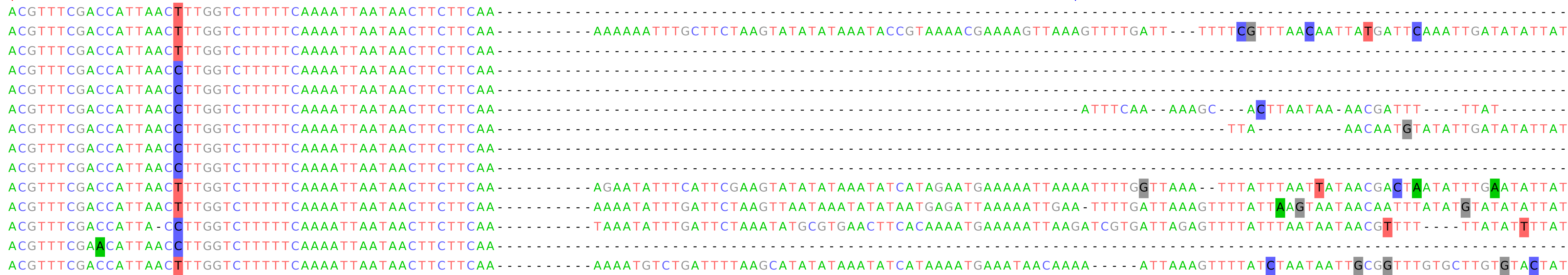
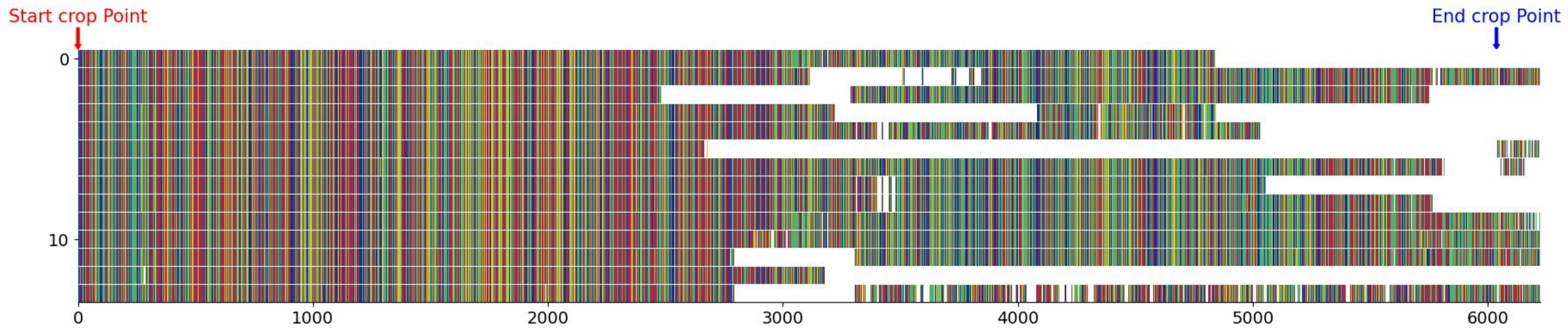


