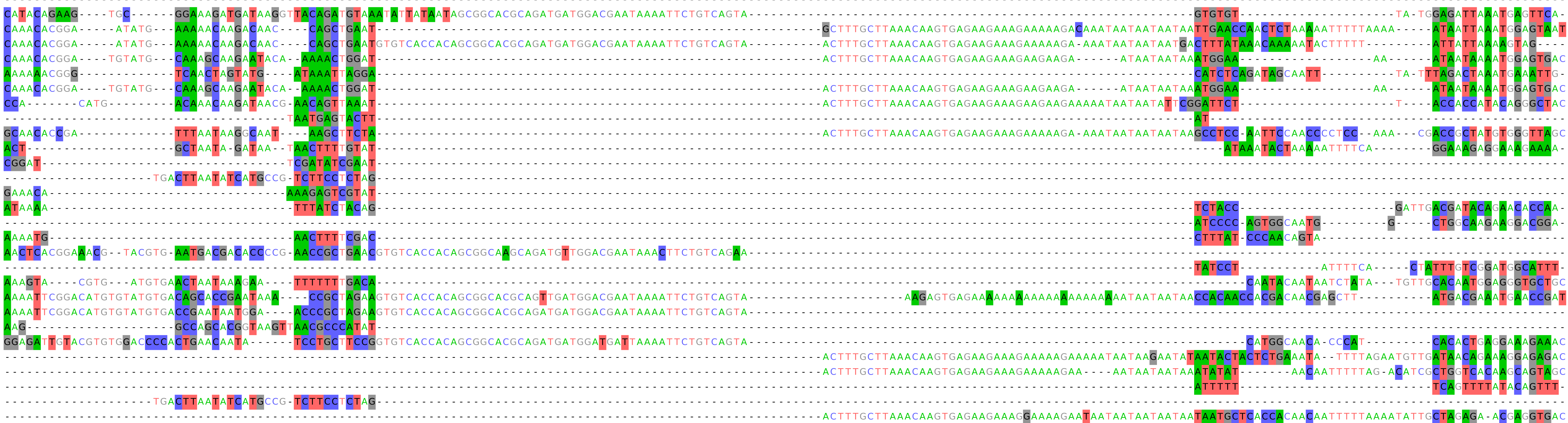
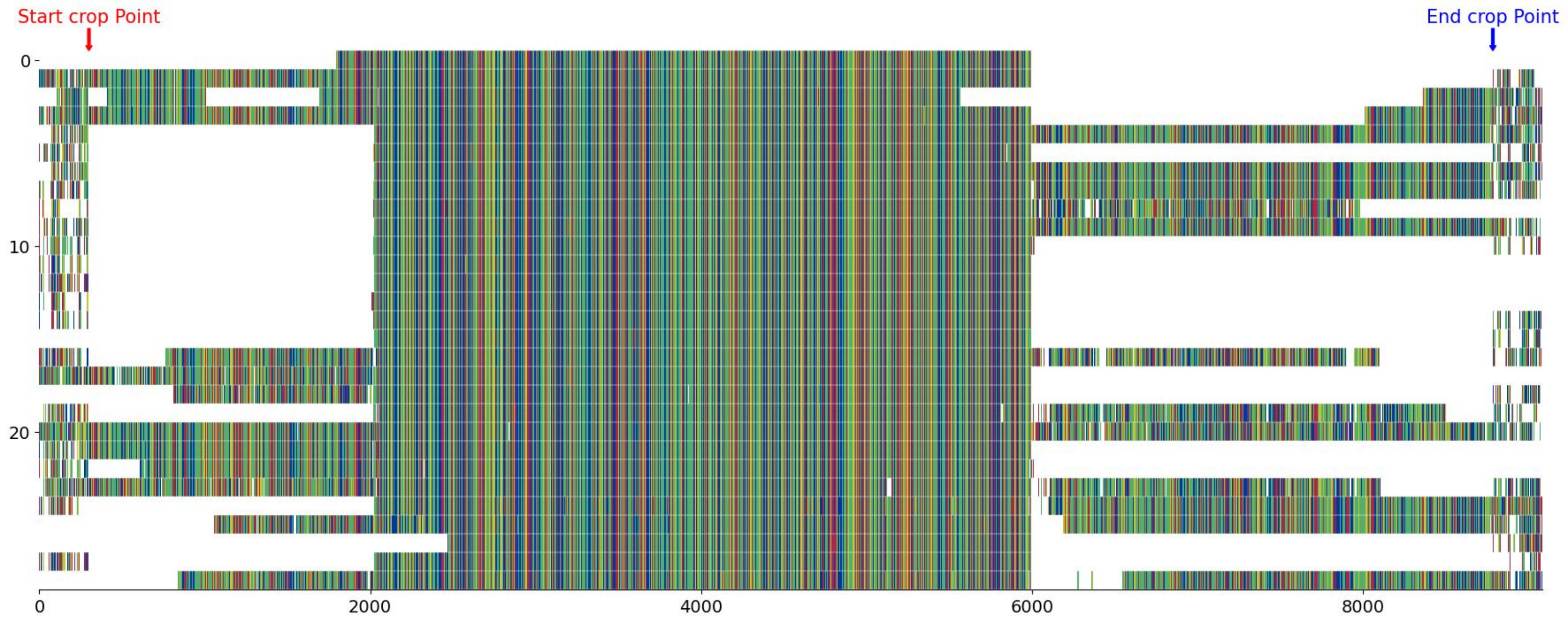


MSA length = 8487

Start crop Point

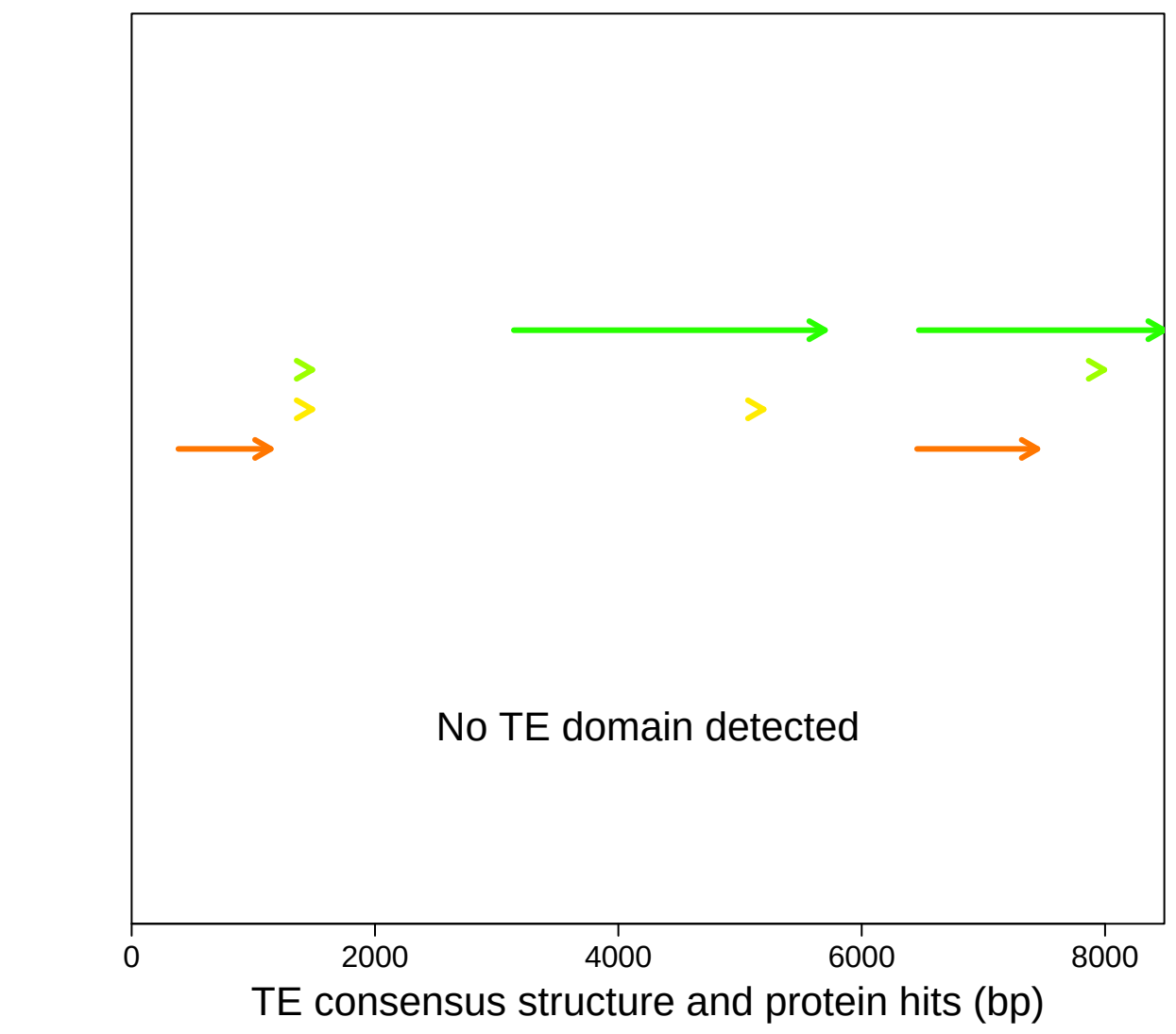
