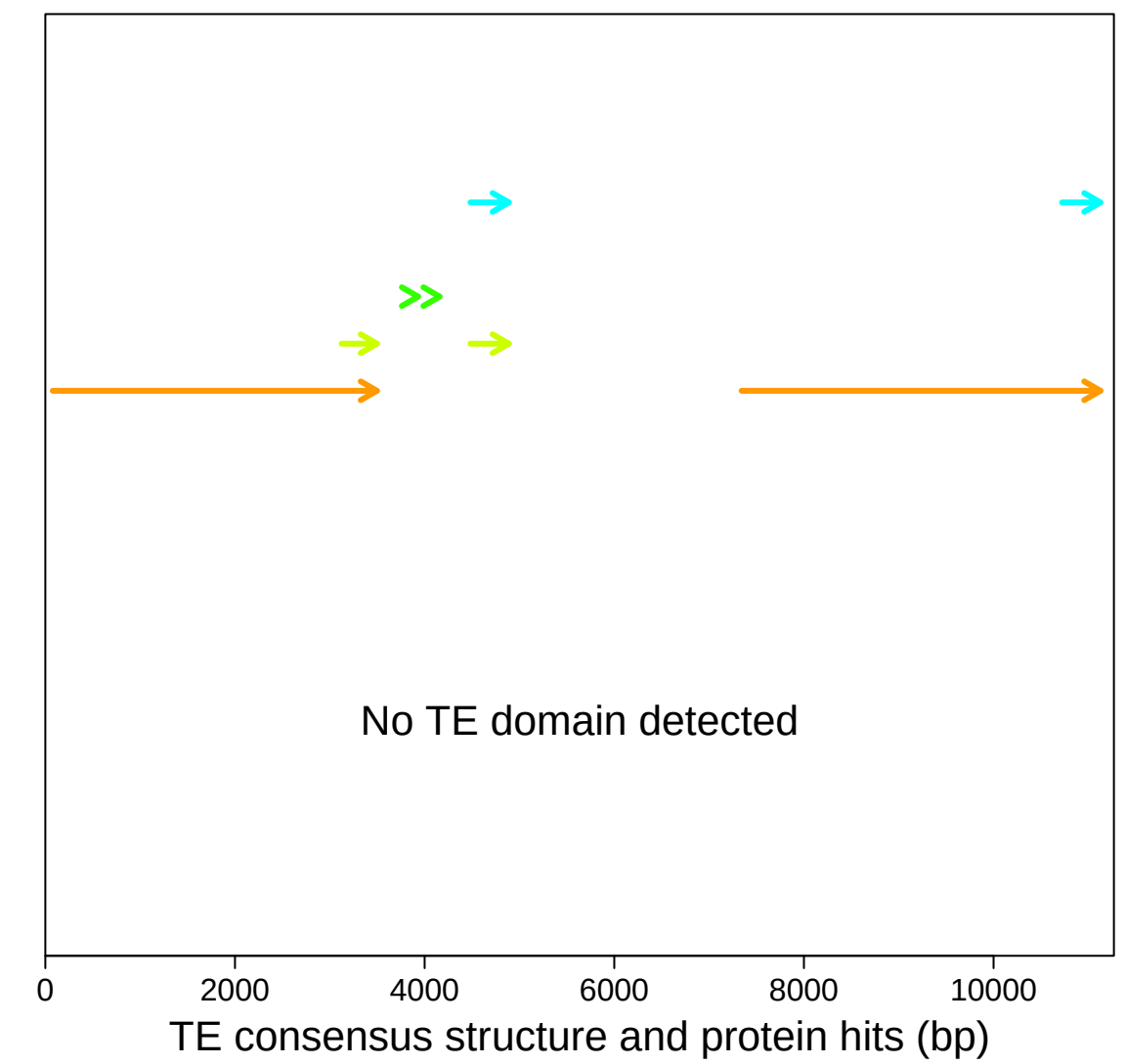
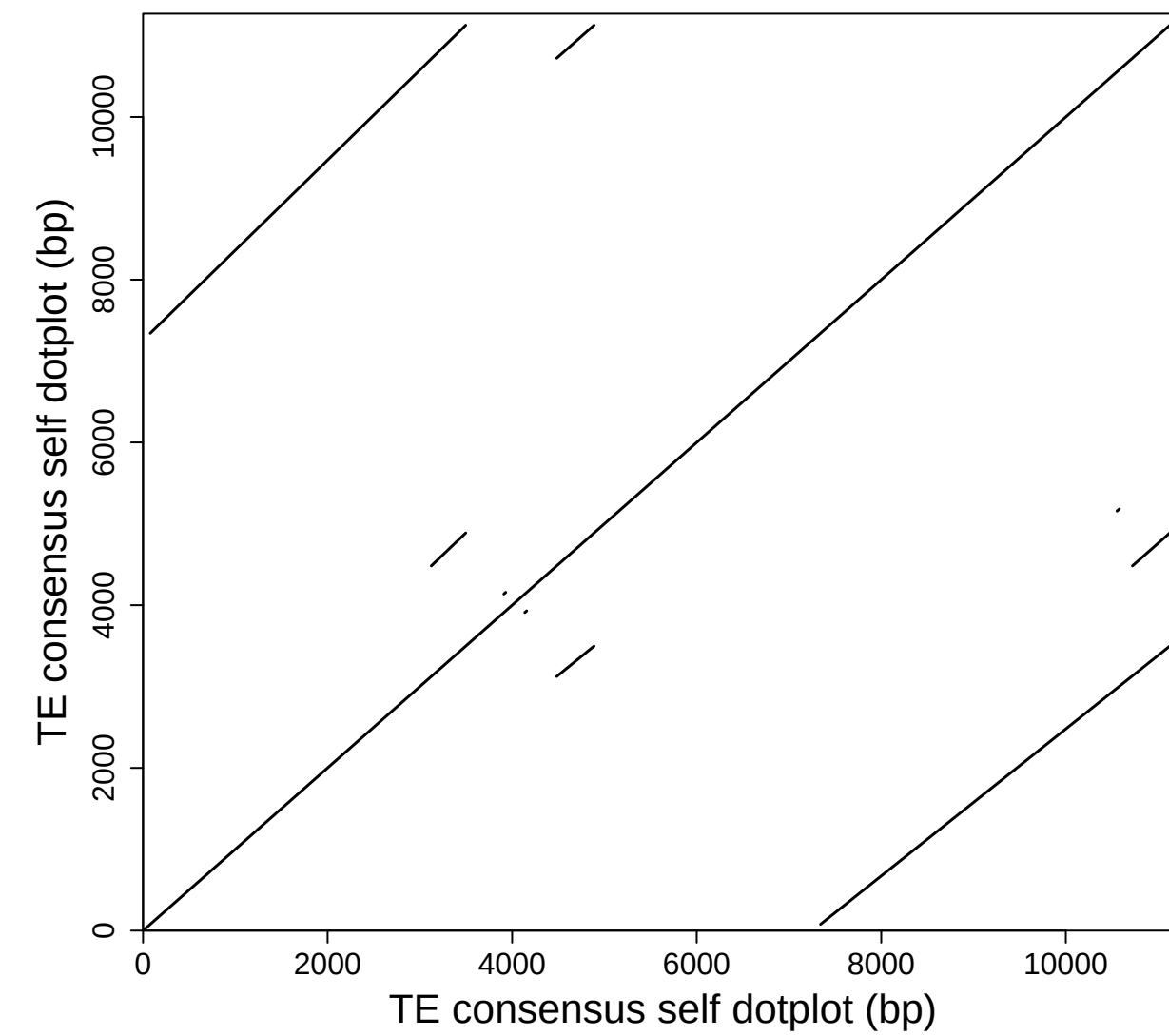
