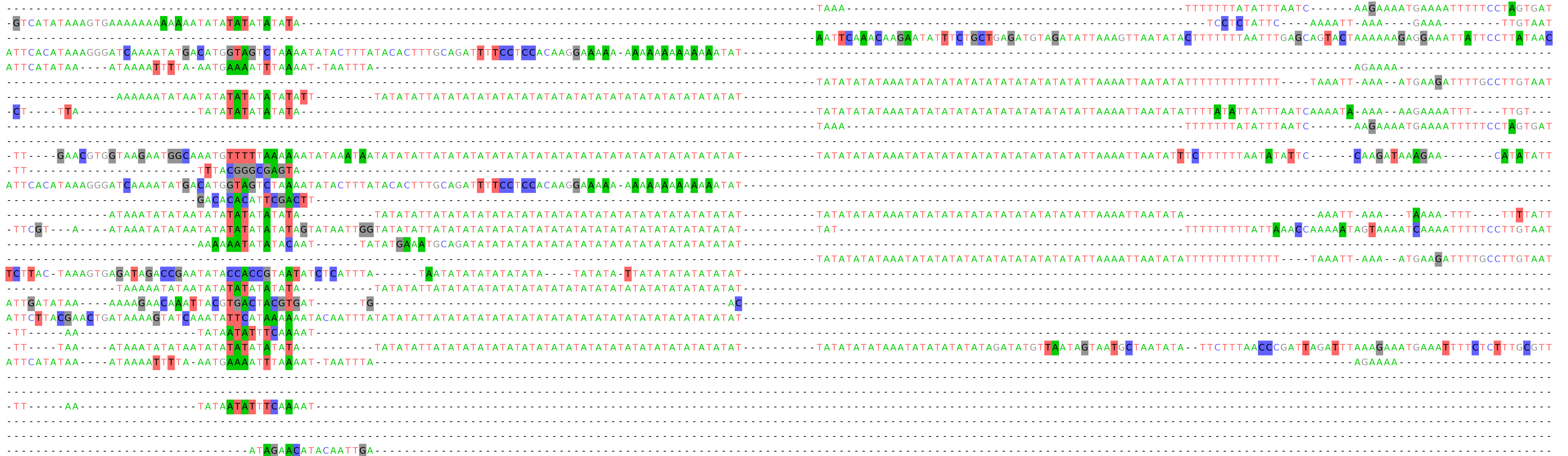
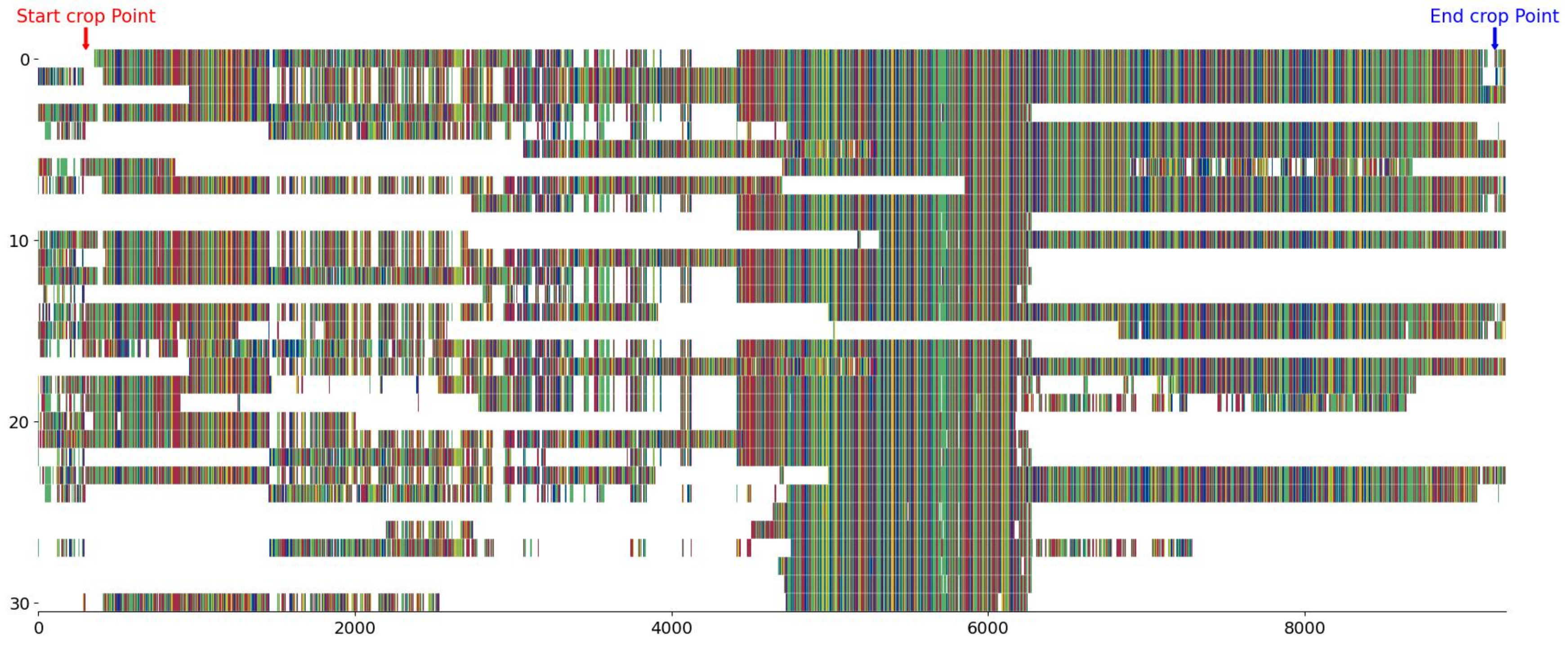


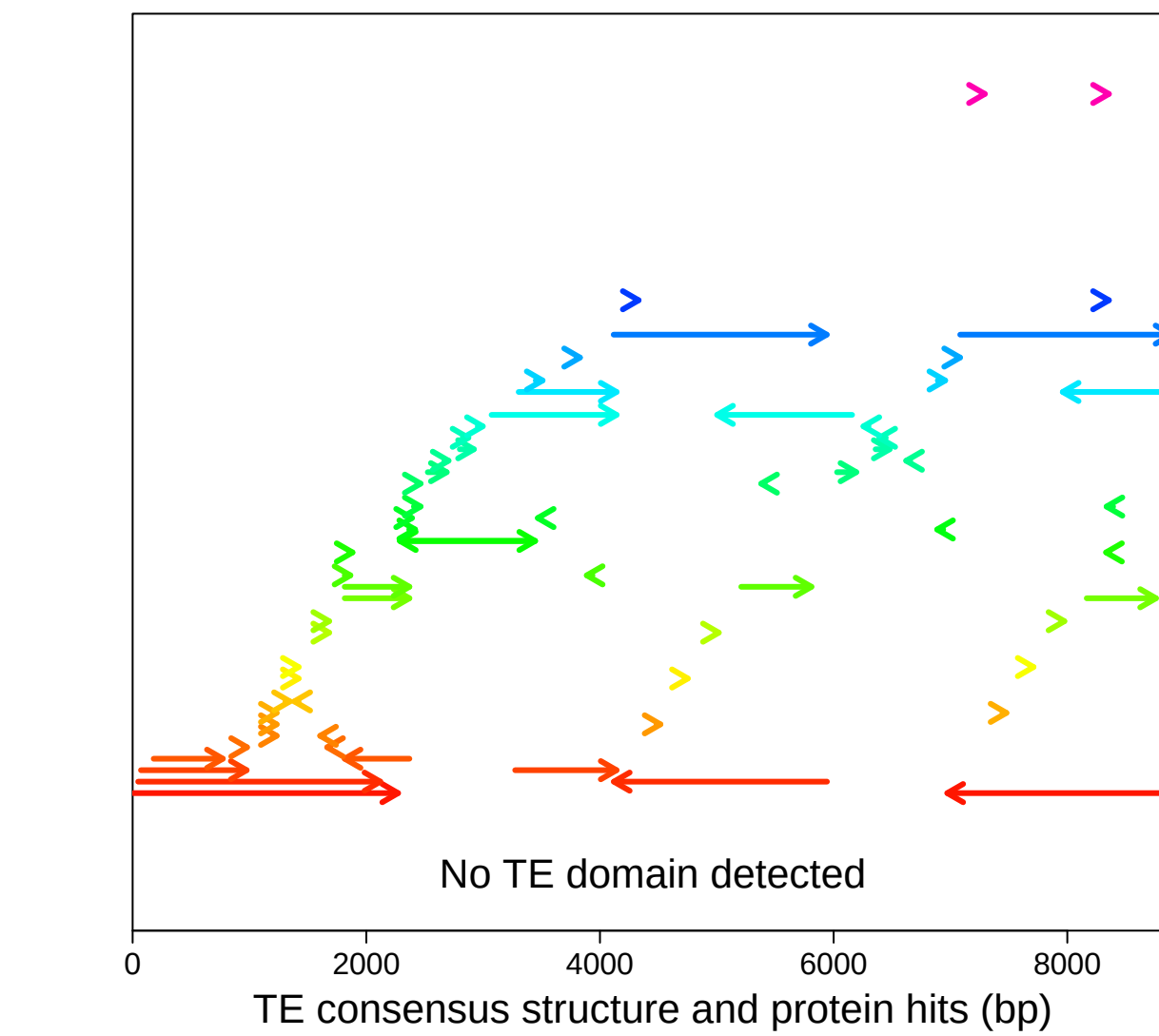
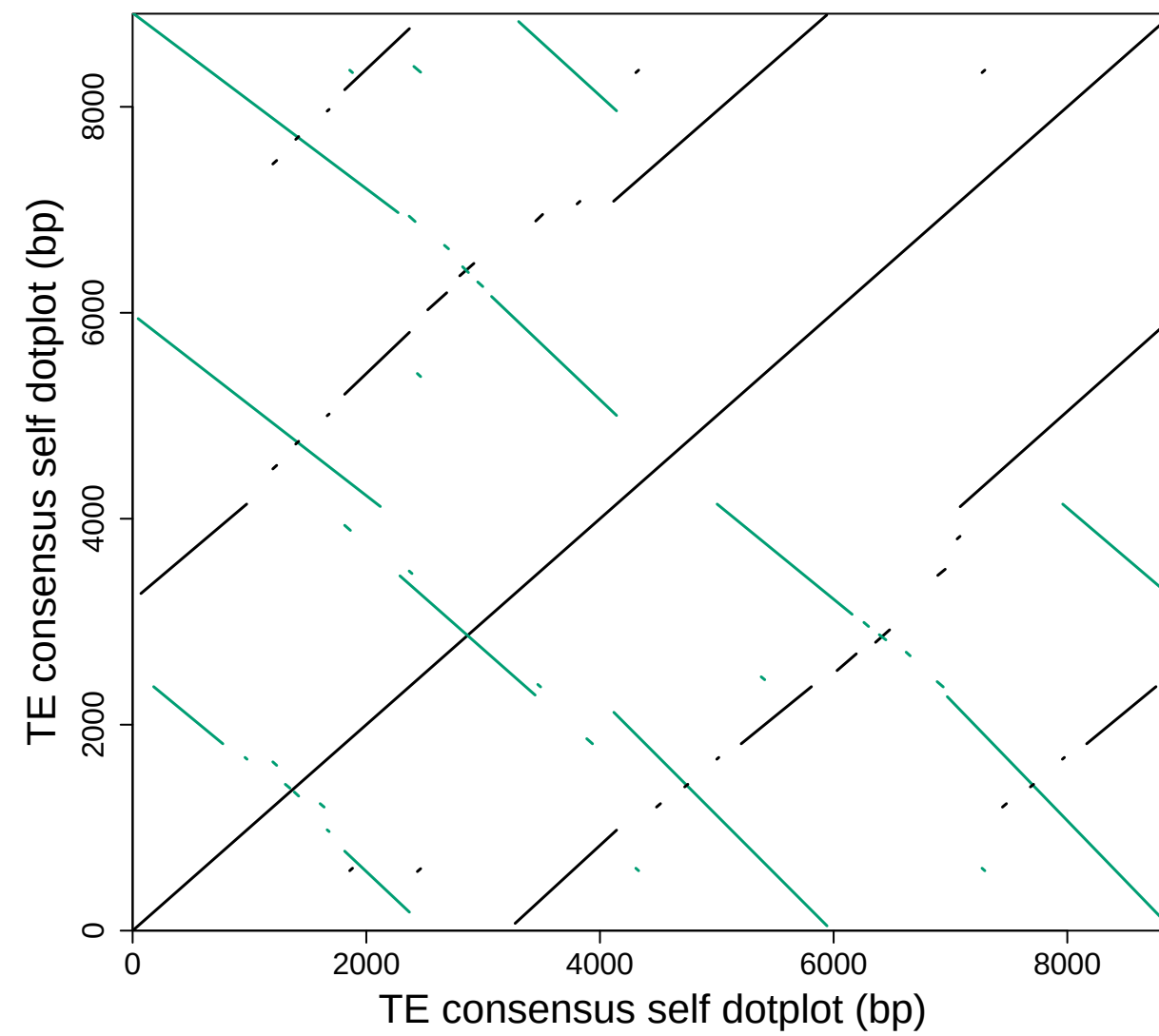