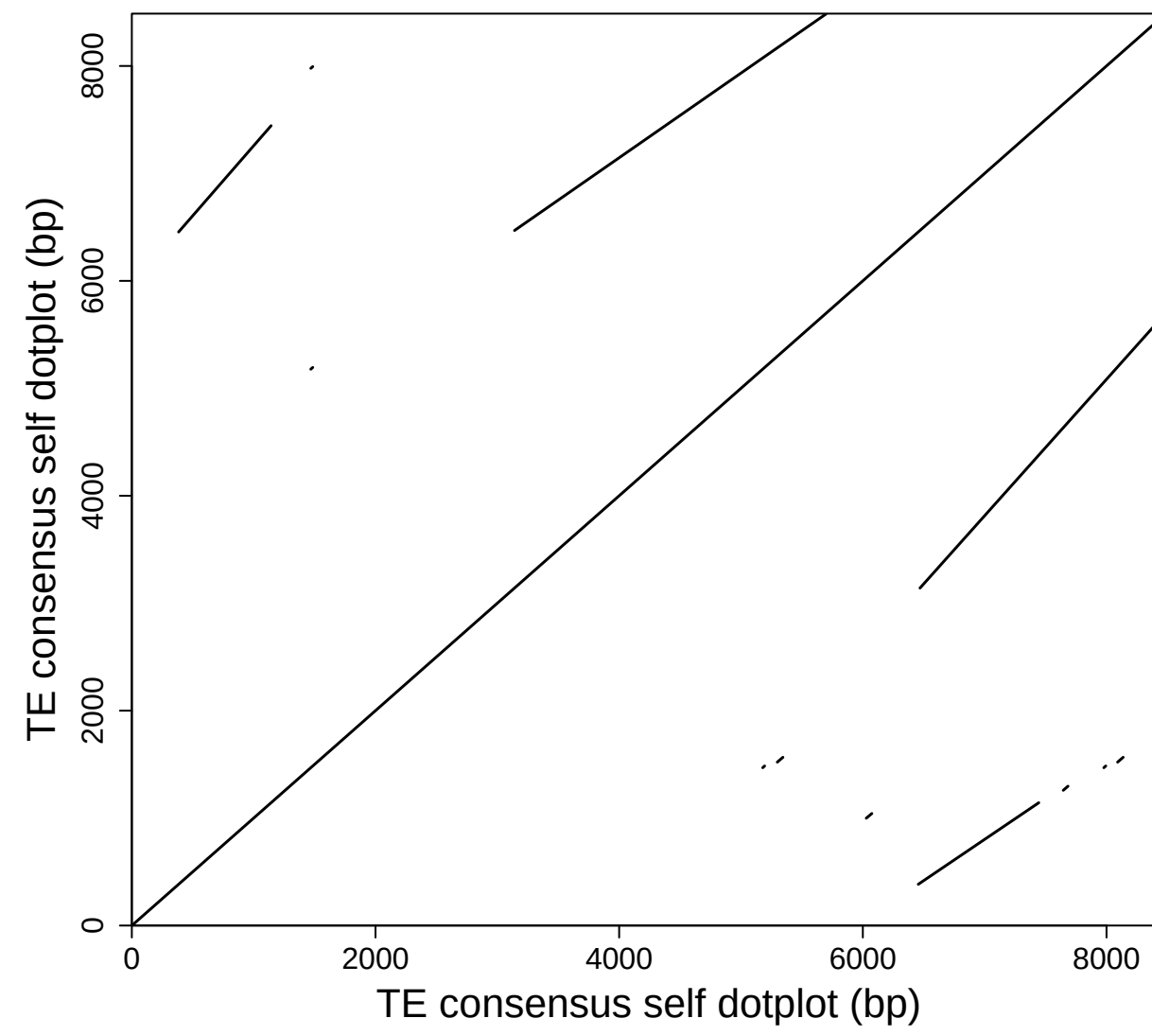
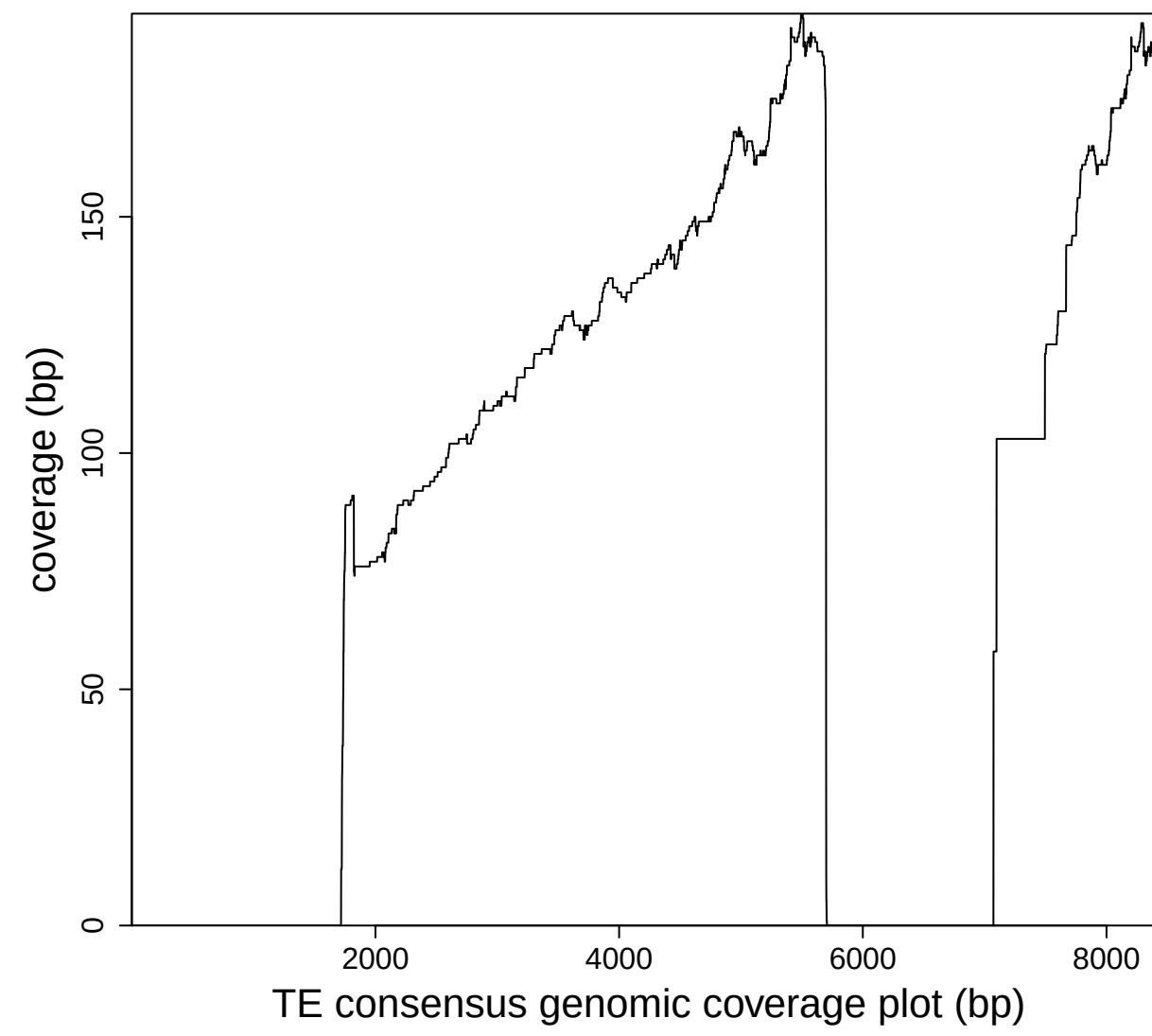
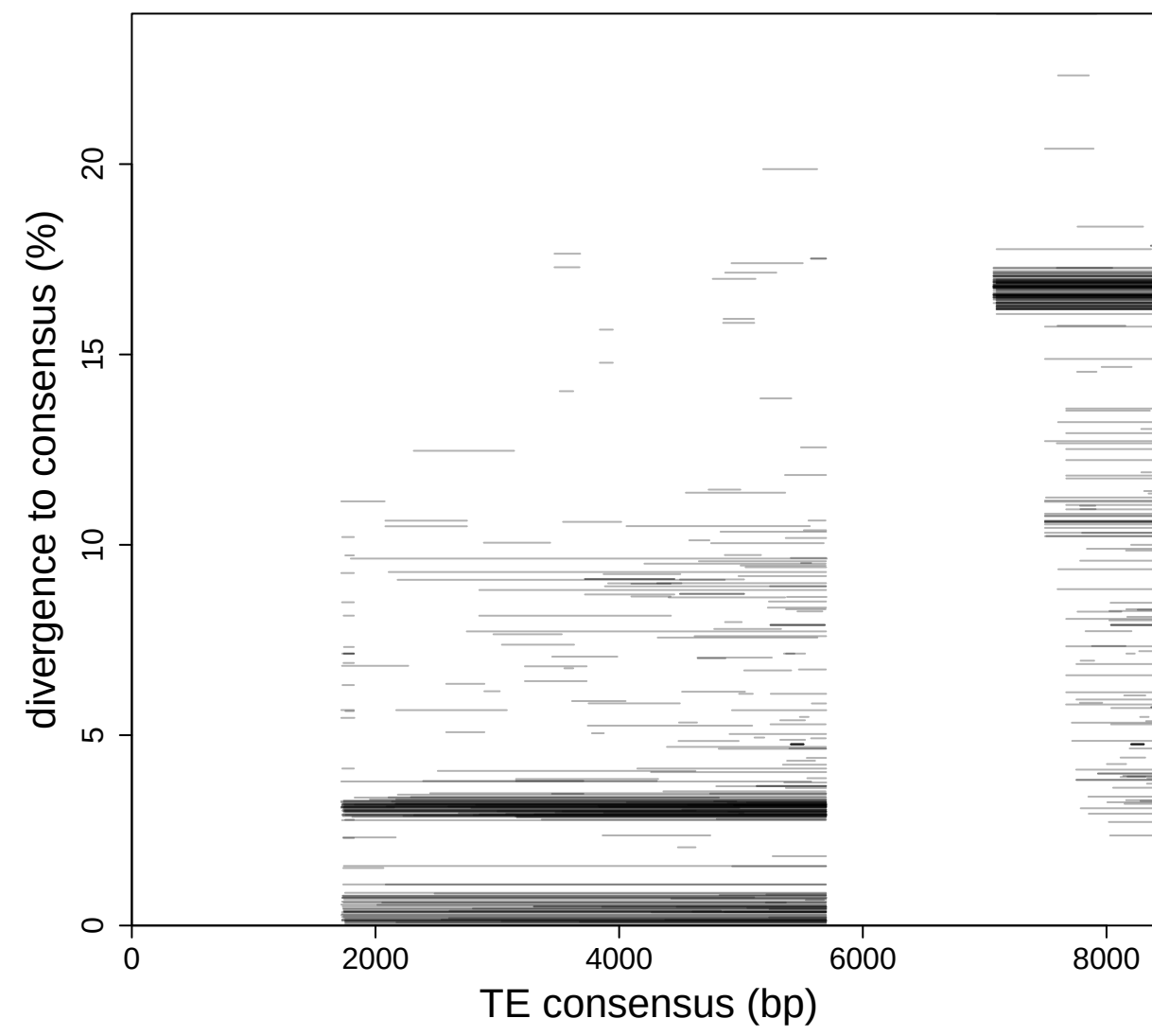
End crop Point