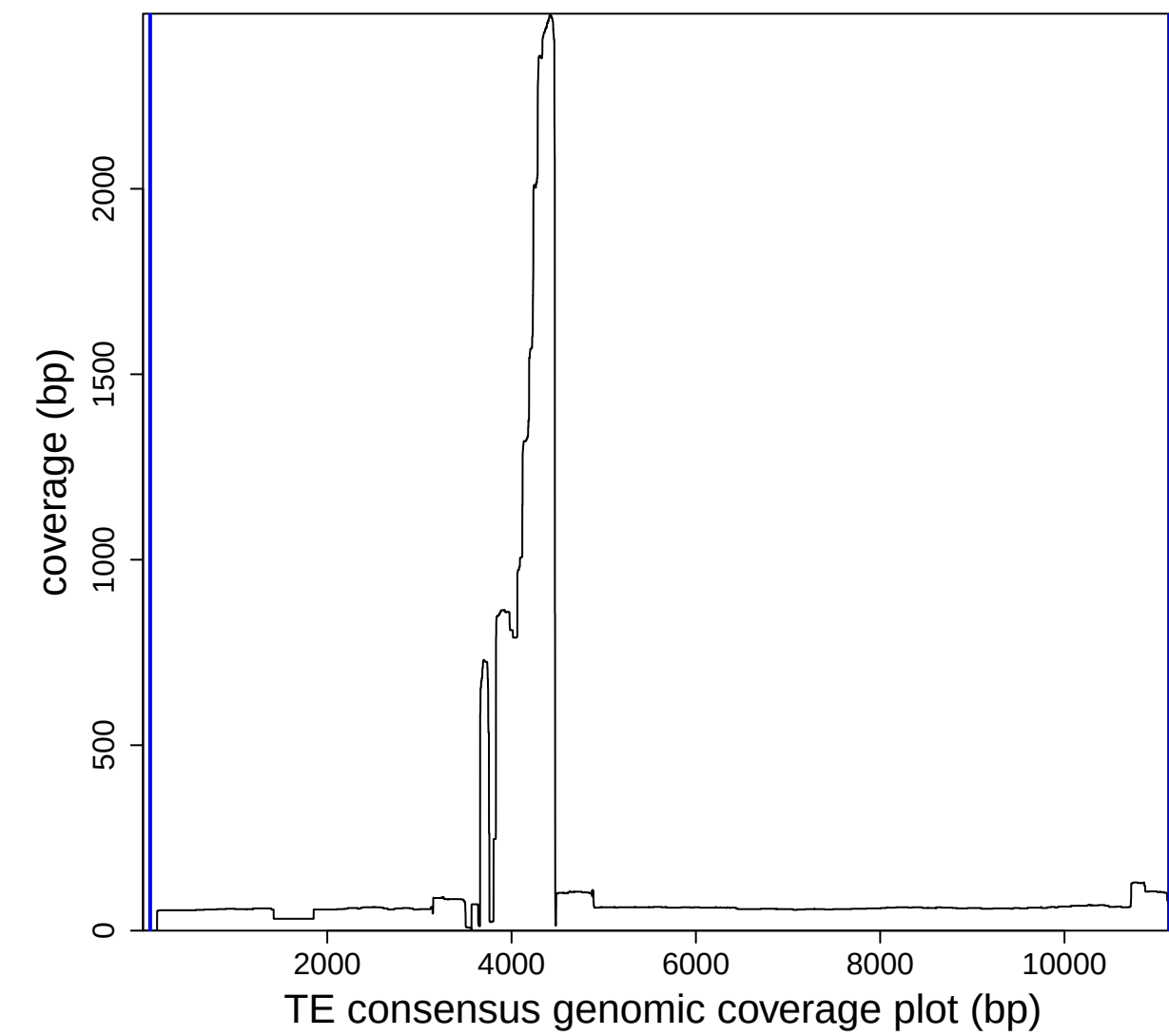
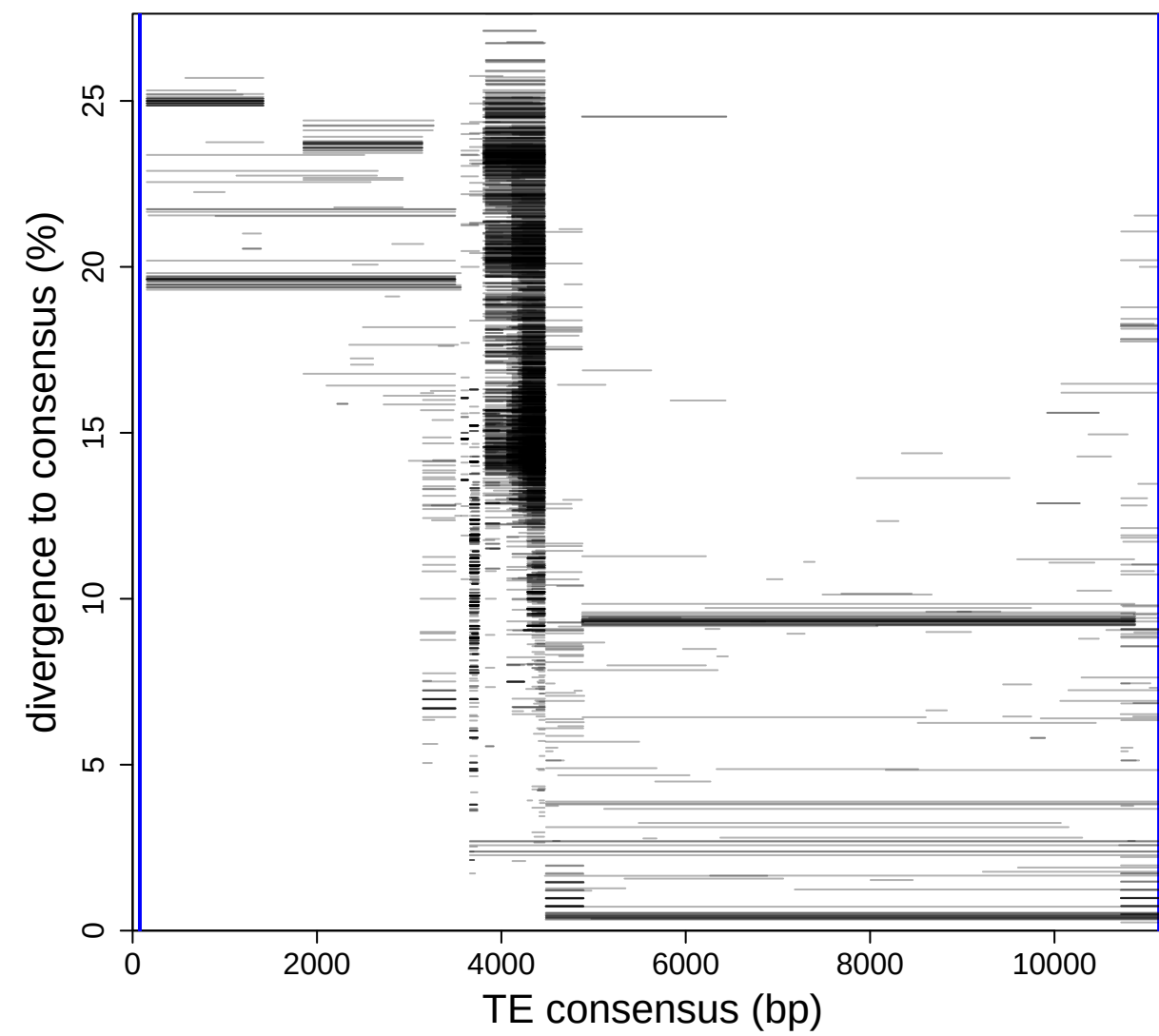
ed_uf.bed g_1.bed fm_1.bed 0 0 bcln.fa aln.fa cl.fa gs.fa ce.