MSA length = 6038

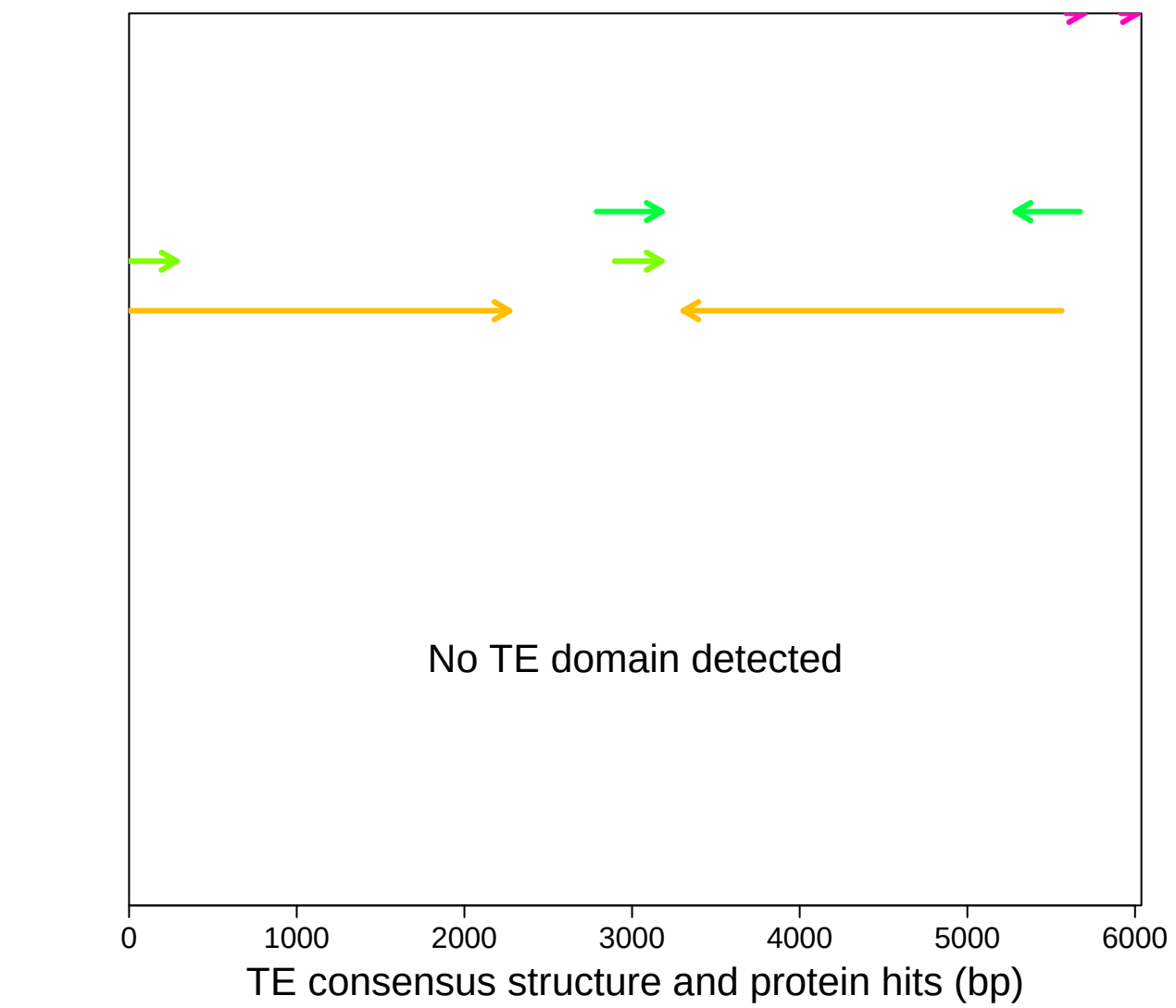
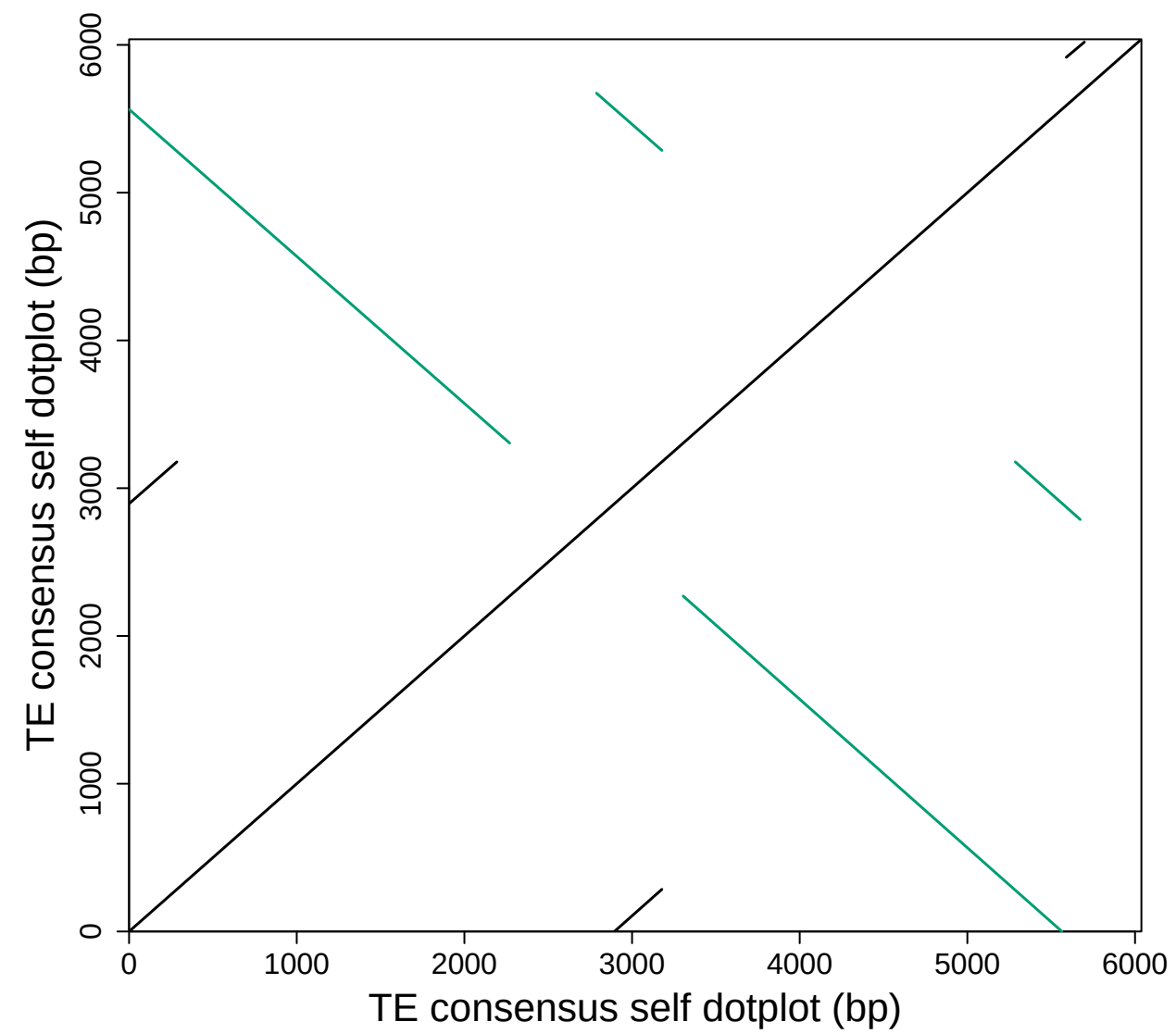