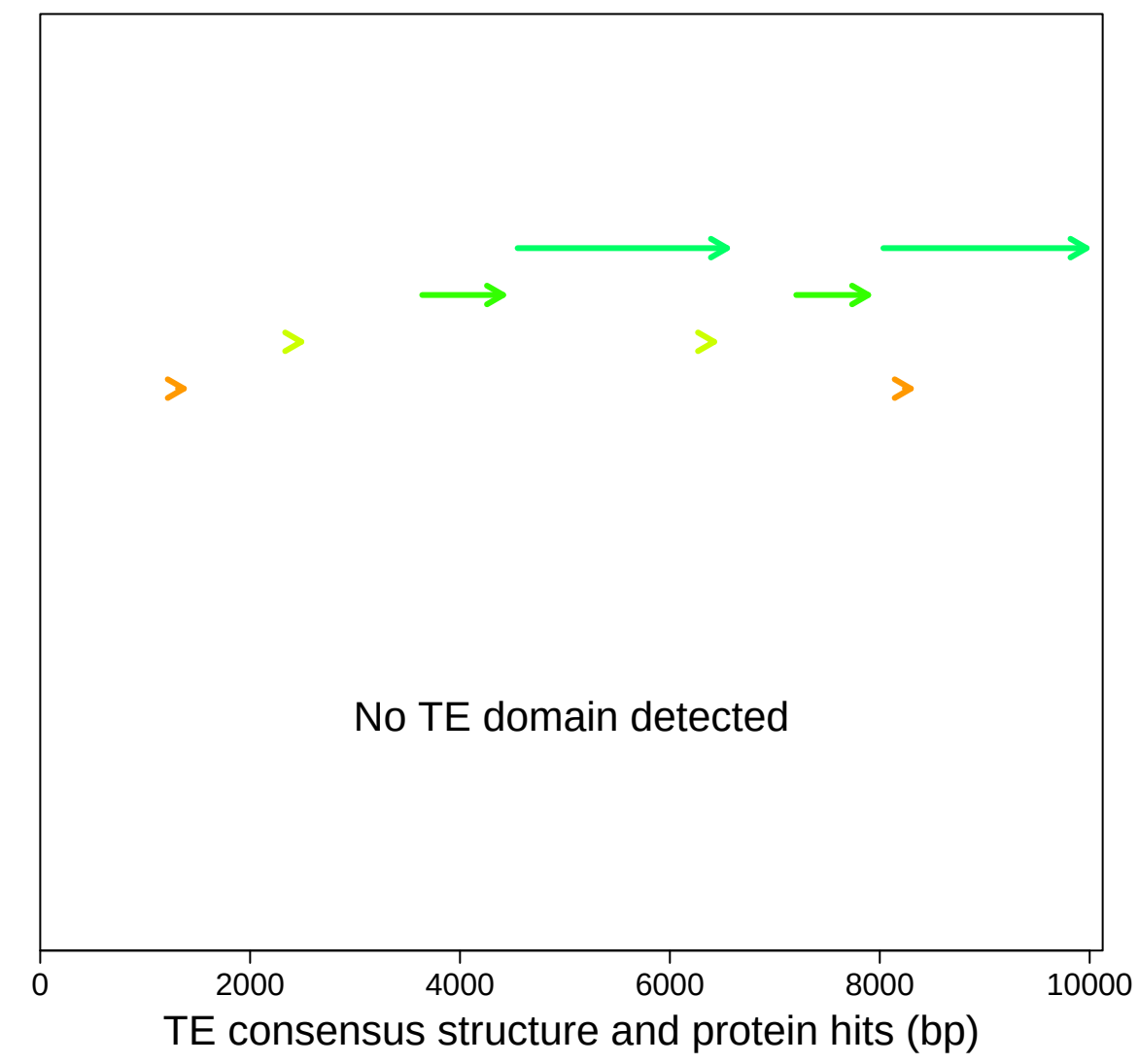
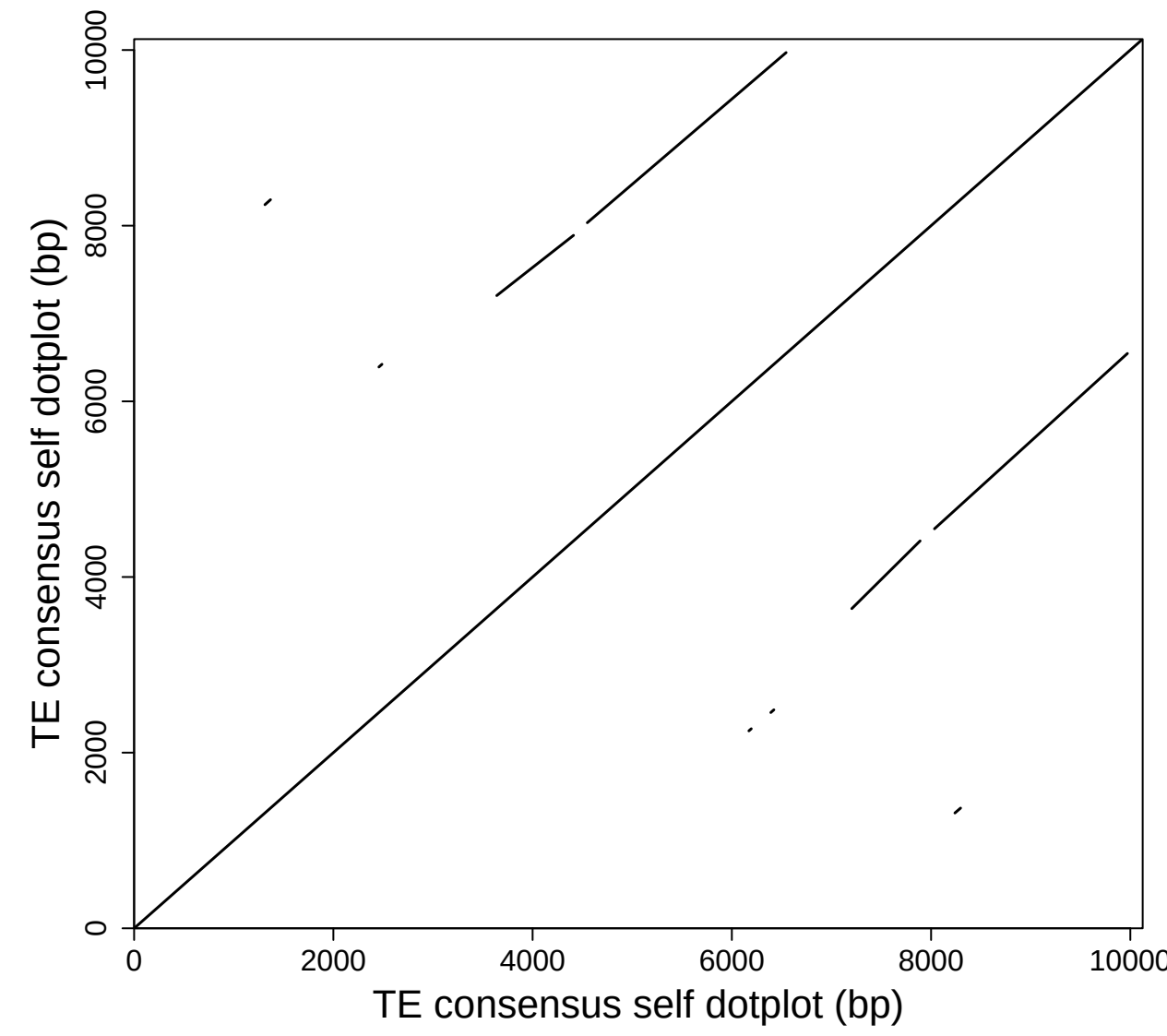
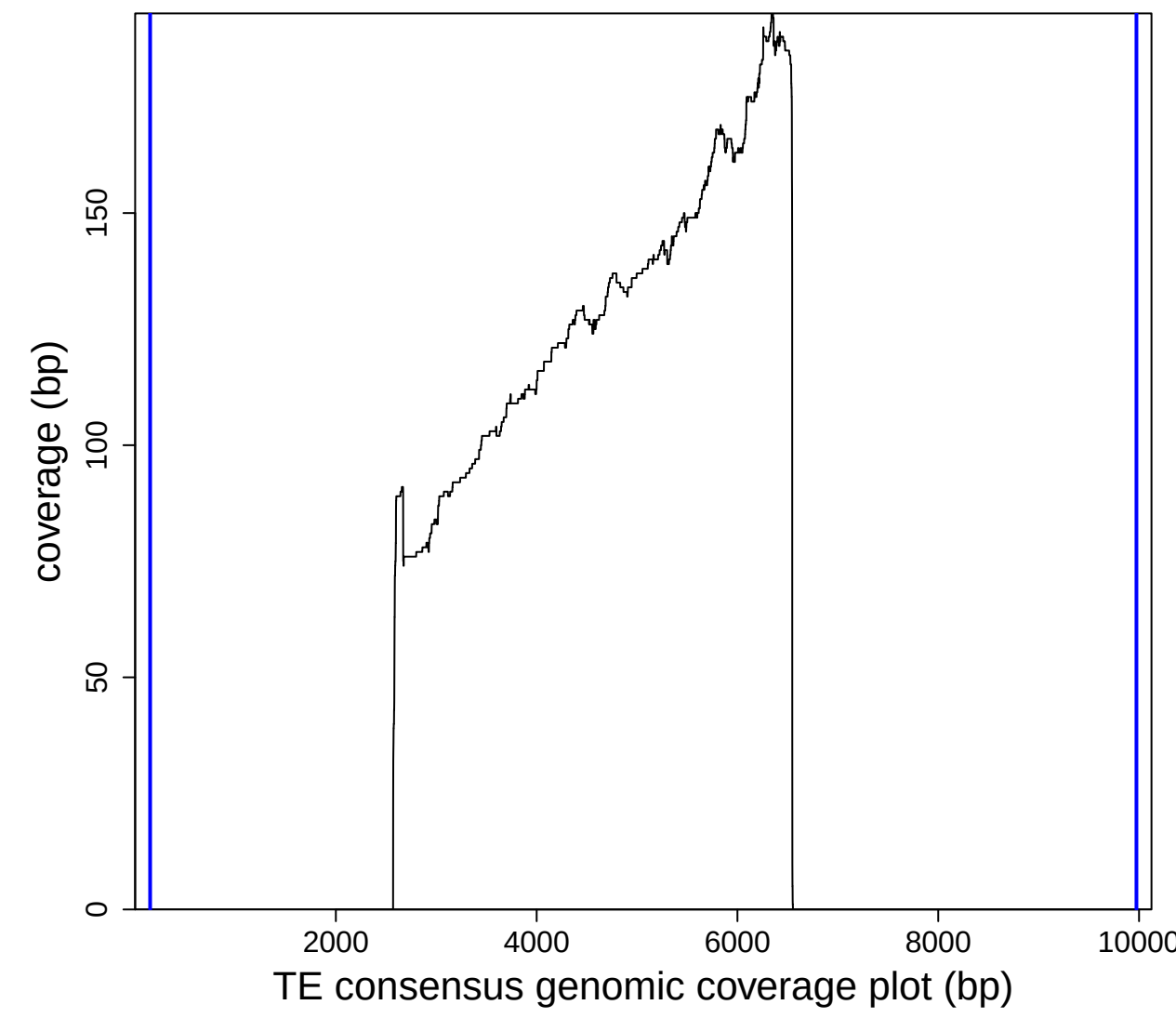
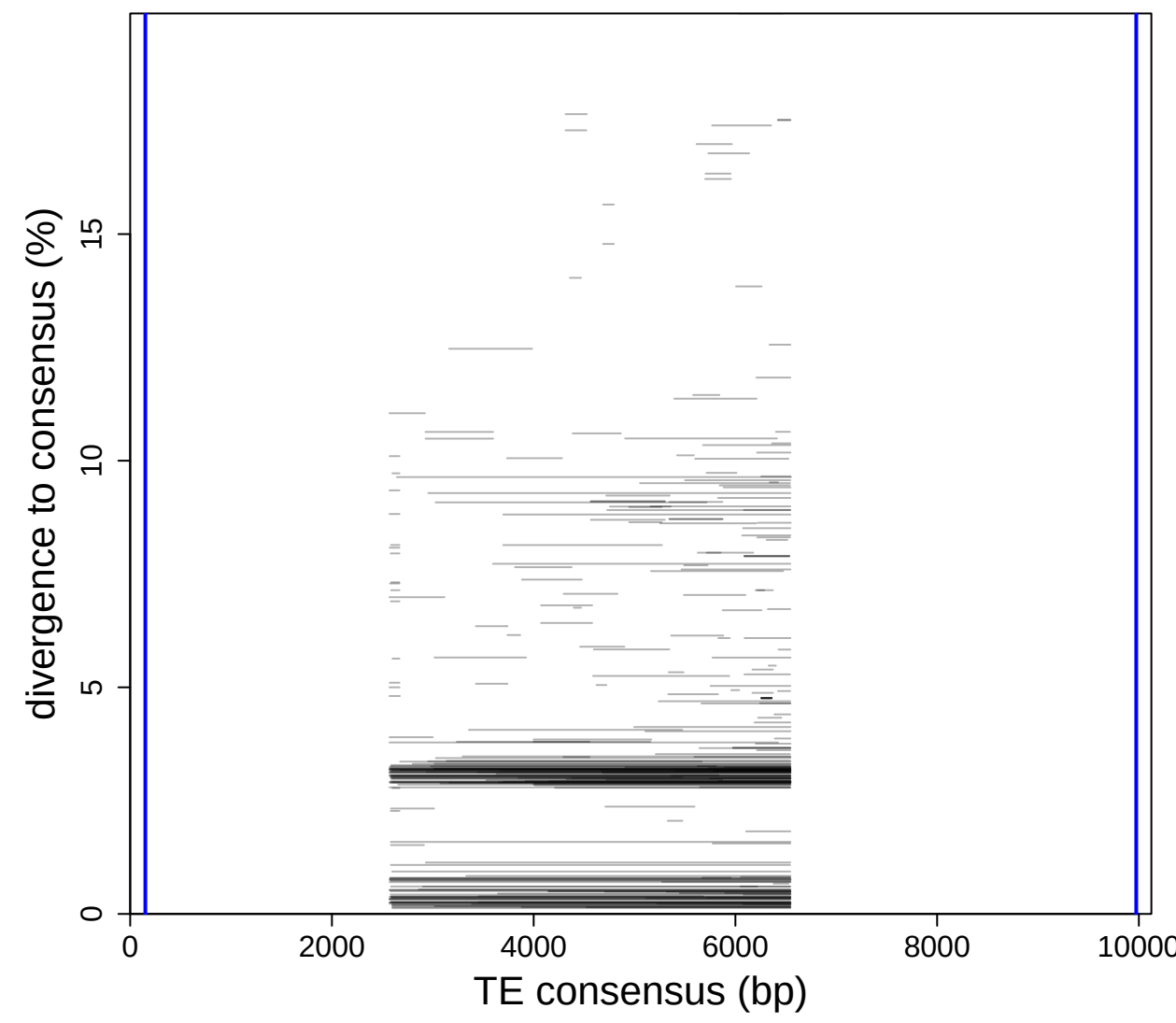