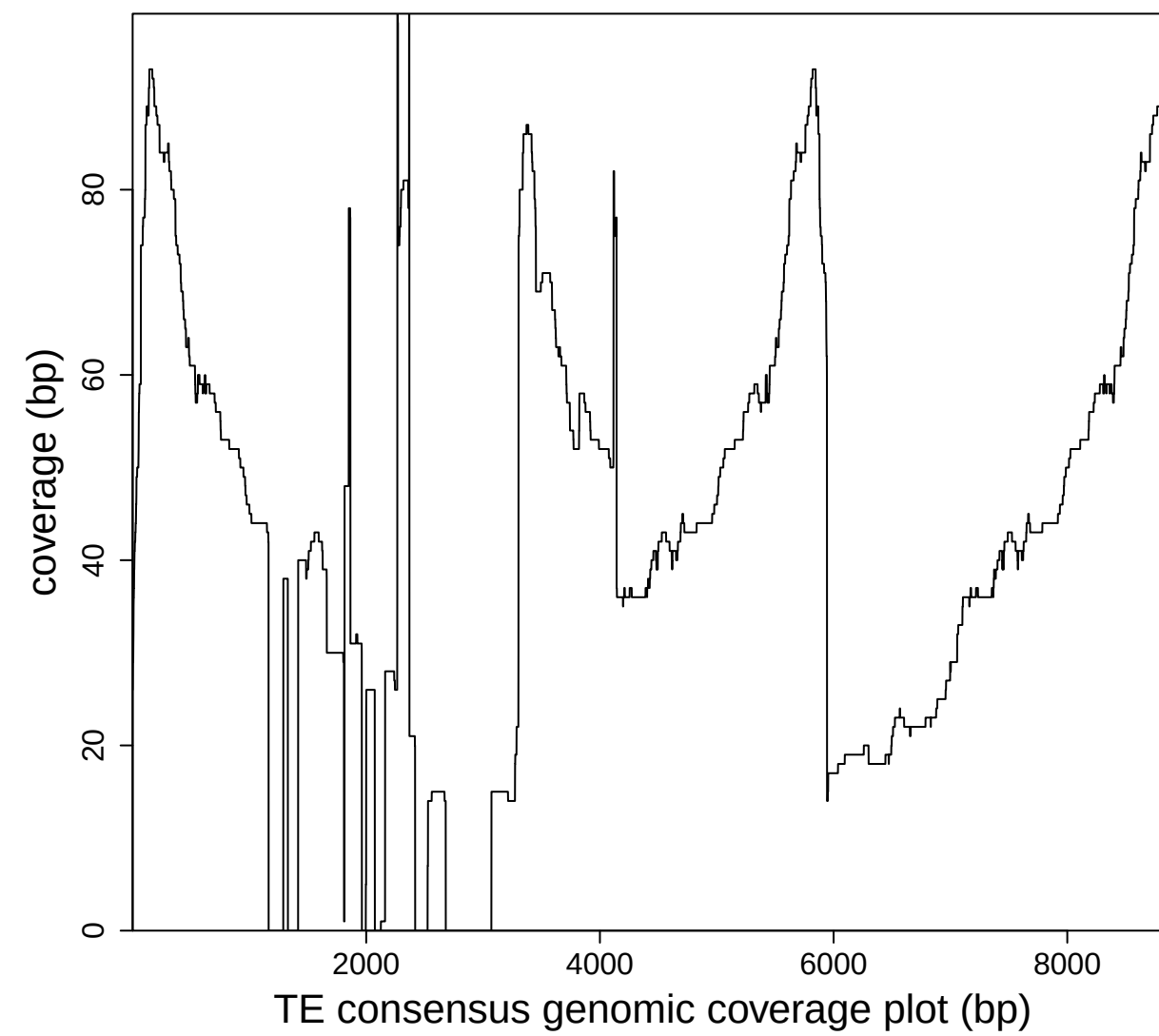
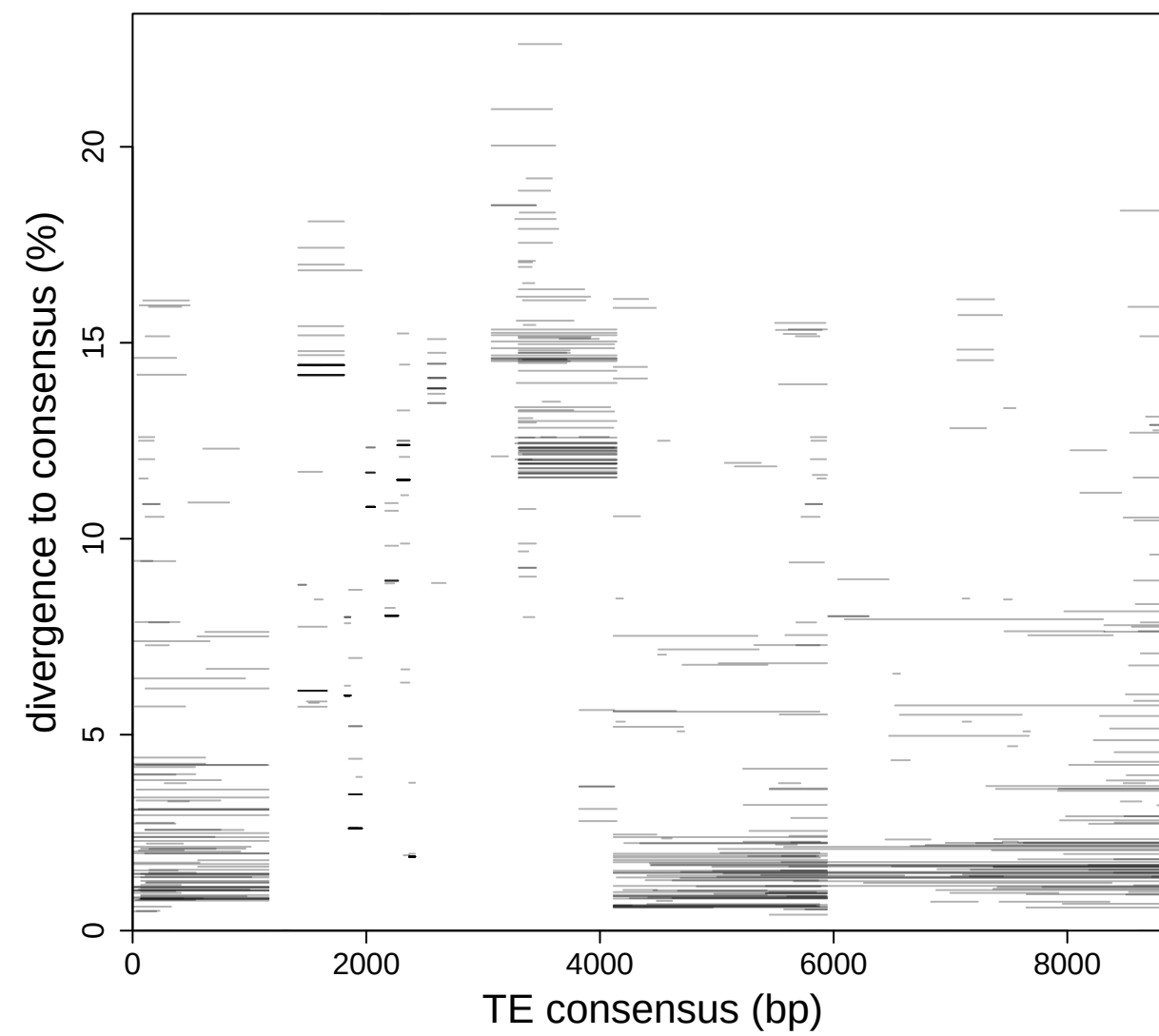
MSA length = 8904





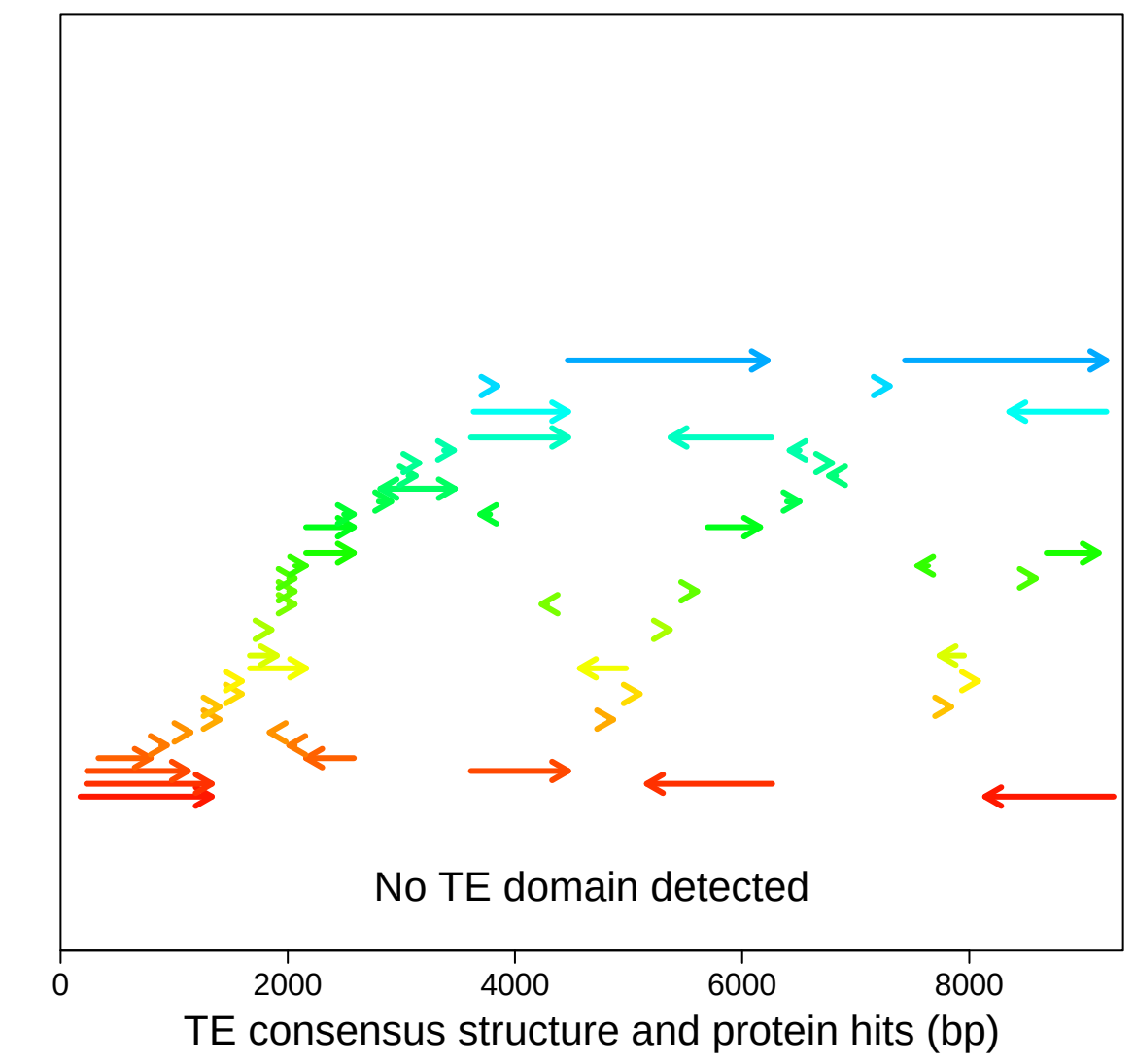