size: 10870bp; fragments: 3893; full length: 0 (>=9783bp)

After TEtrimmer 10870 bp

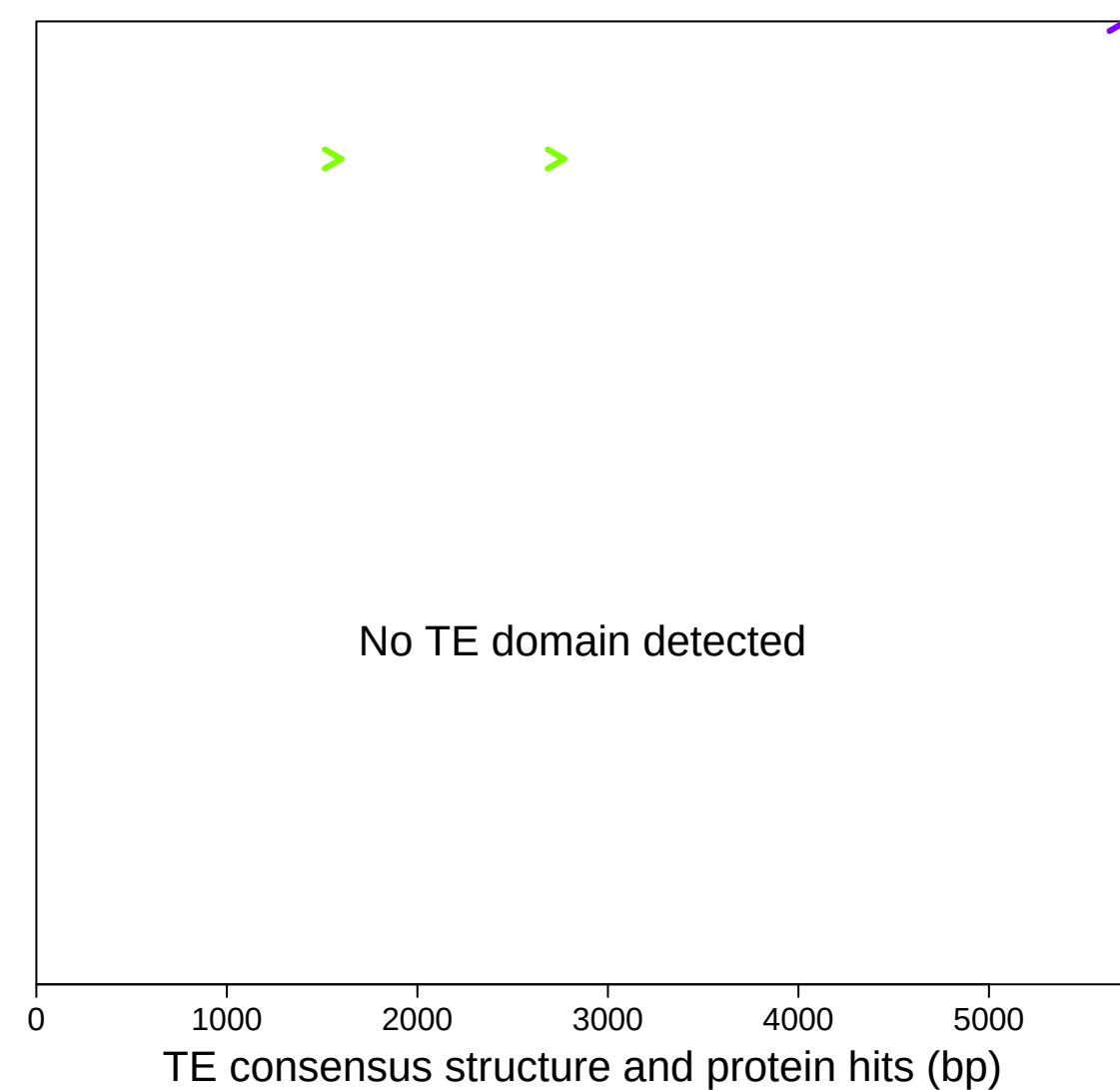
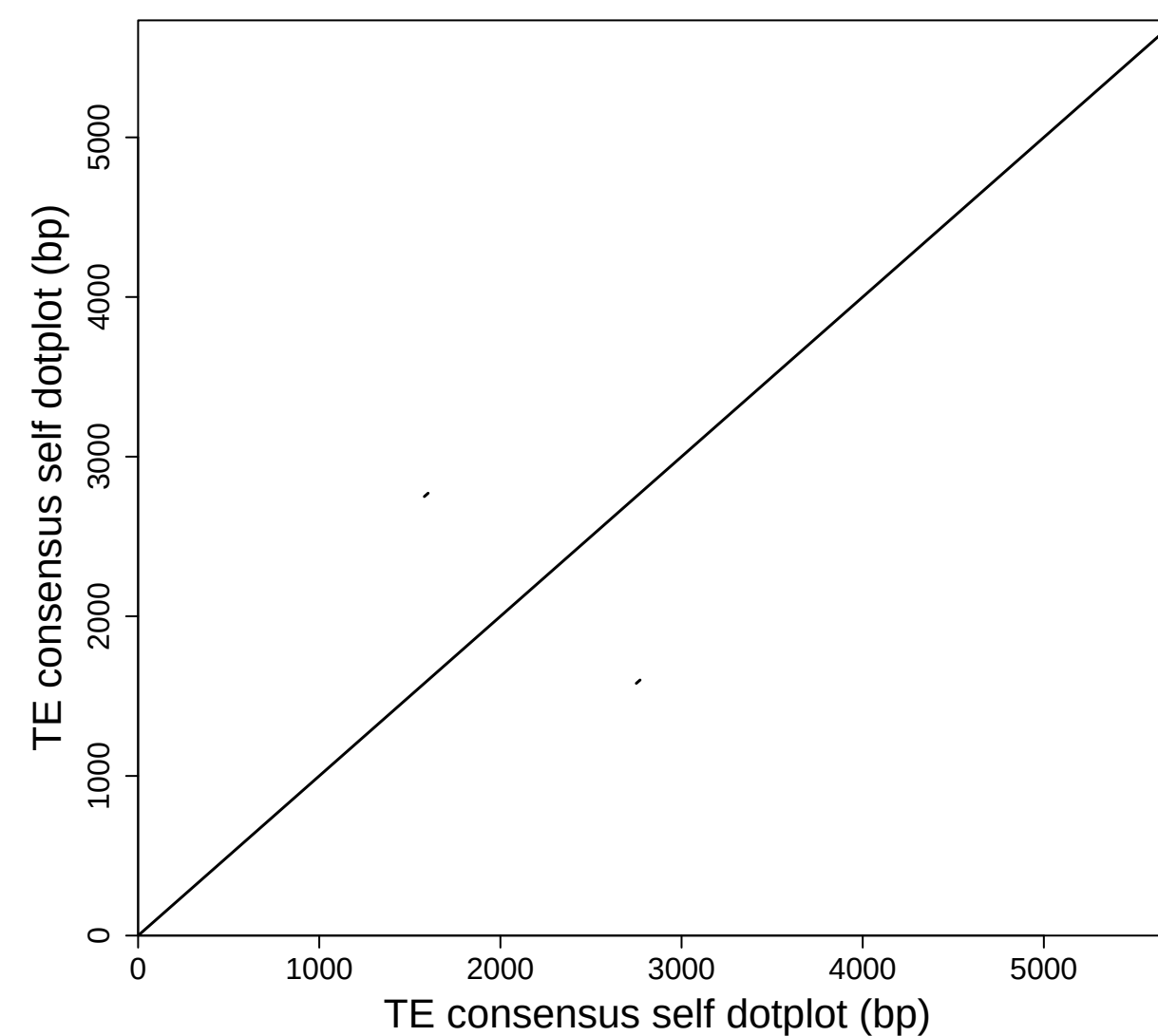