1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcln.fa_aln.
size: 8487bp; fragments: 513; full length: 0 (>=7638.3bp)

After TEtrimmer 8487 bp



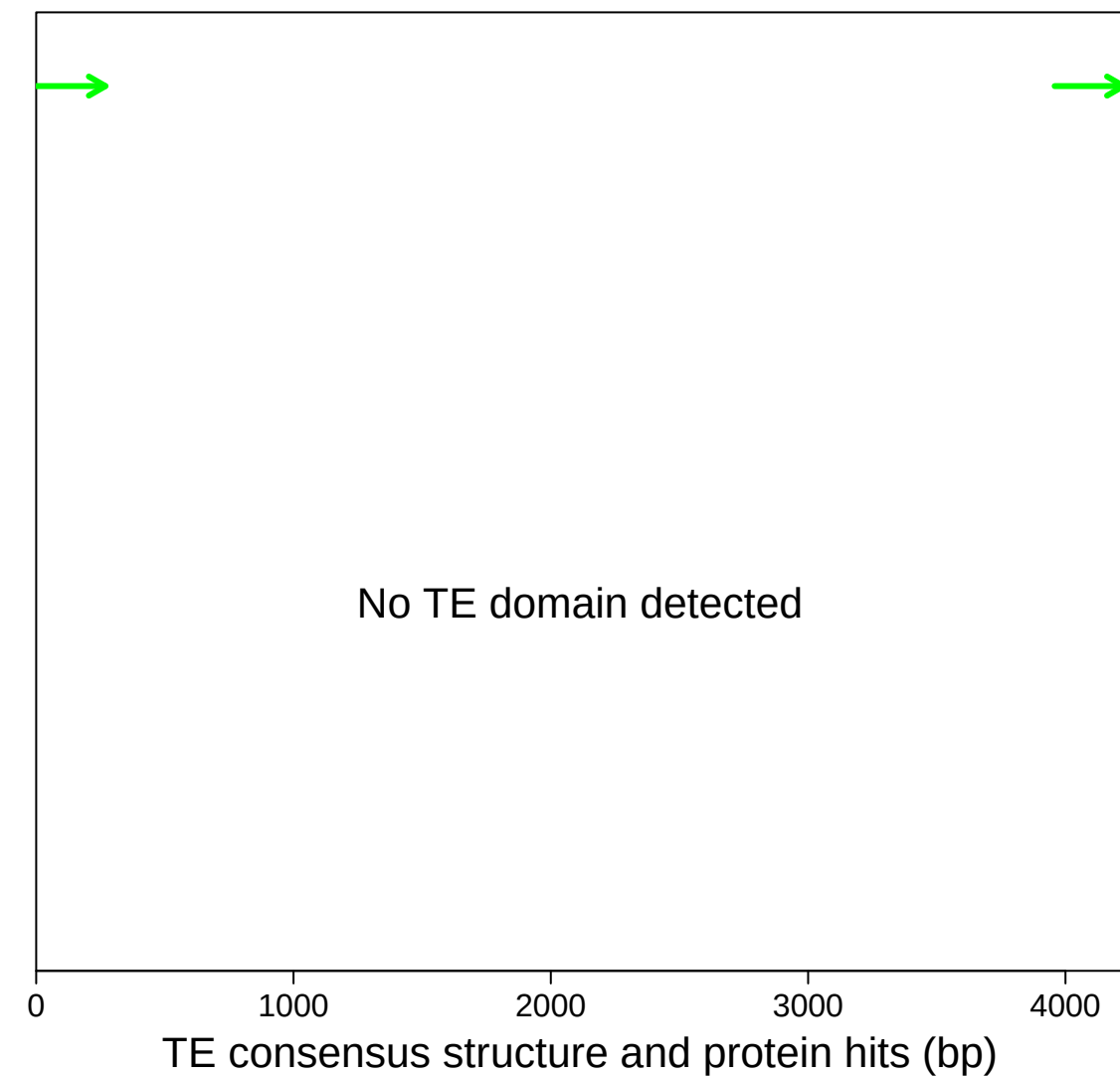