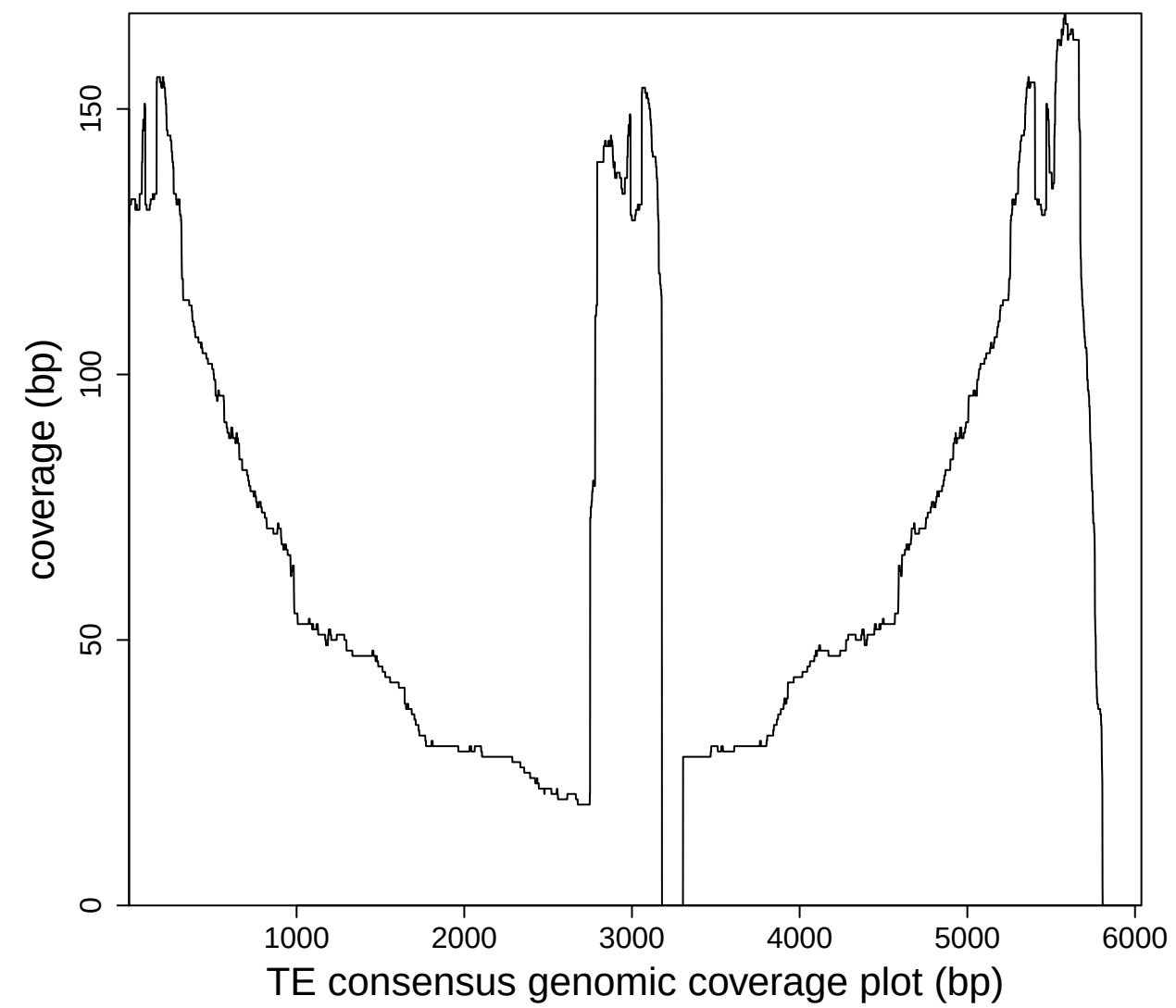
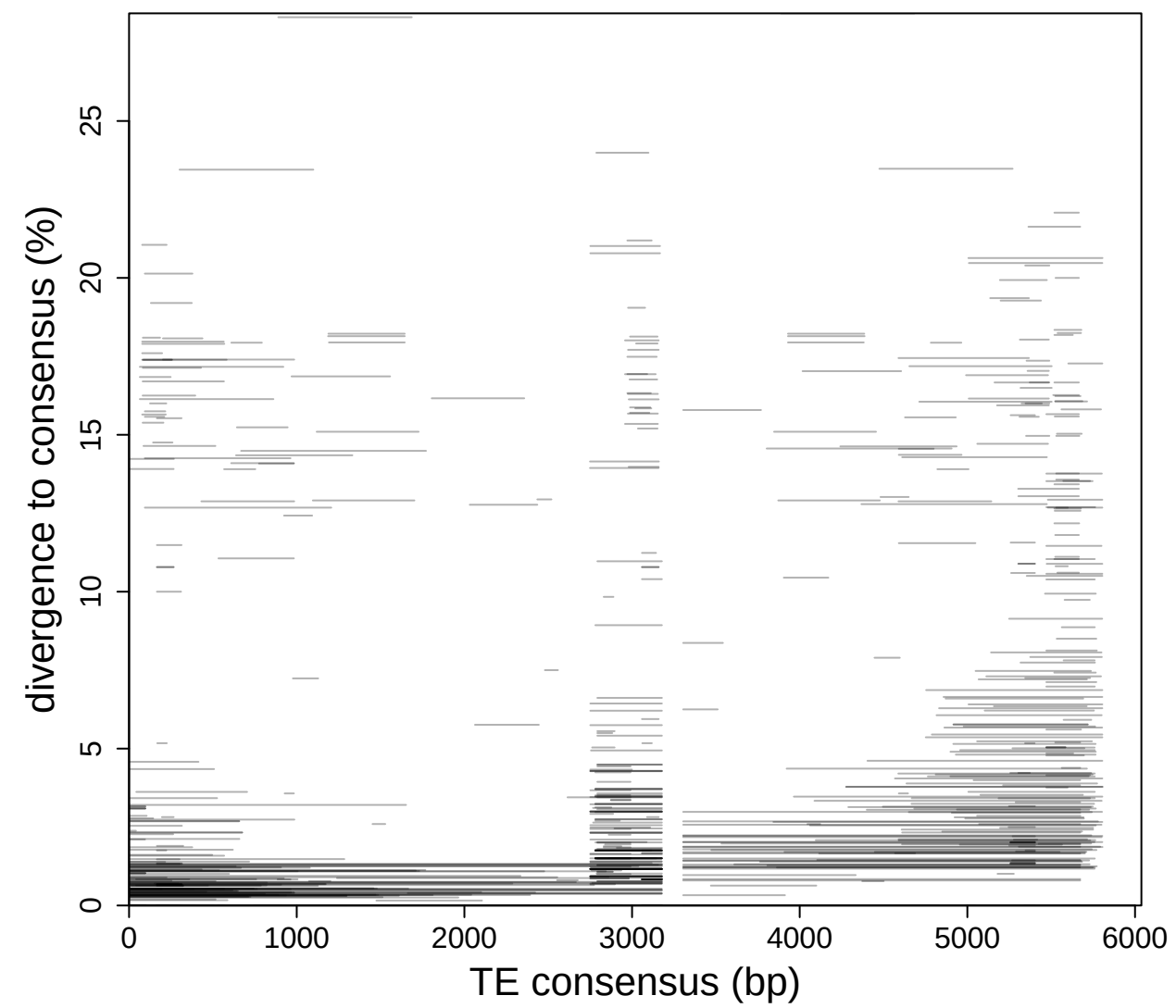