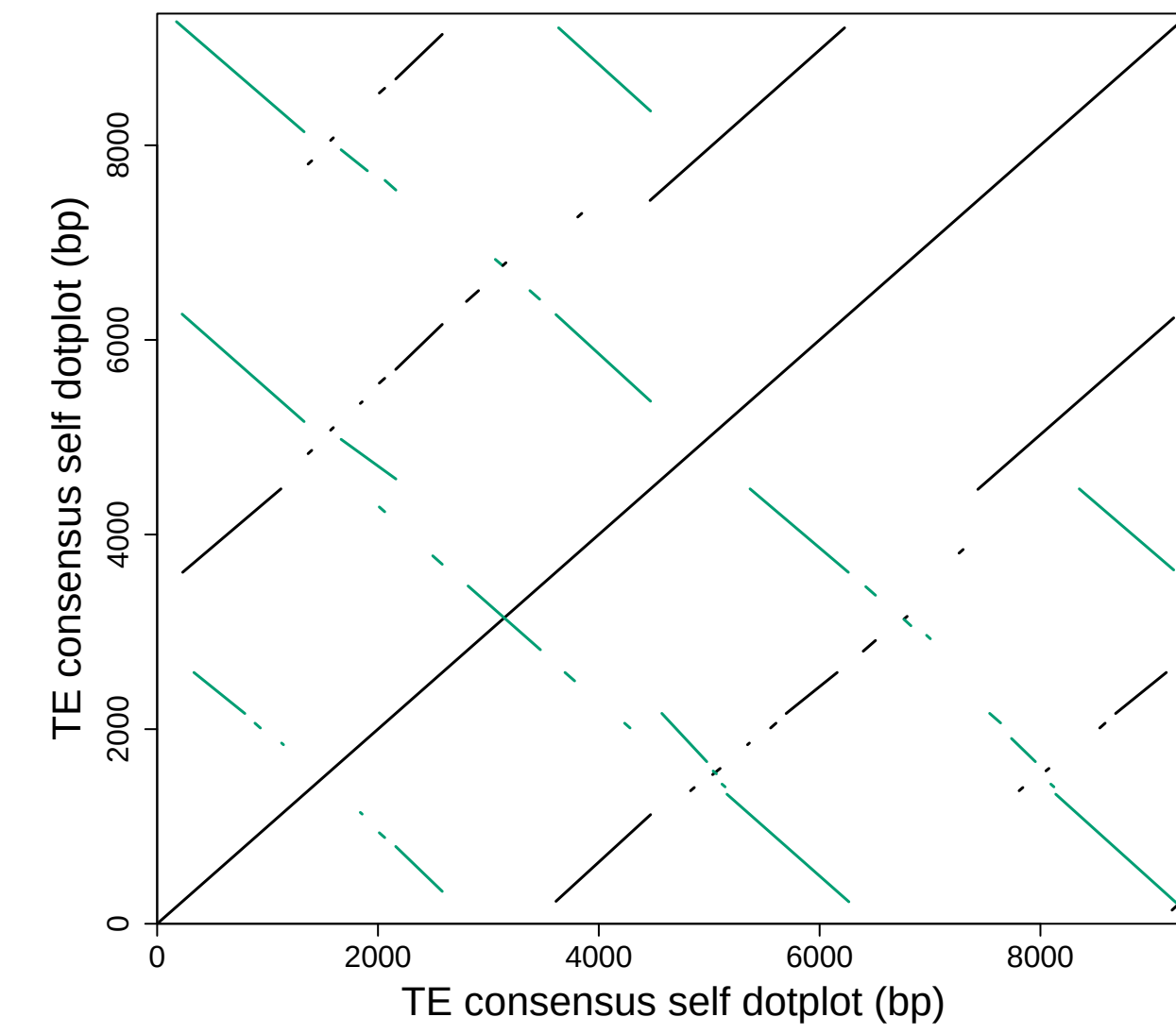
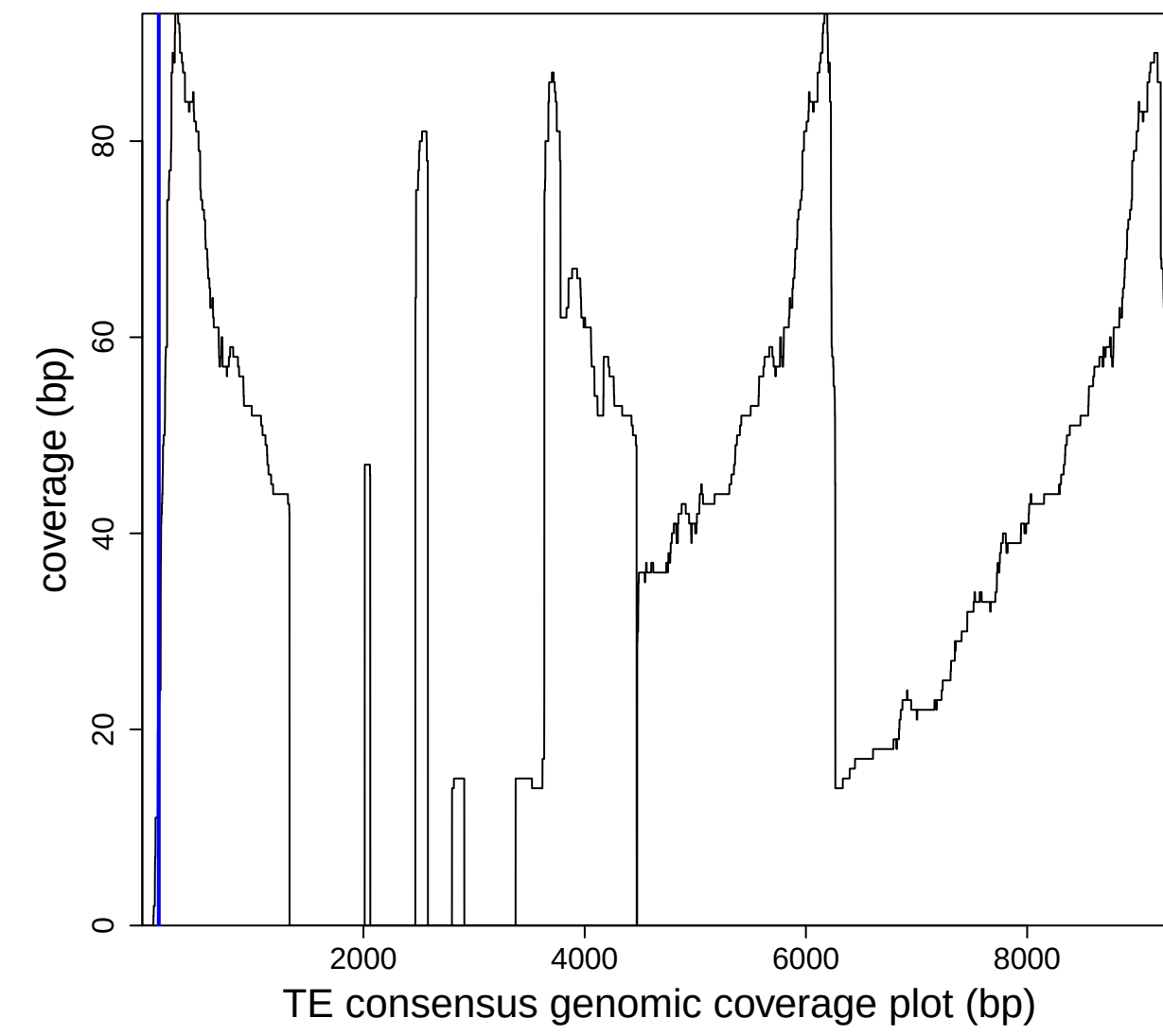
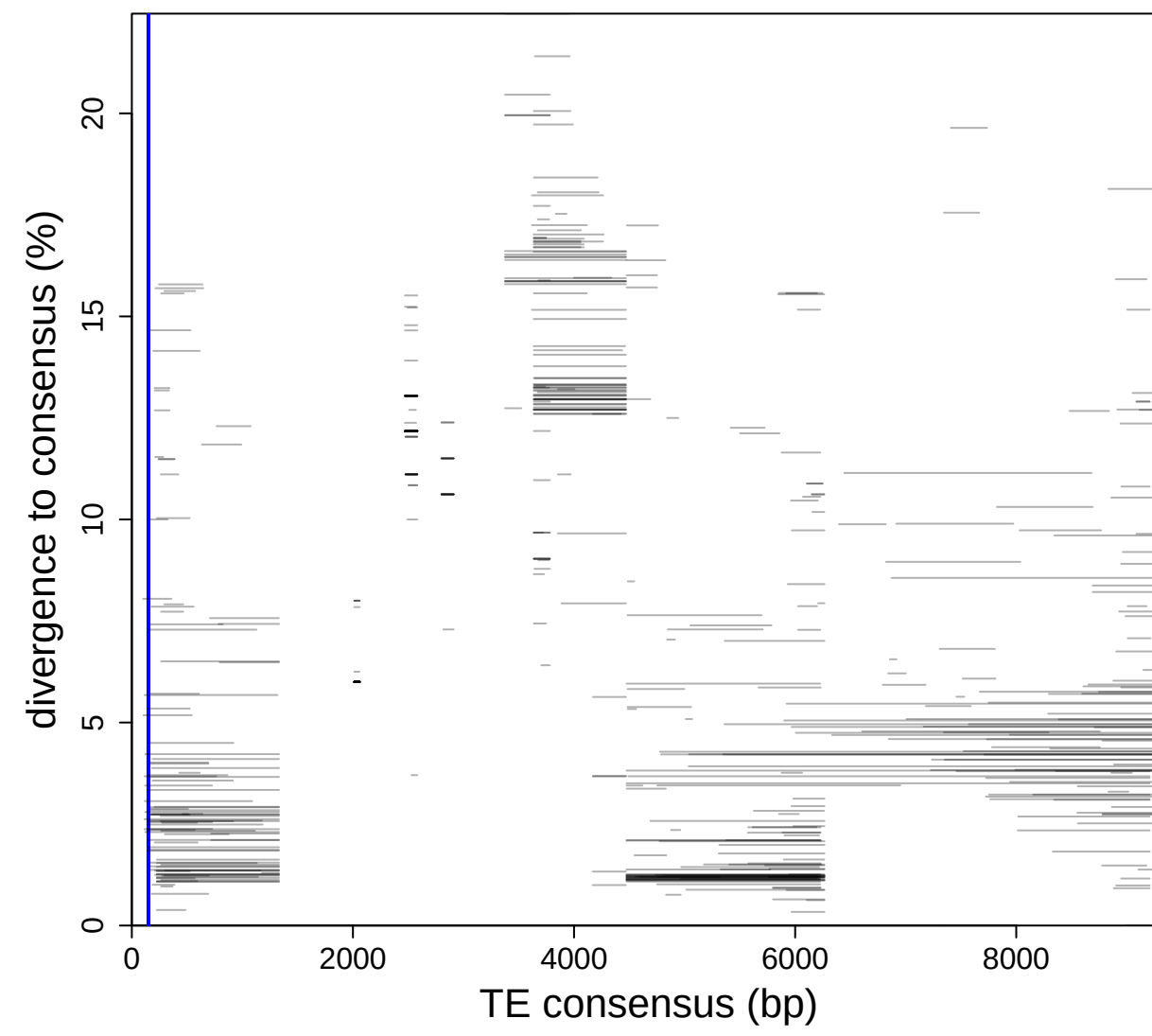
bed_fm_1.bed 0 0 n.bed g 1.bed fm_2.bed 0 0 bcln.fa aln.fa