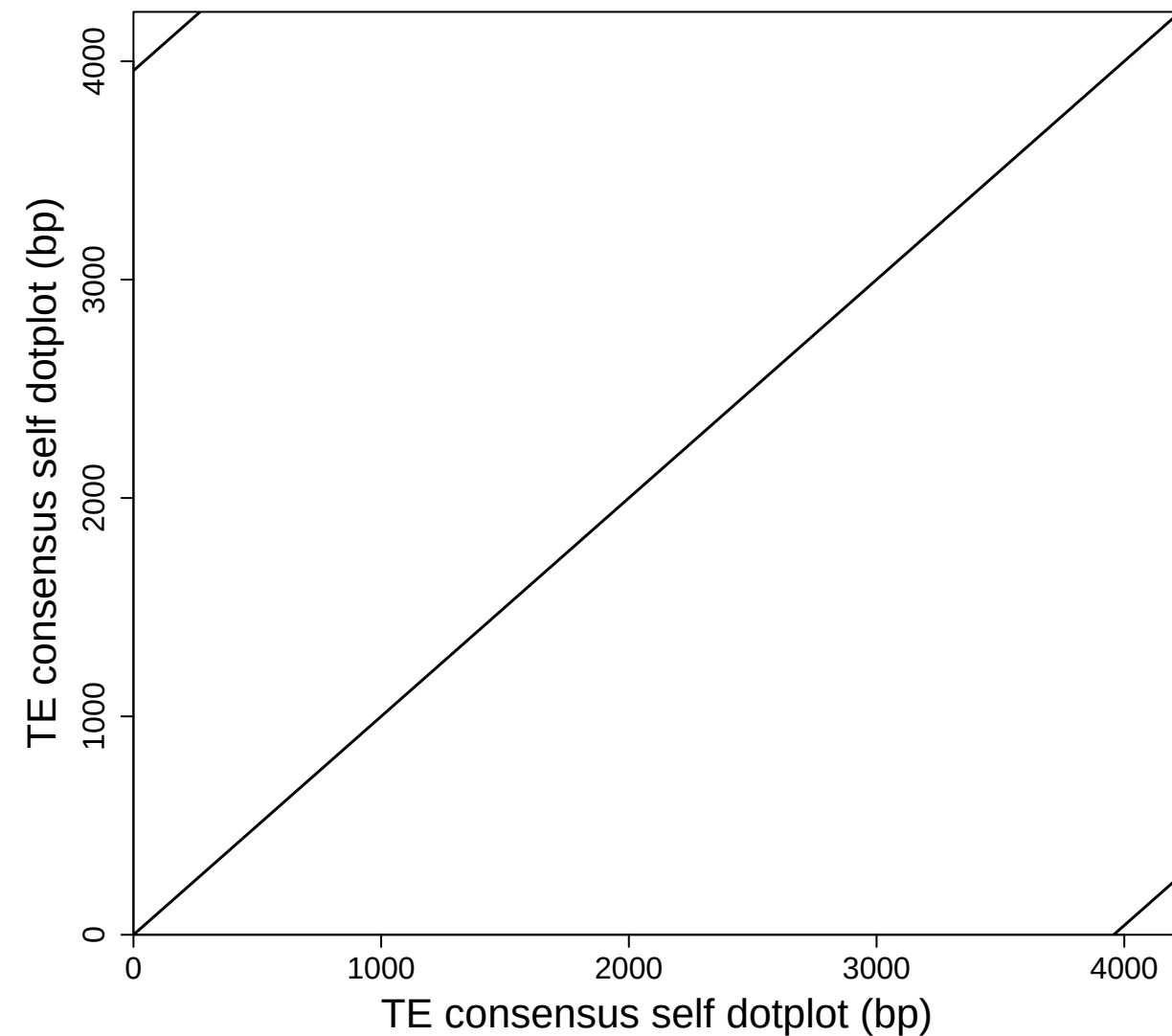
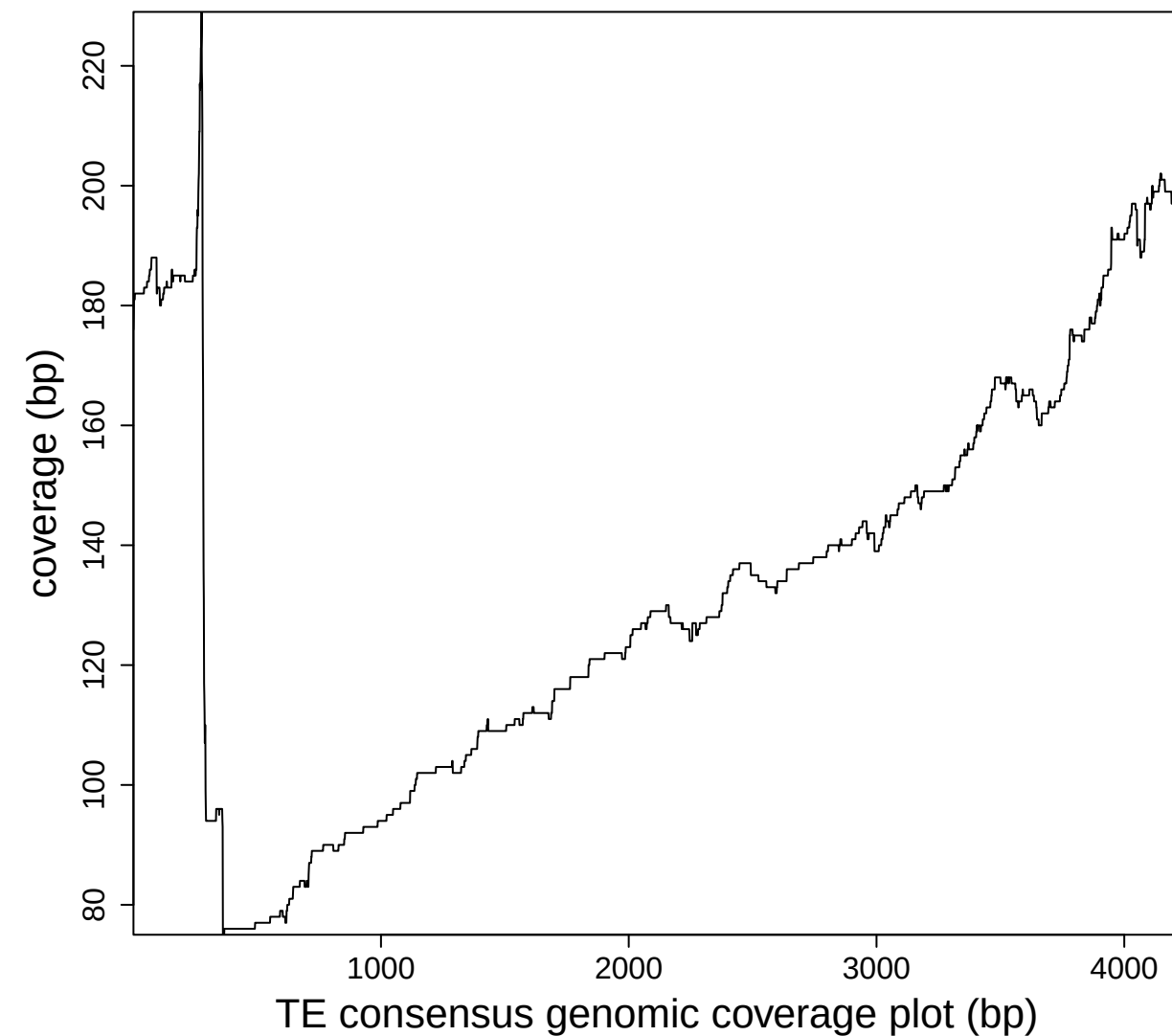
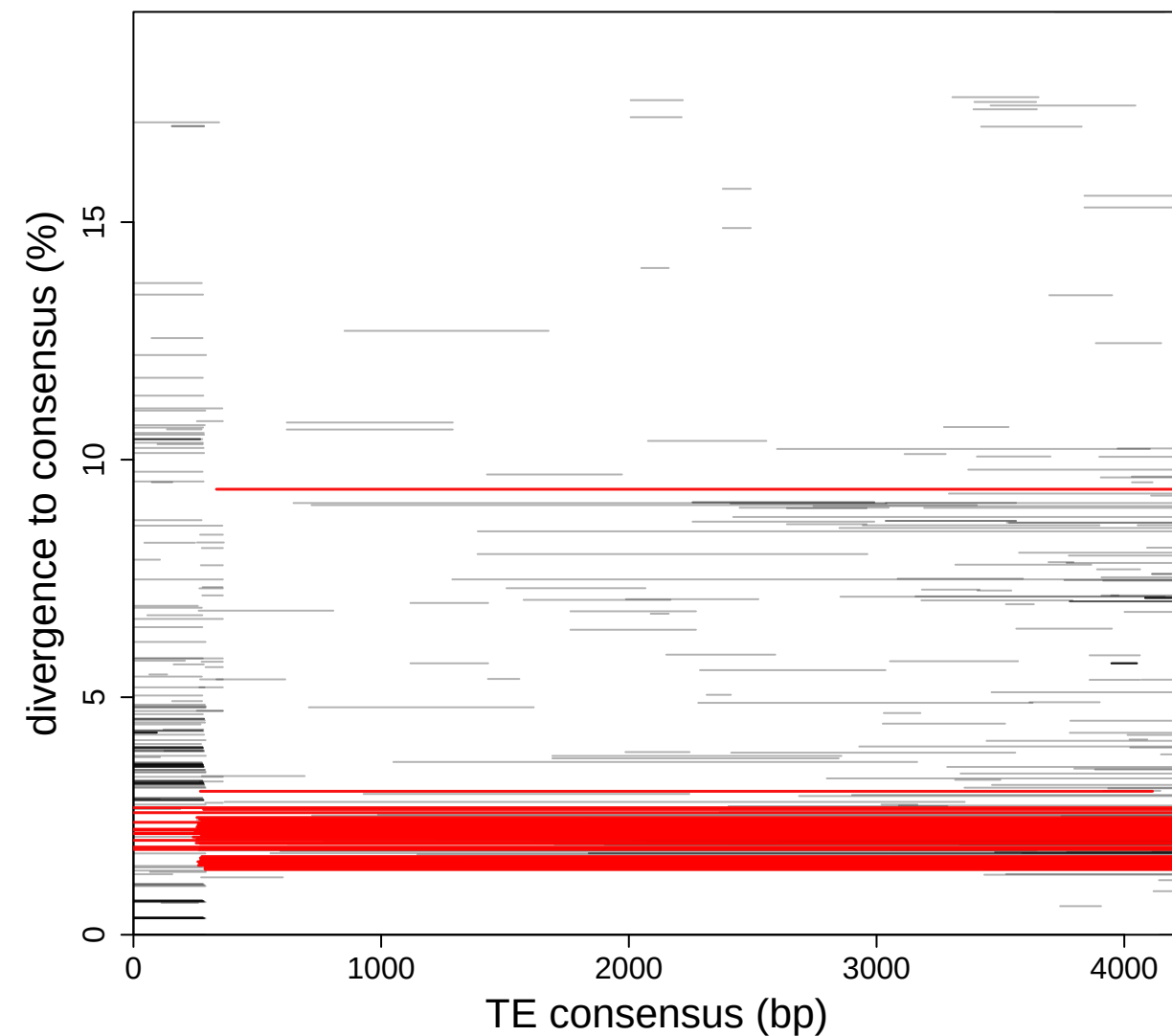
uf.bed g 1.bed fm 1.bed 