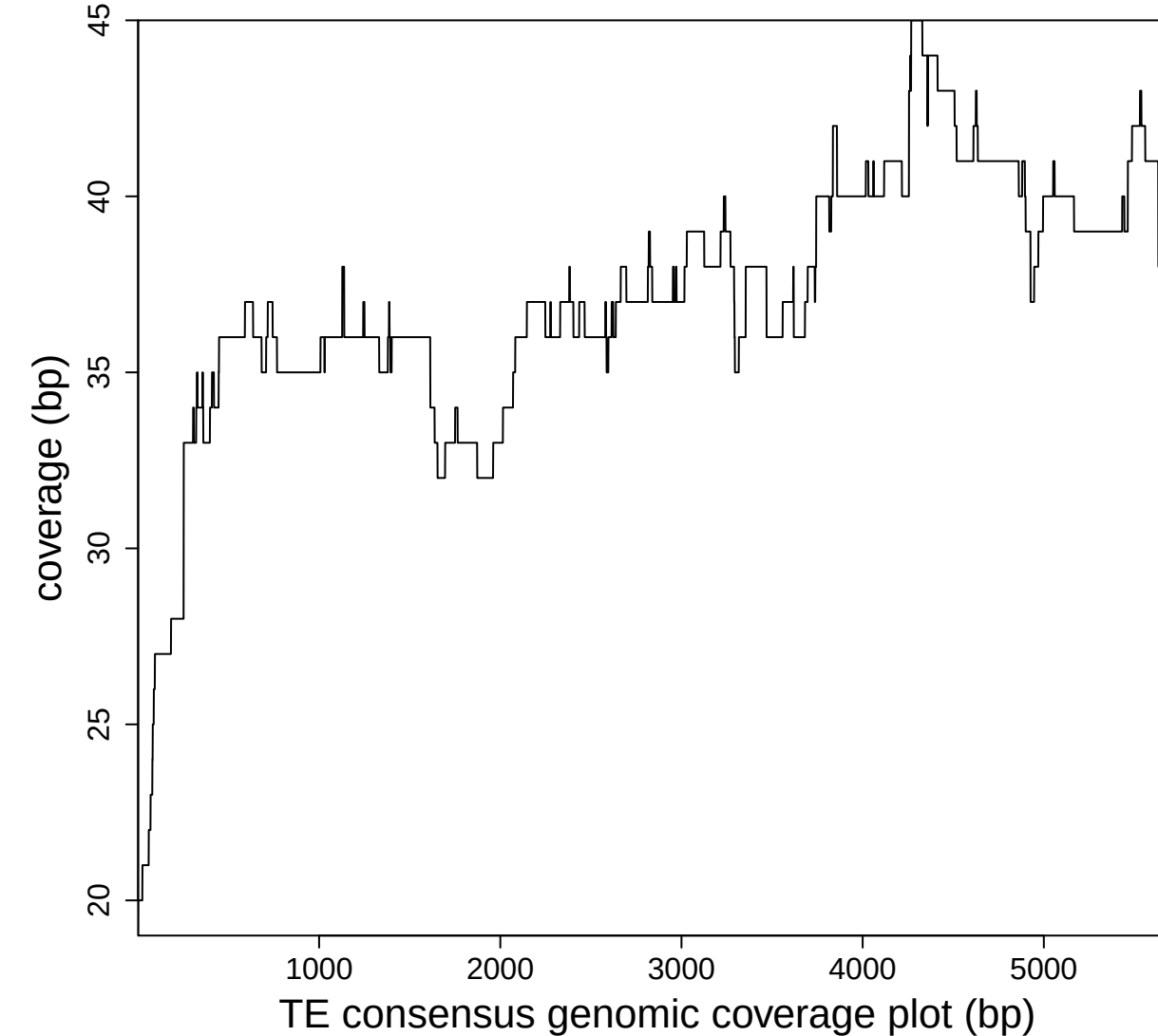
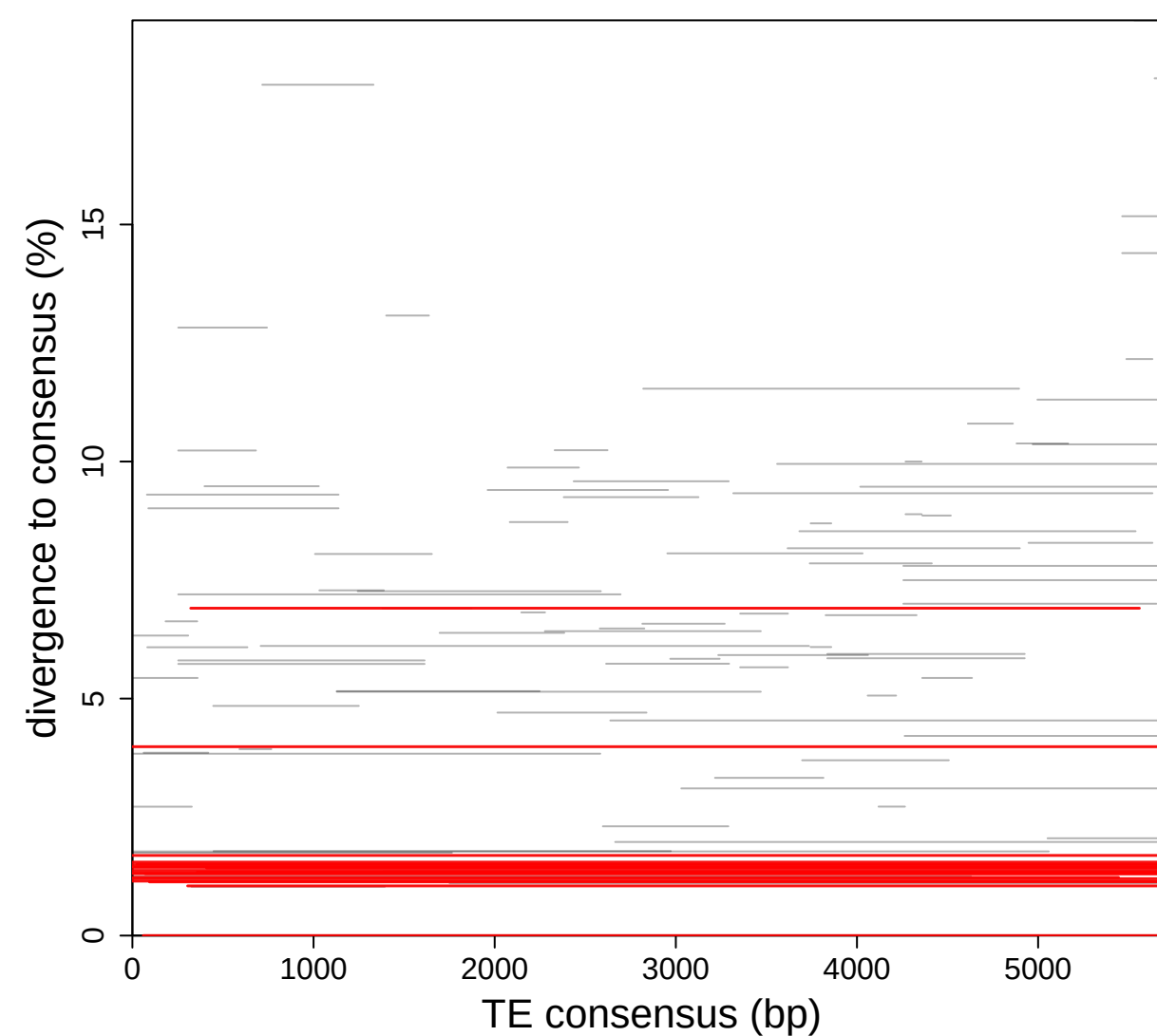


ly_577.fasta.b.bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa
size: 11285bp; fragments: 