End crop Point





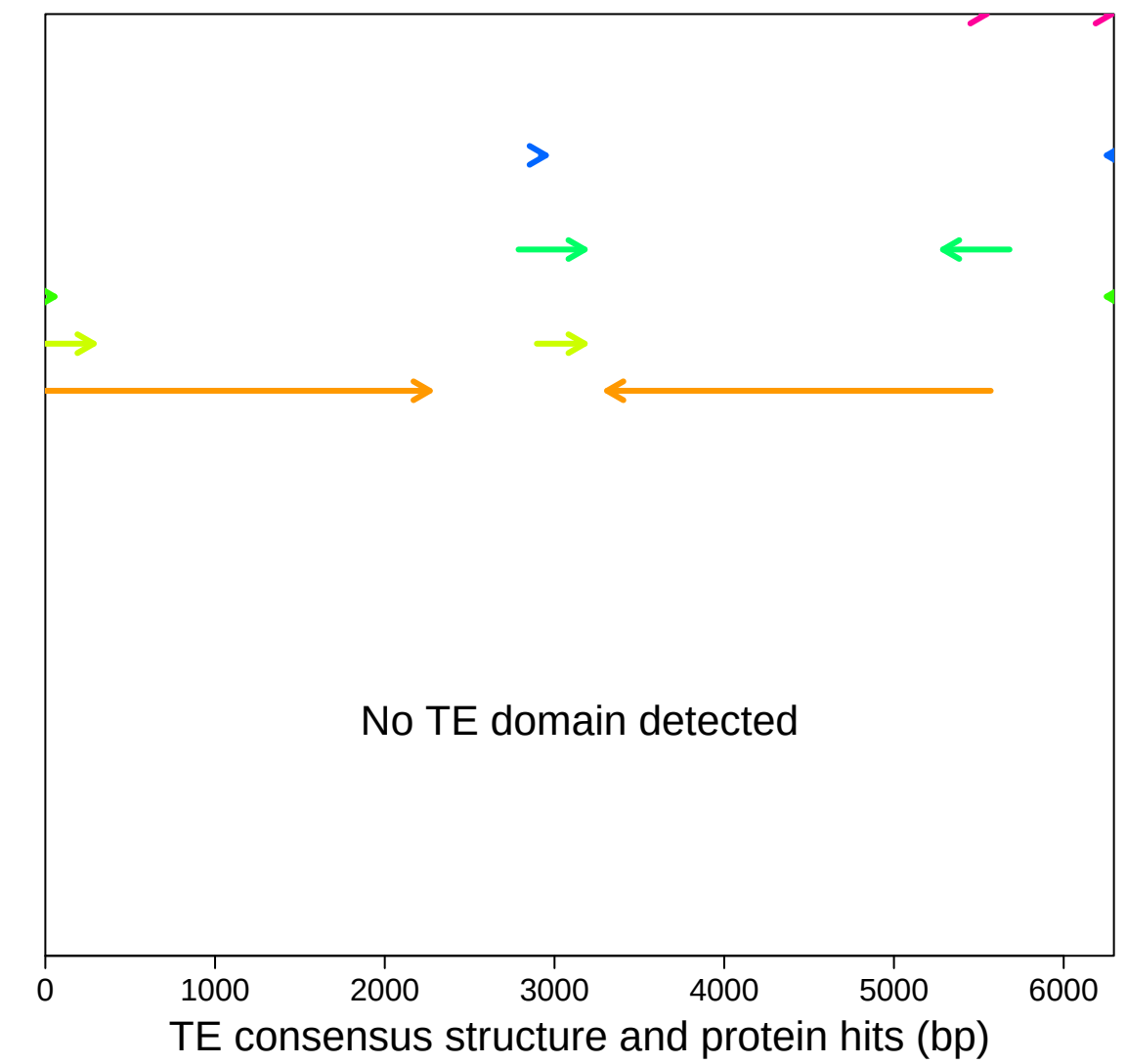
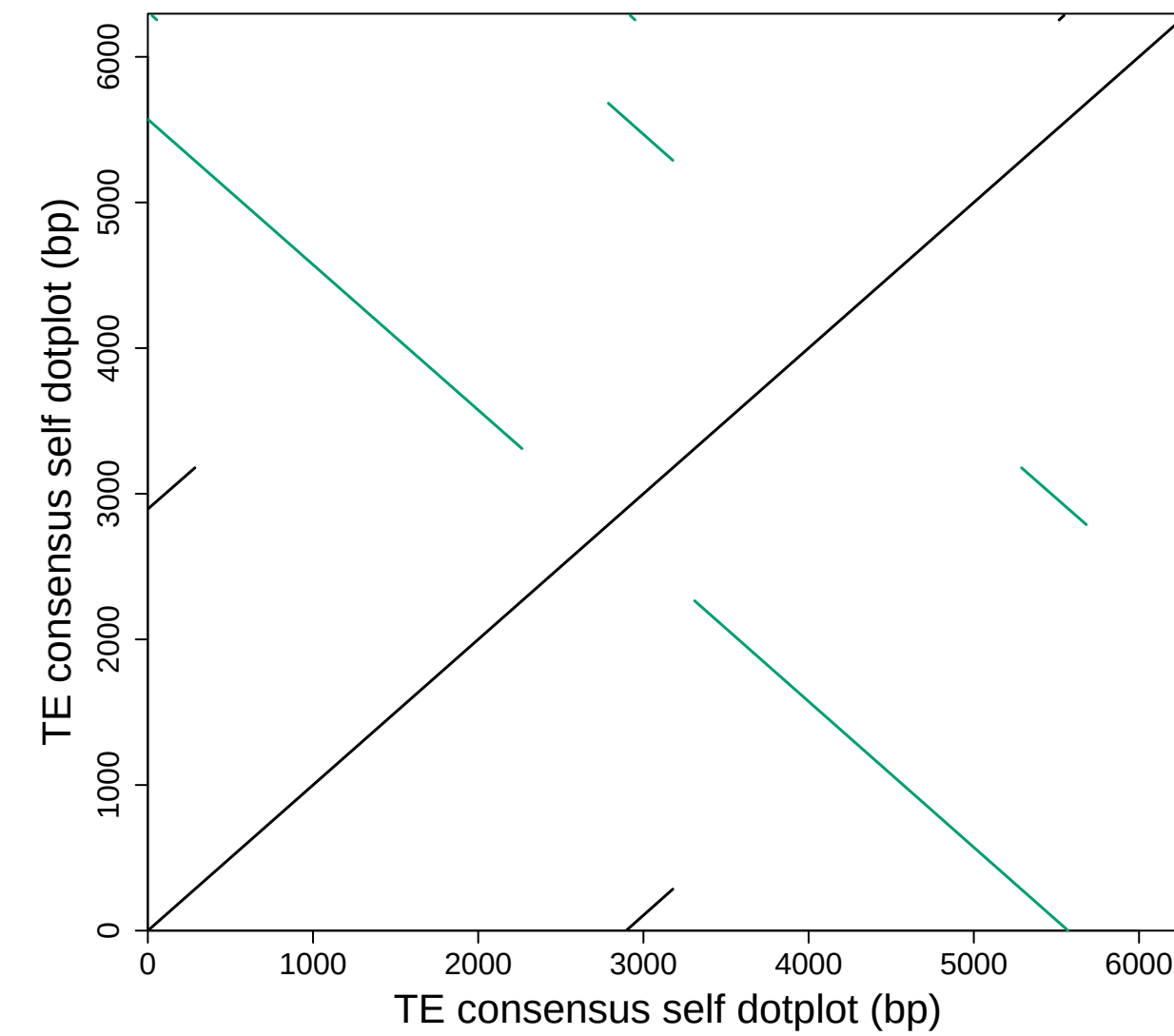