0 0 n.bed g 1.bed fm 2.bed 0 0 bc
size: 10124bp; fragments: 296; full length: 0 (>=9111.6bp)

After TEtrimmer Extended plot Blue lines are boundaries

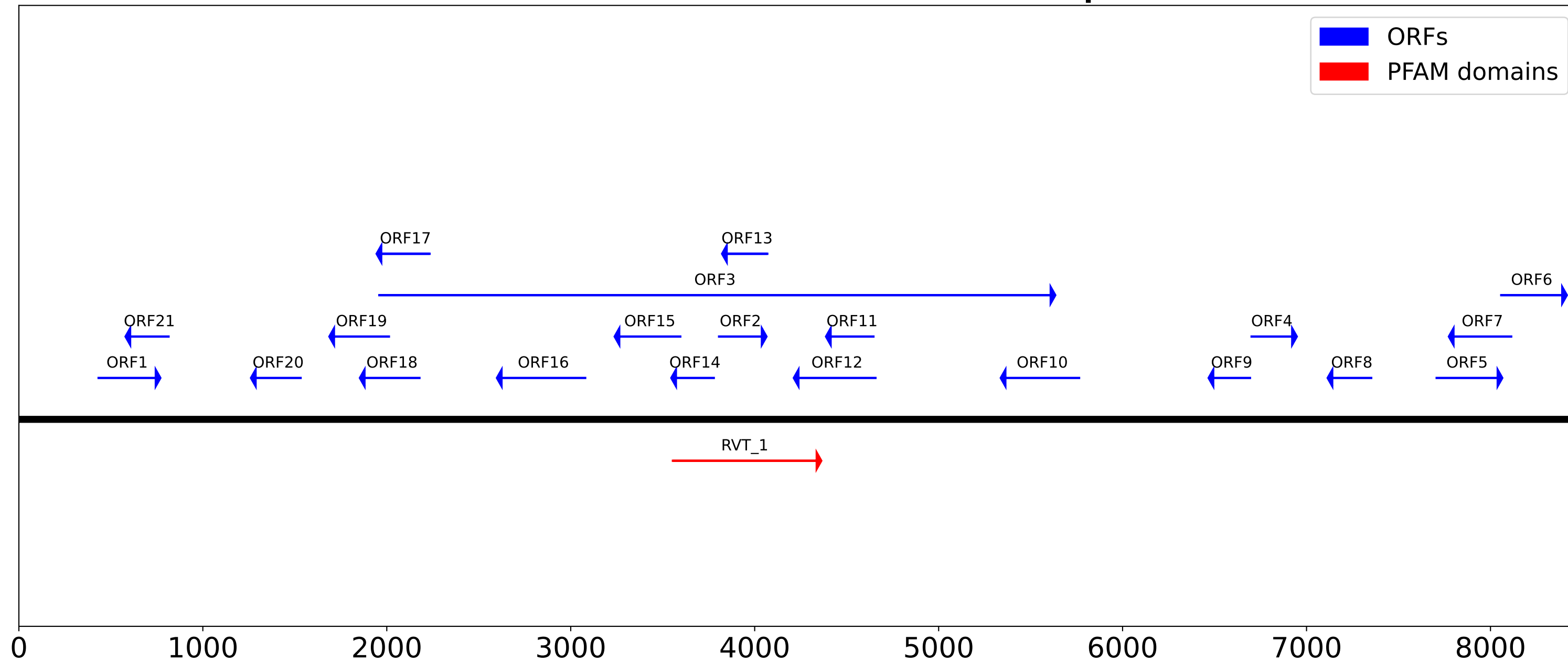


TE: rnd_5_family_11215
size: 4226bp; fragments: 497; full length: 66 (≥ 3803.4 bp)

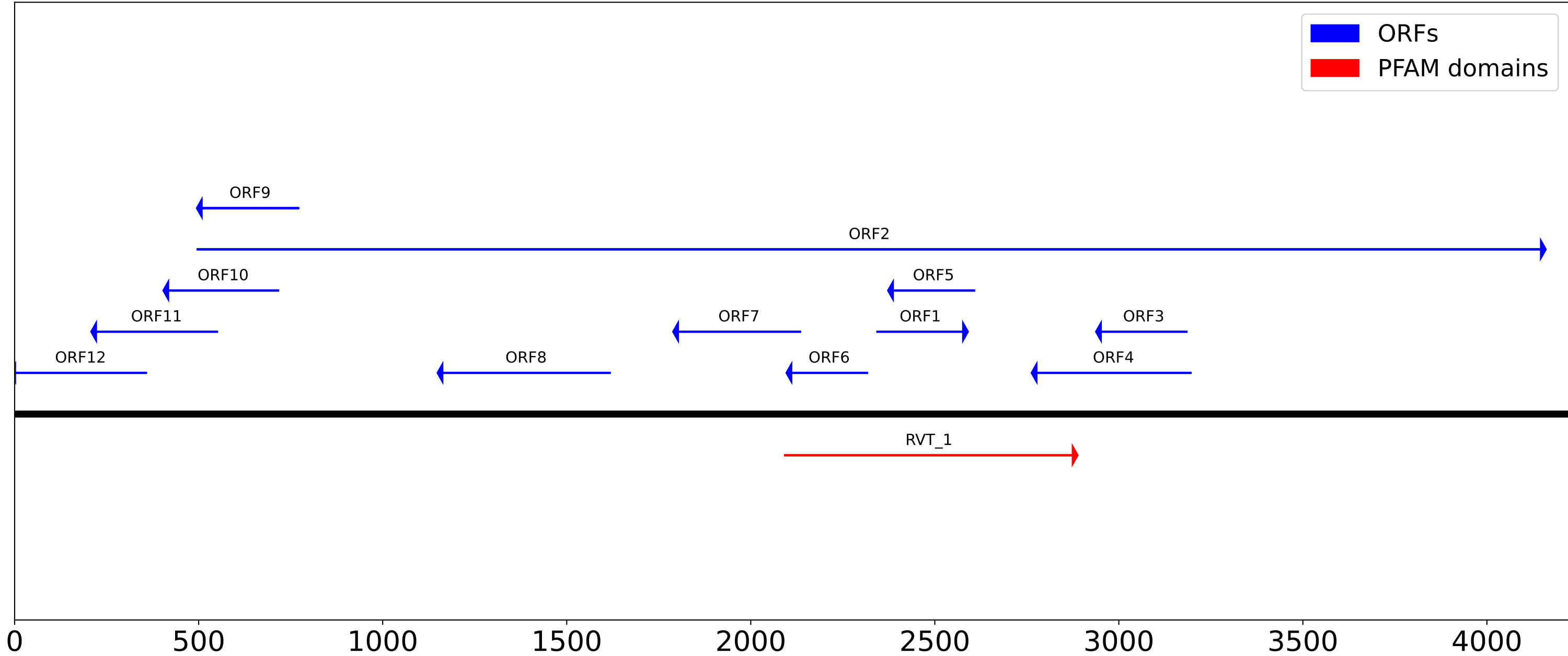