3806; full length: 0 (>=10156.5bp)

After TEtrimmer Extended plot Blue lines are boundaries

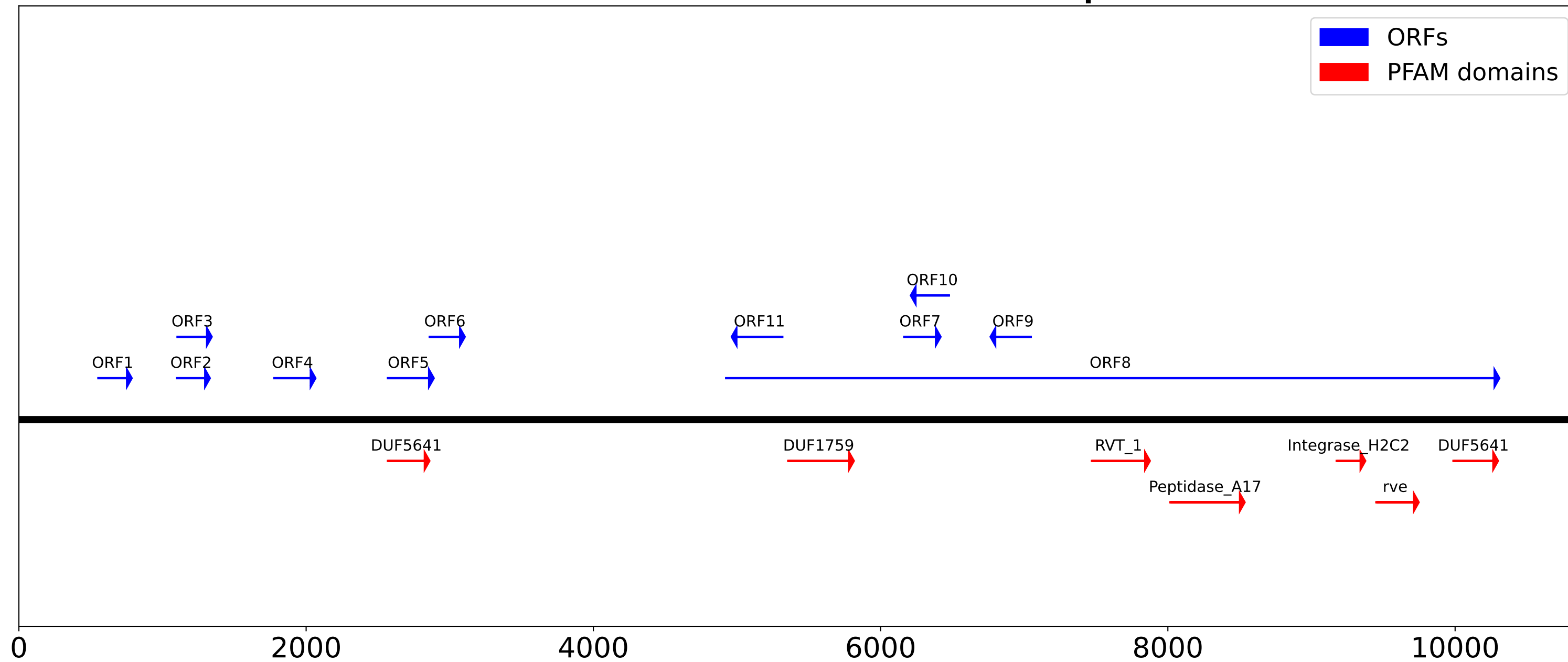


TE: ltr_1_family_577