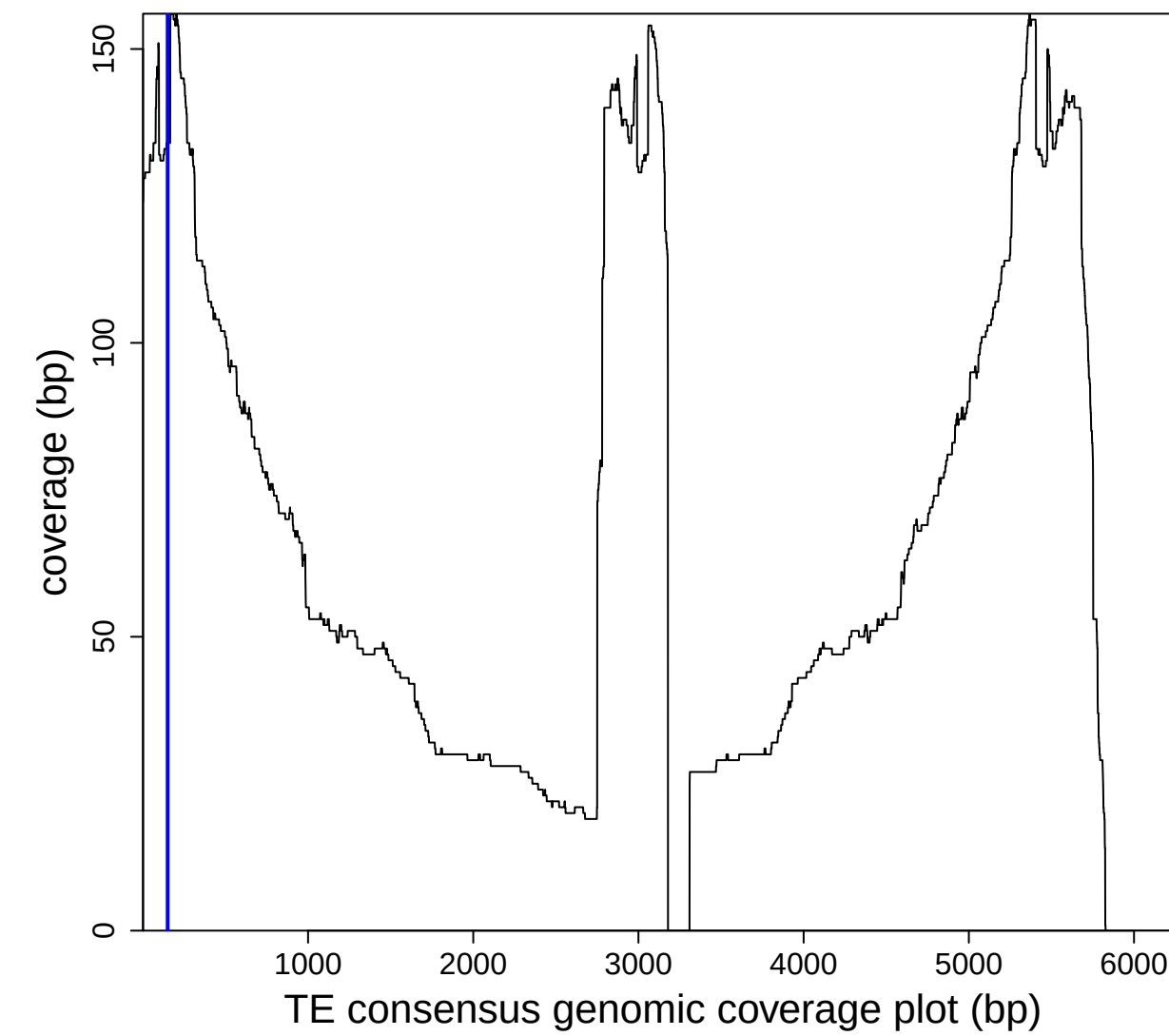
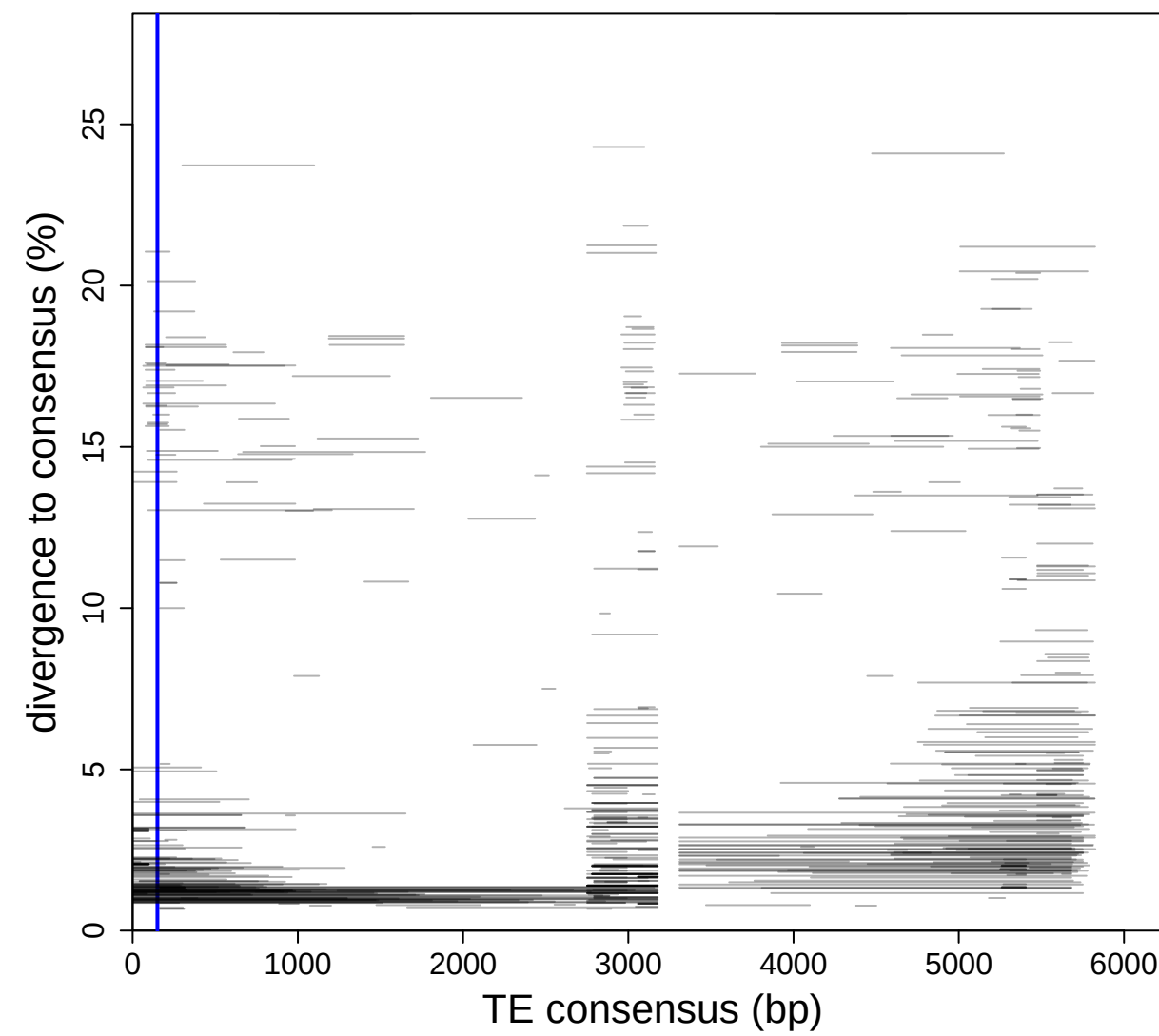