size: 8904bp; fragments: 763; full length: 0 (>=8013.6bp)

After TETrimmer 8904 bp



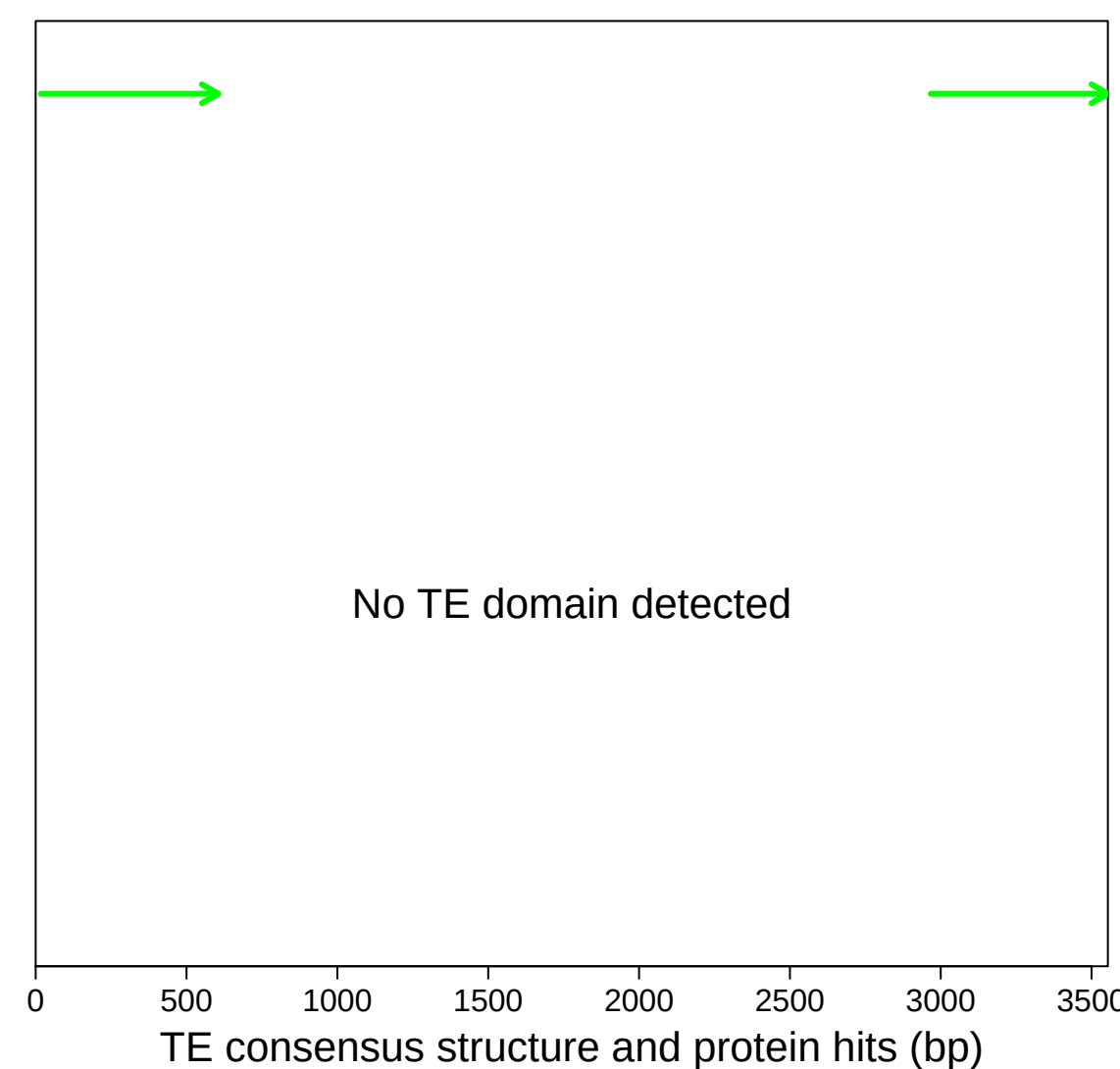