Before TEtrimmer 4226 bp



After TEtrimmer ORF and PFAM domain plot



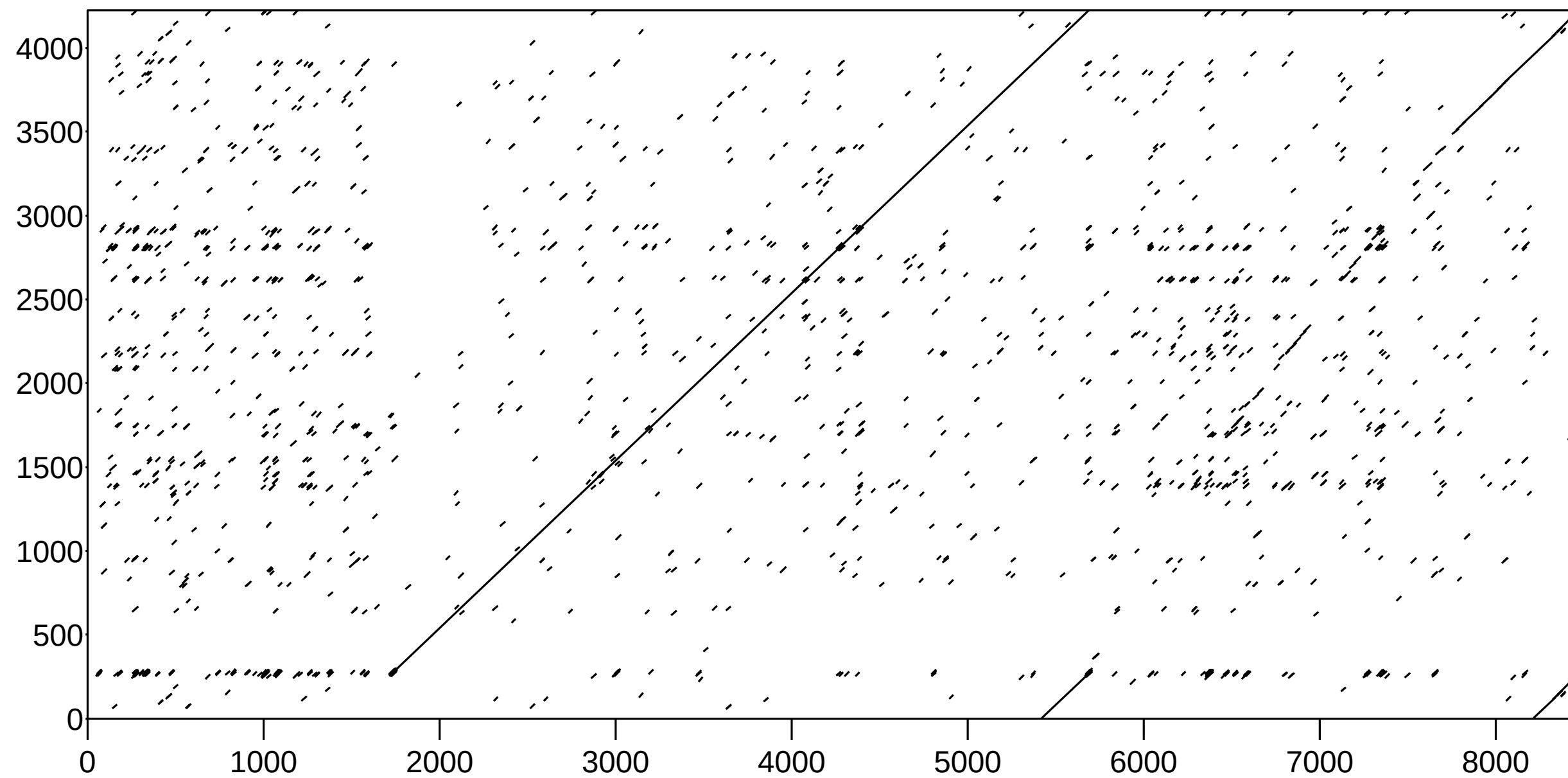
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)