bed_uf.bed g_3.bed fm_1.bed 0_0_bcln.fa aln.fa cl.fa gs.fa c
size: 6038bp; fragments: 651; full length: 0 (>=5434.2bp)

After TETrimmer 6038 bp



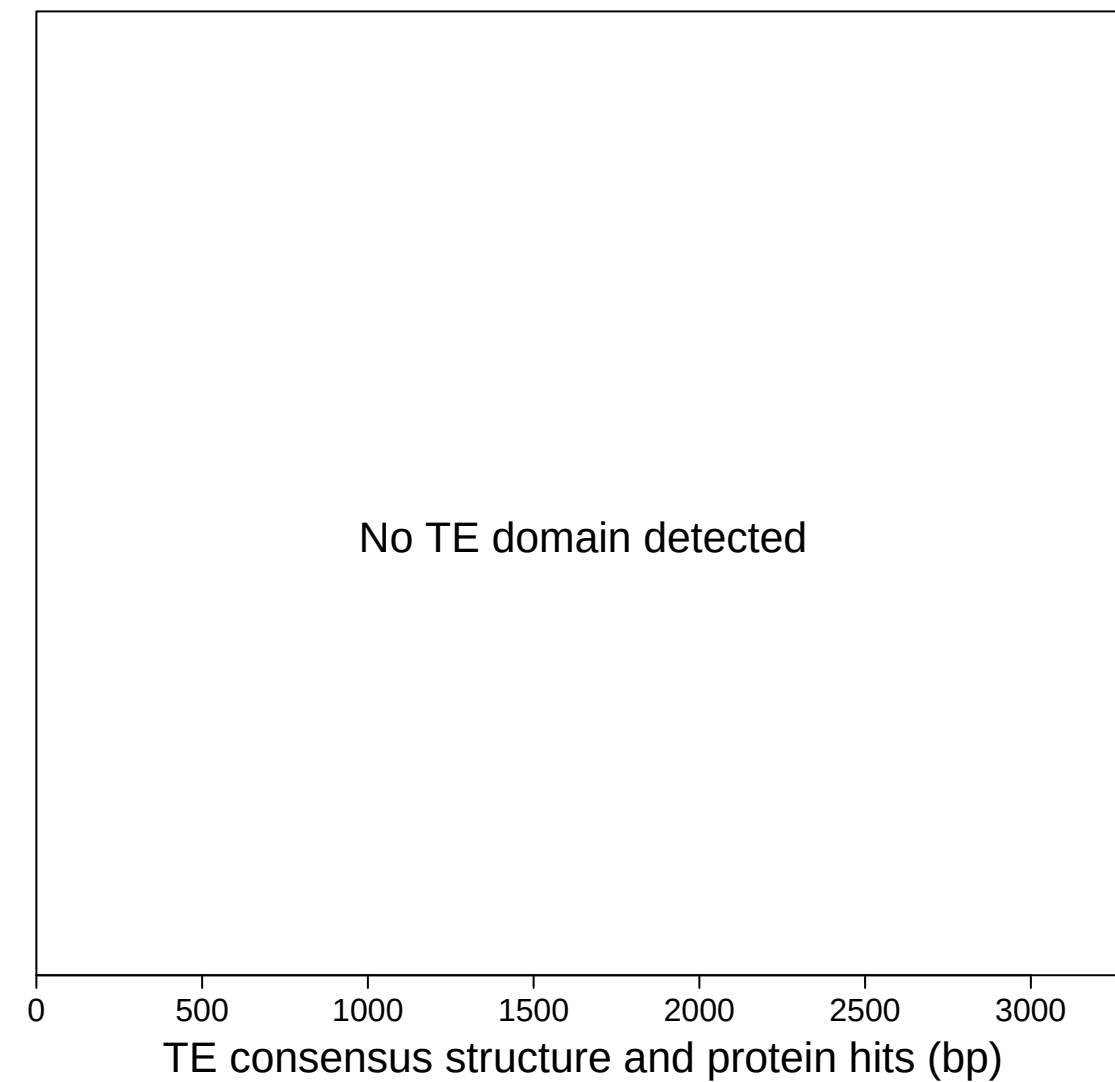