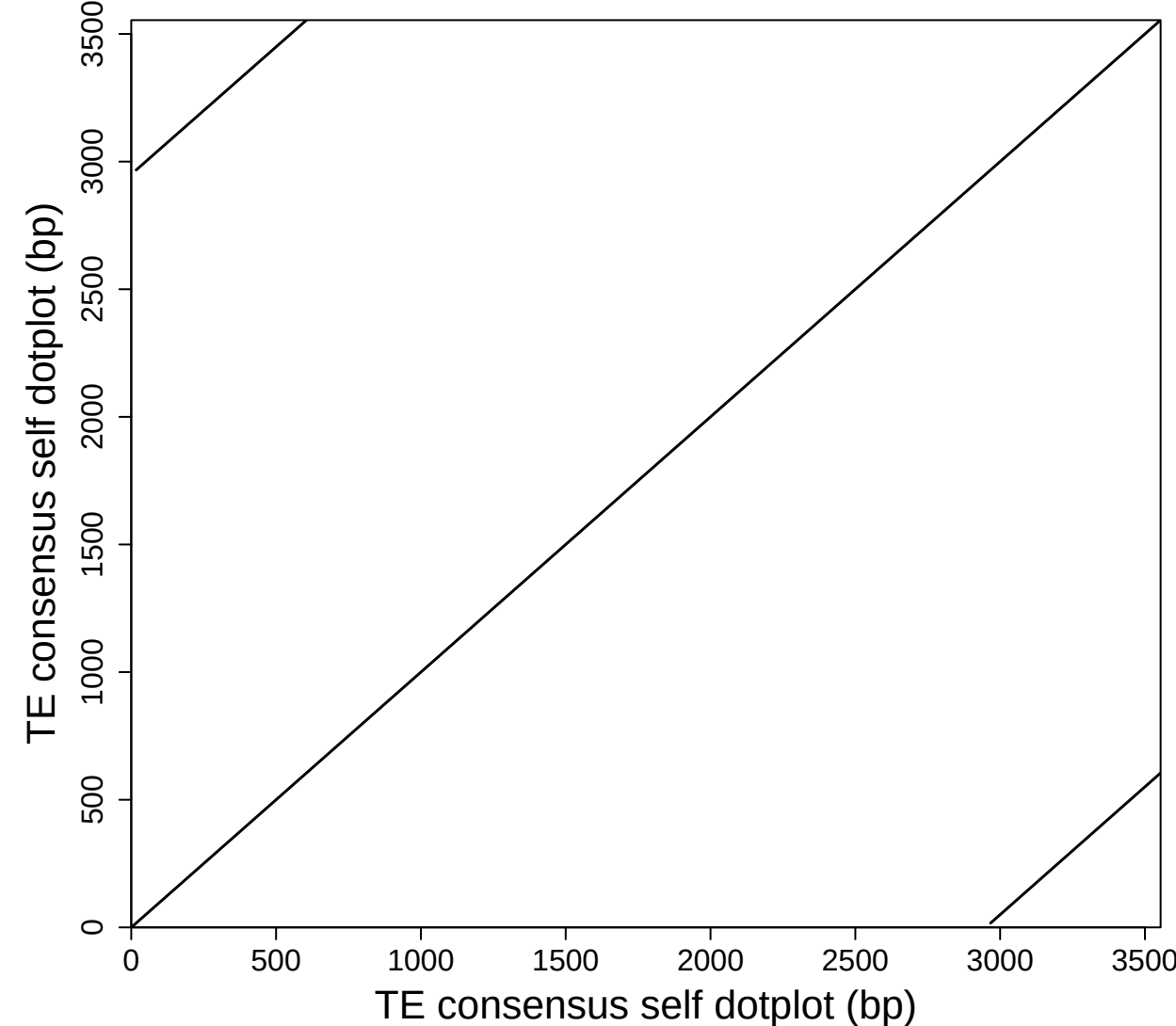
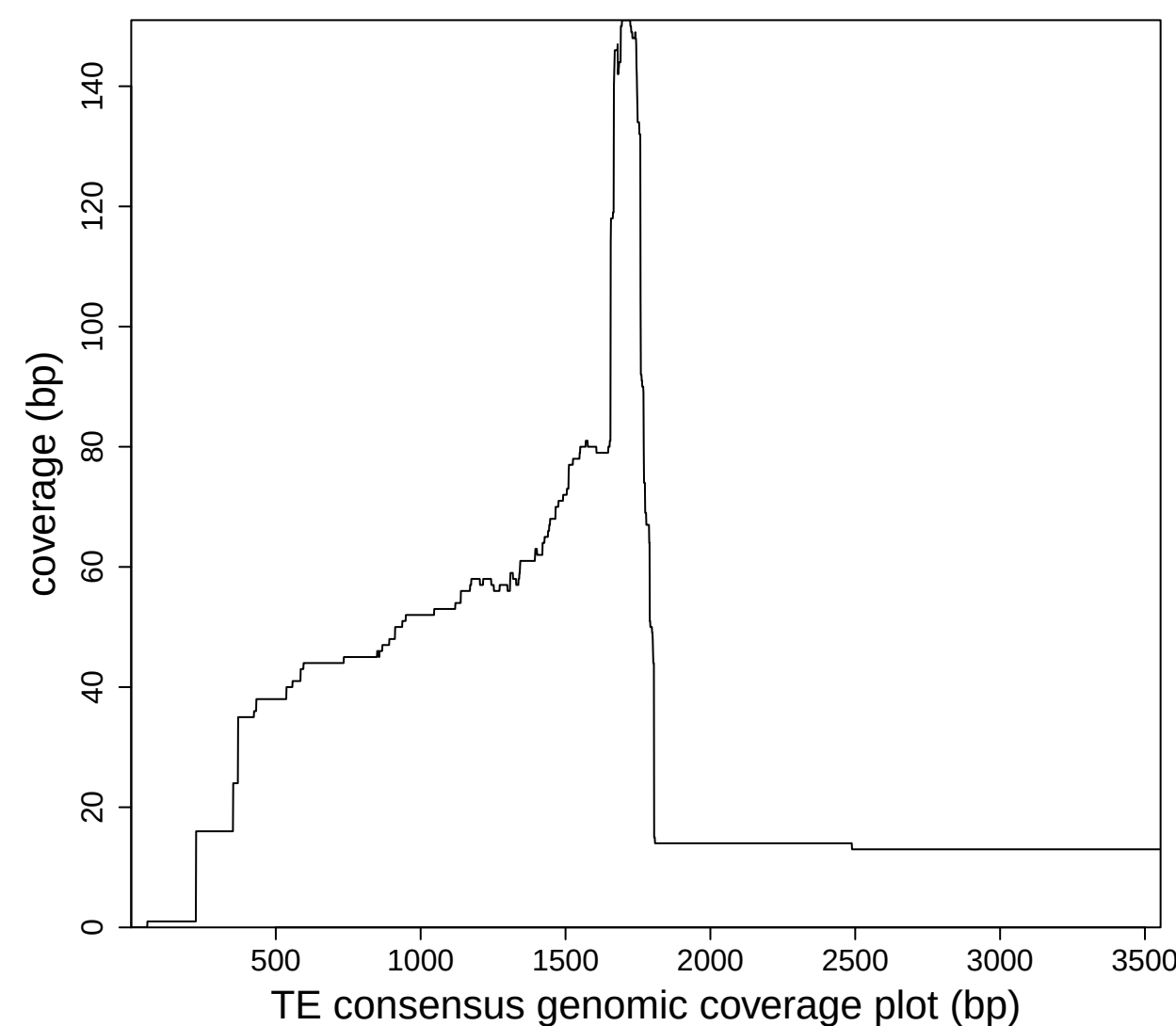
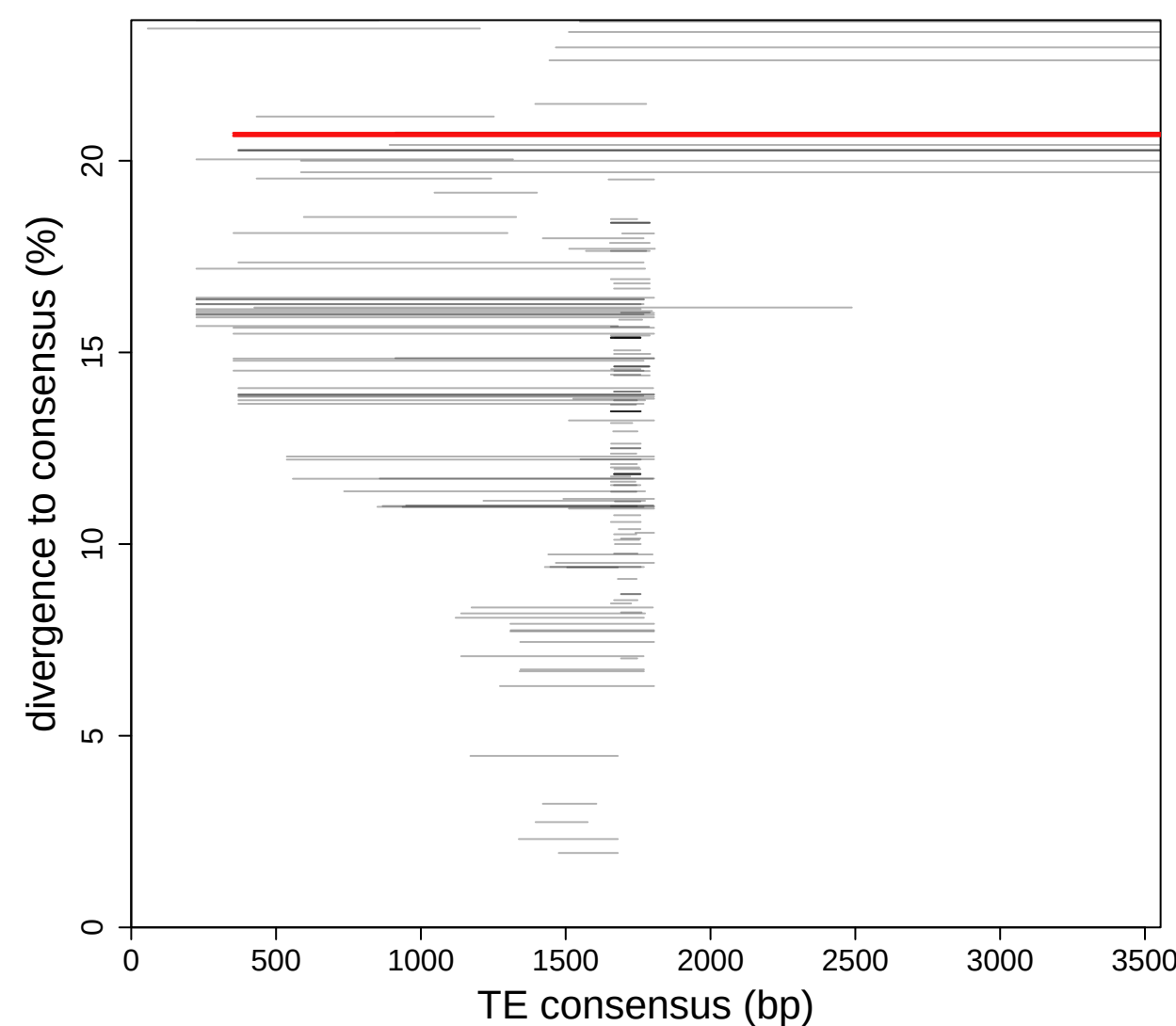
f.bed_g_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcl
size: 9353bp; fragments: 566; full length: 0 (>=8417.7bp)

After TEtrimmer Extended plot Blue lines are boundaries

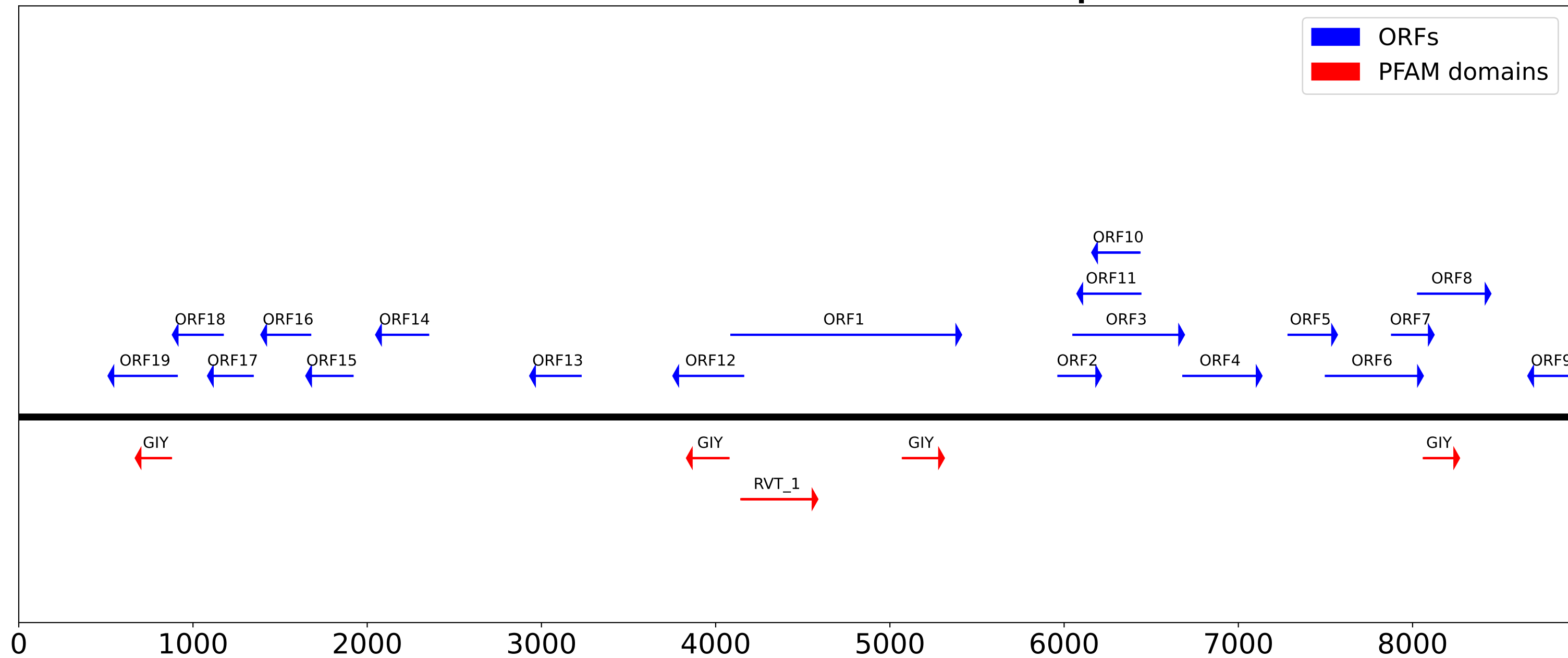


TE: rnd_5_family_946
size: 3554bp; fragments: 167; full length: 2 (≥ 3198.6 bp)

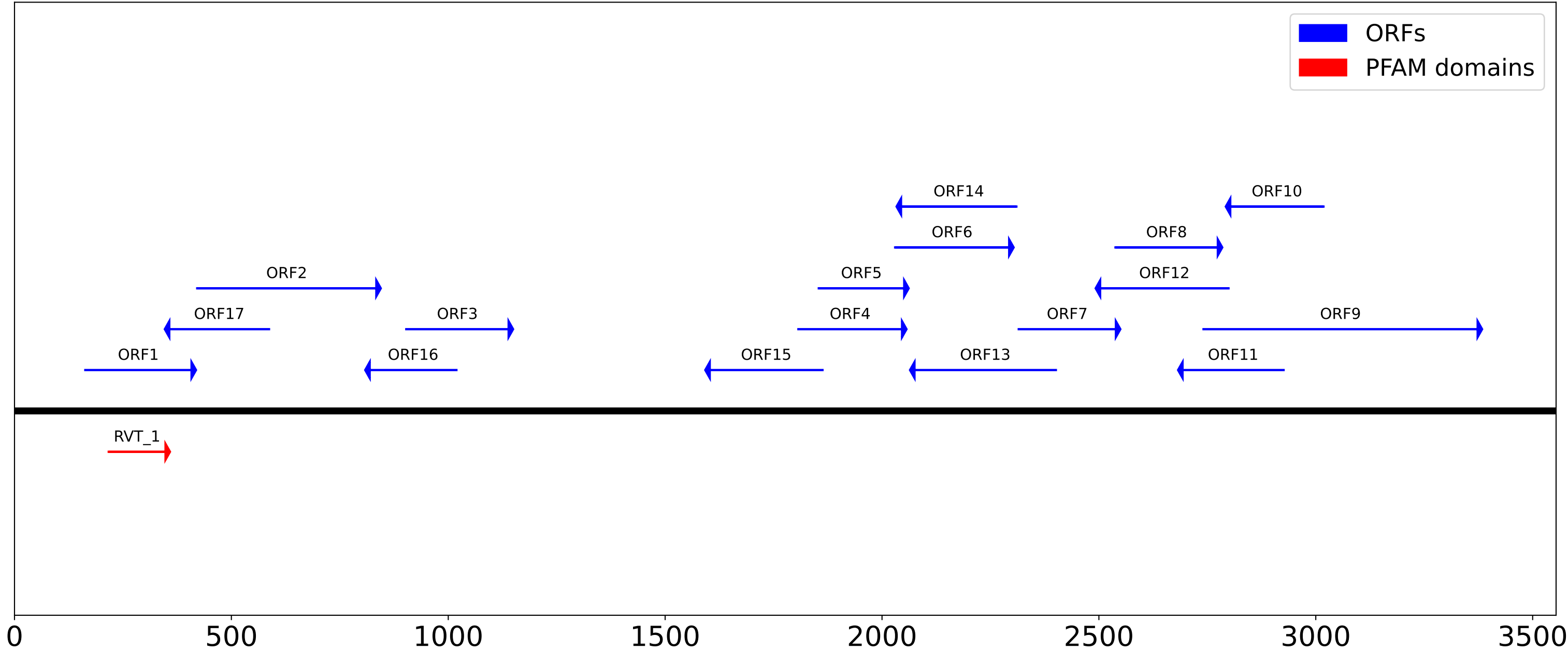