size: 5733bp; fragments: 114; full length: 20 (≥ 5159.7 bp)

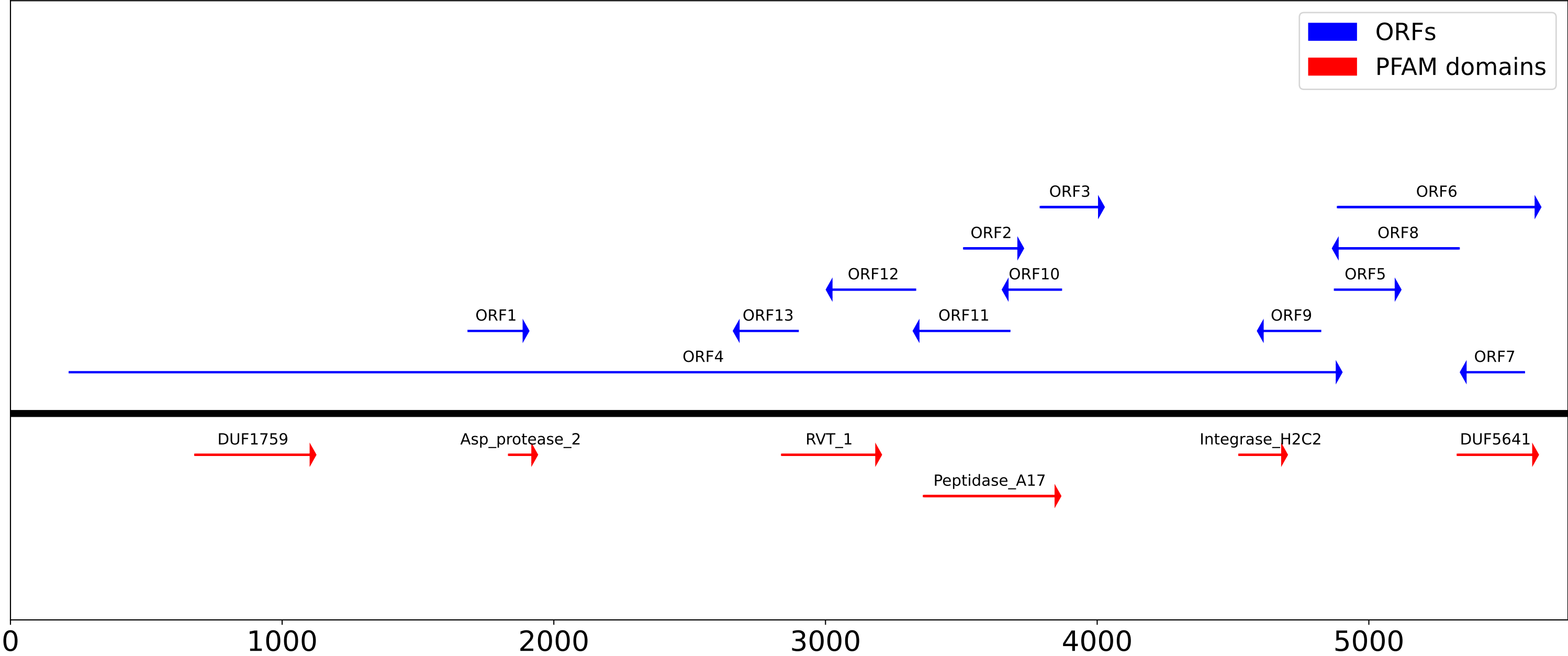


Before TEtrimmer 5733 bp

After TEtrimmer ORF and PFAM domain plot



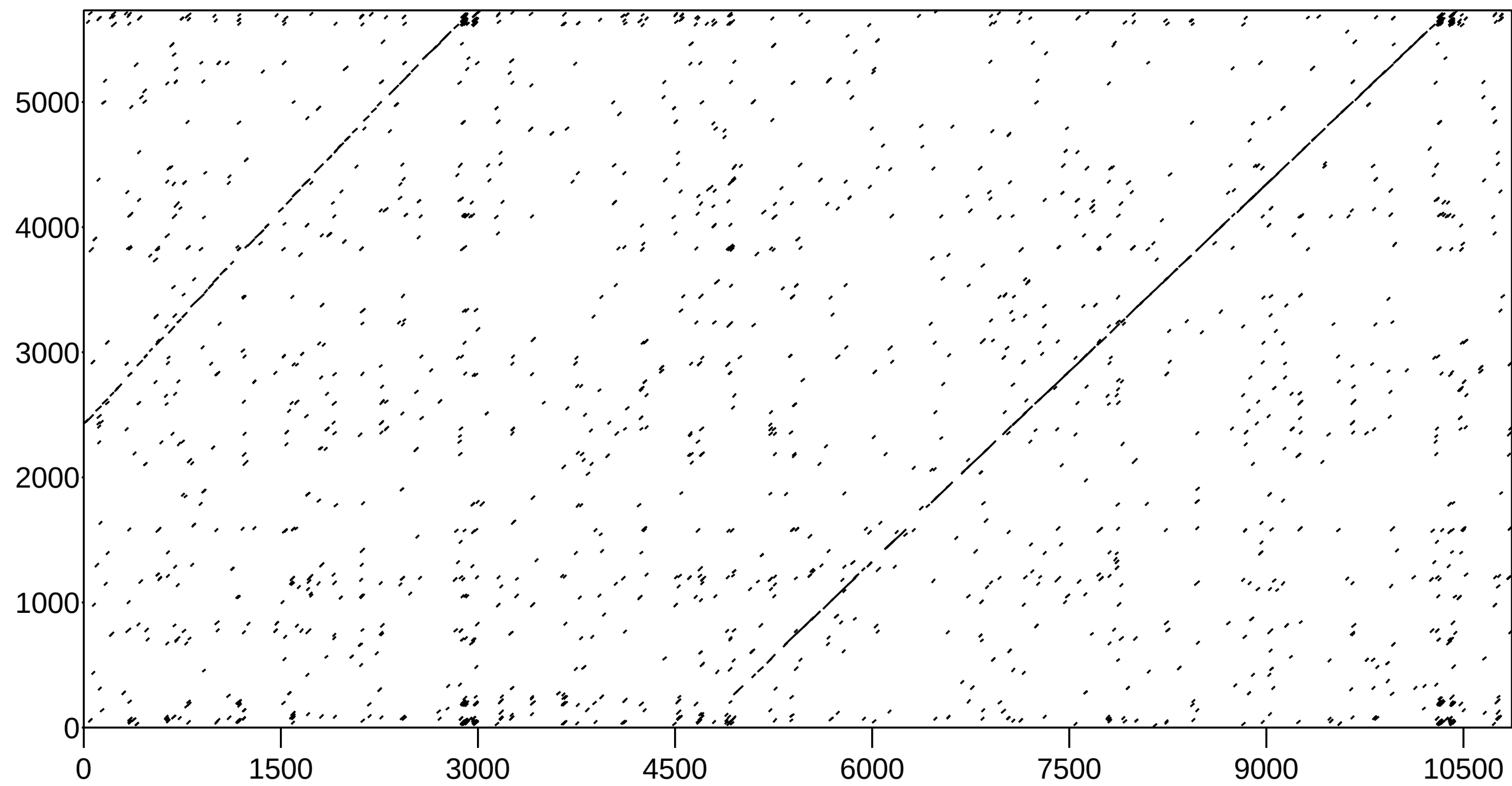
Before TEtrimmer ORF and PFAM domain plot



Dotplot

(windowsize = 25, threshold = 50.00 09/10/25)

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)