Before TEtrimmer 3554 bp



After TEtrimmer ORF and PFAM domain plot



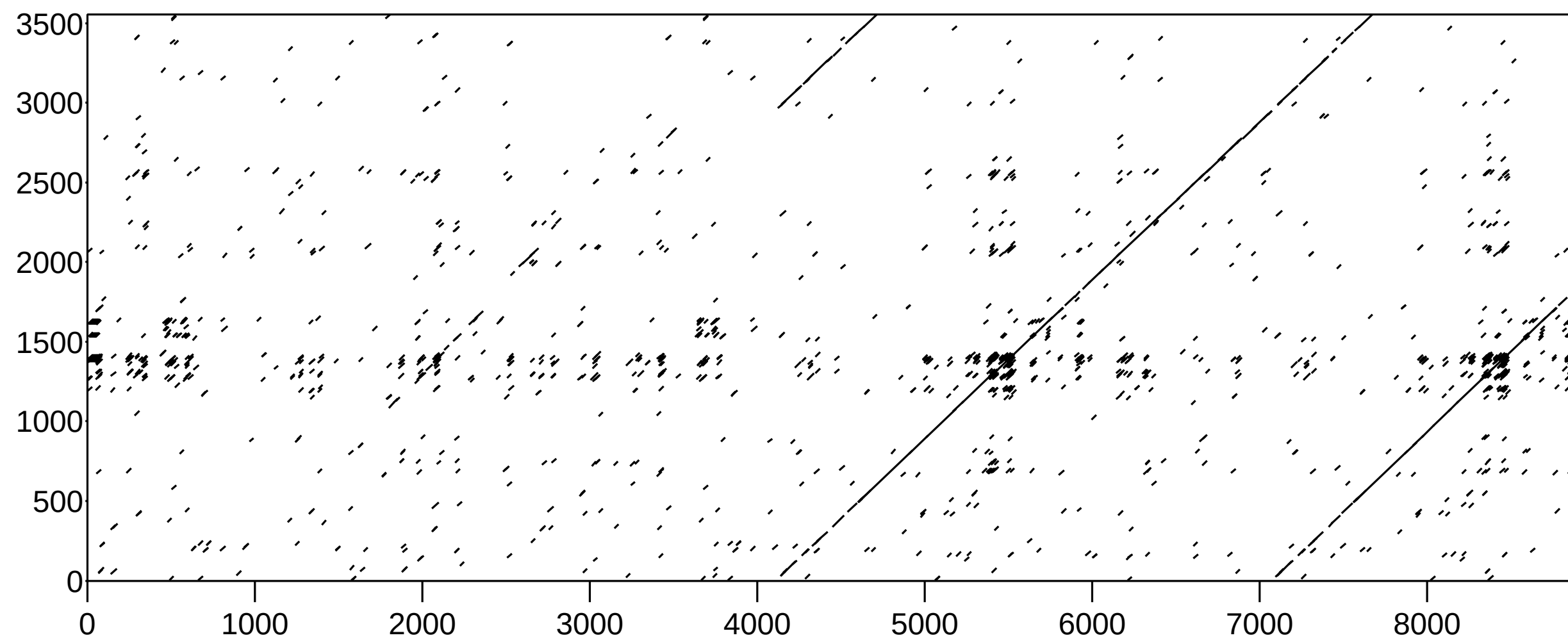
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)