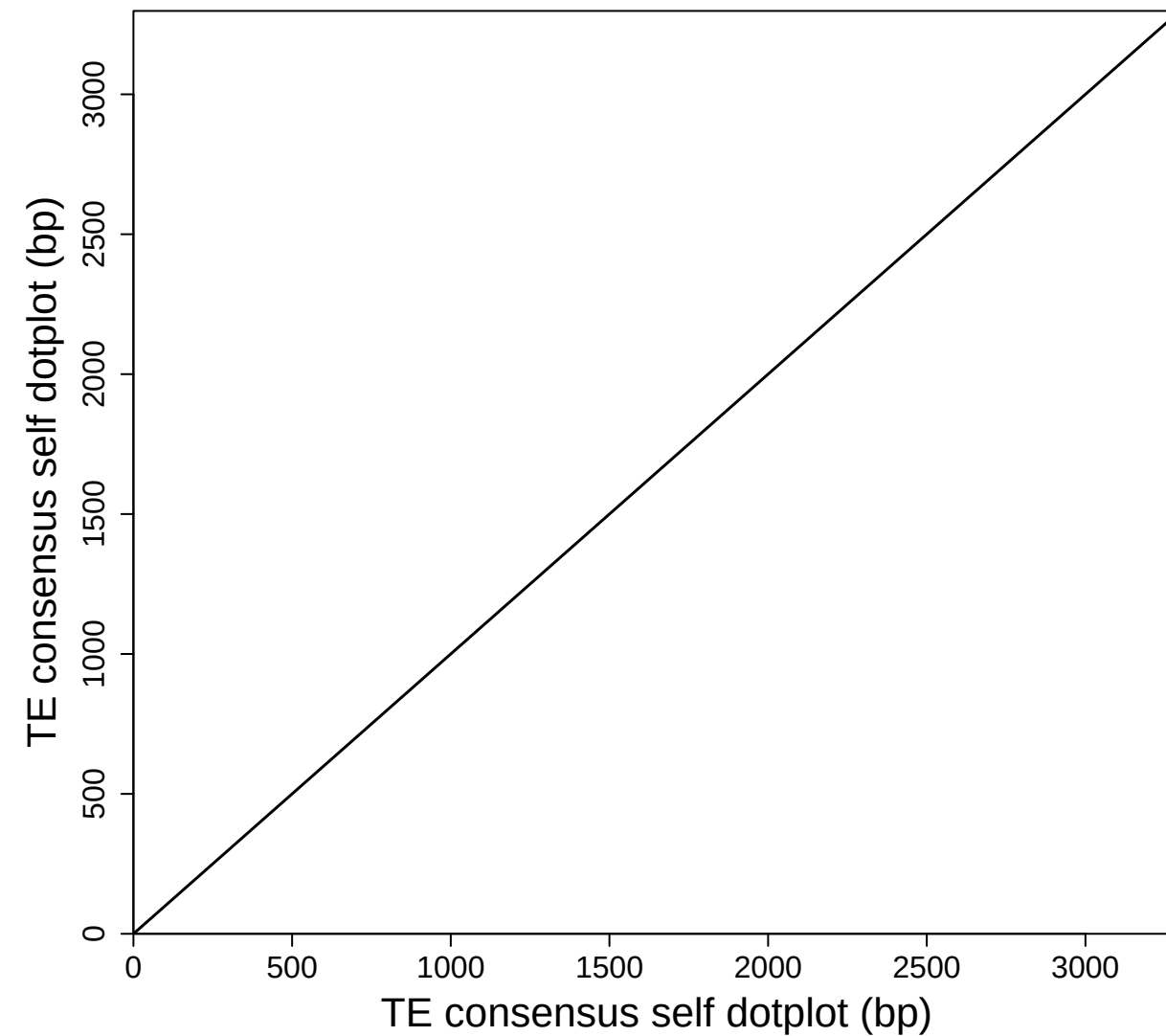
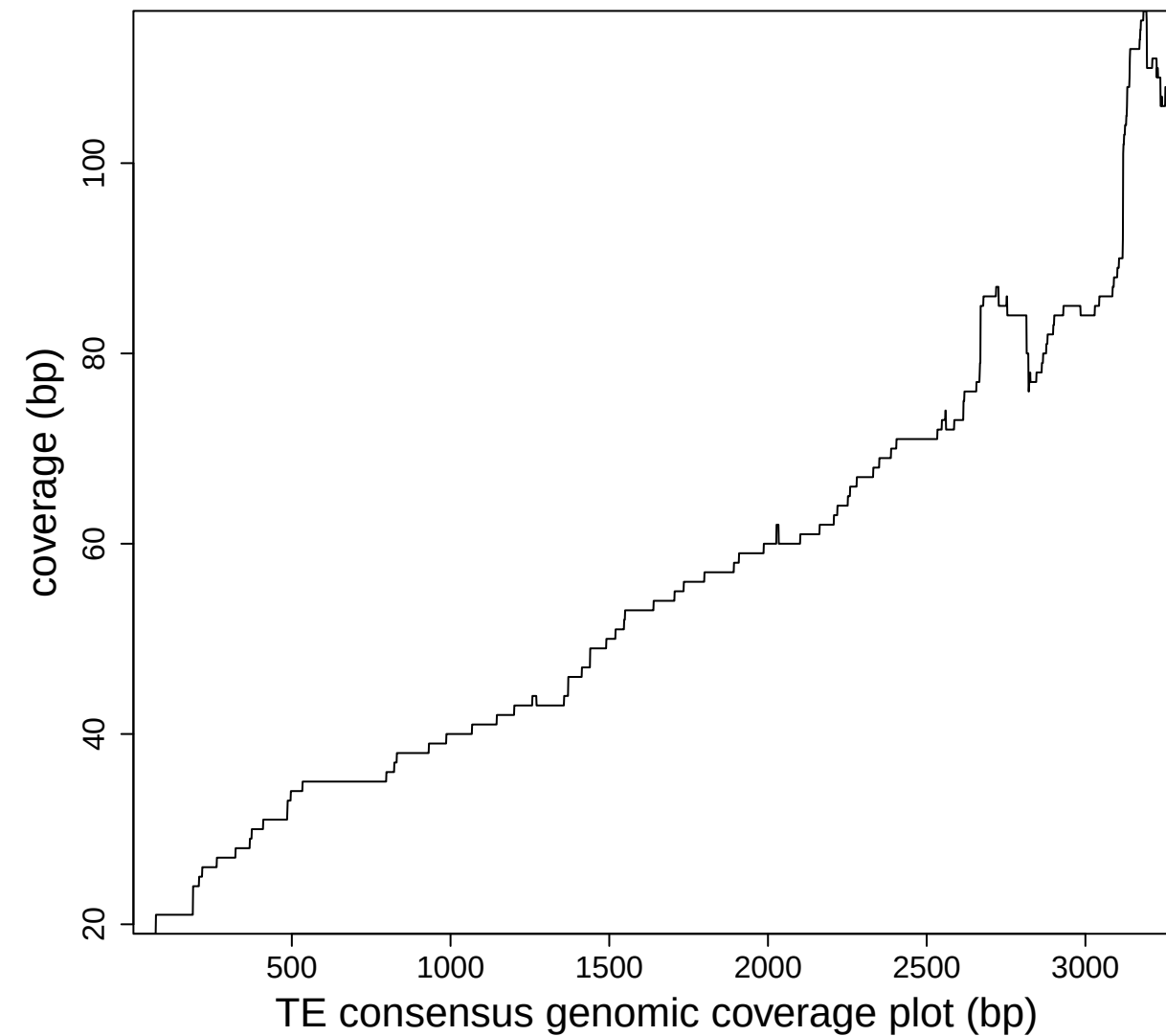
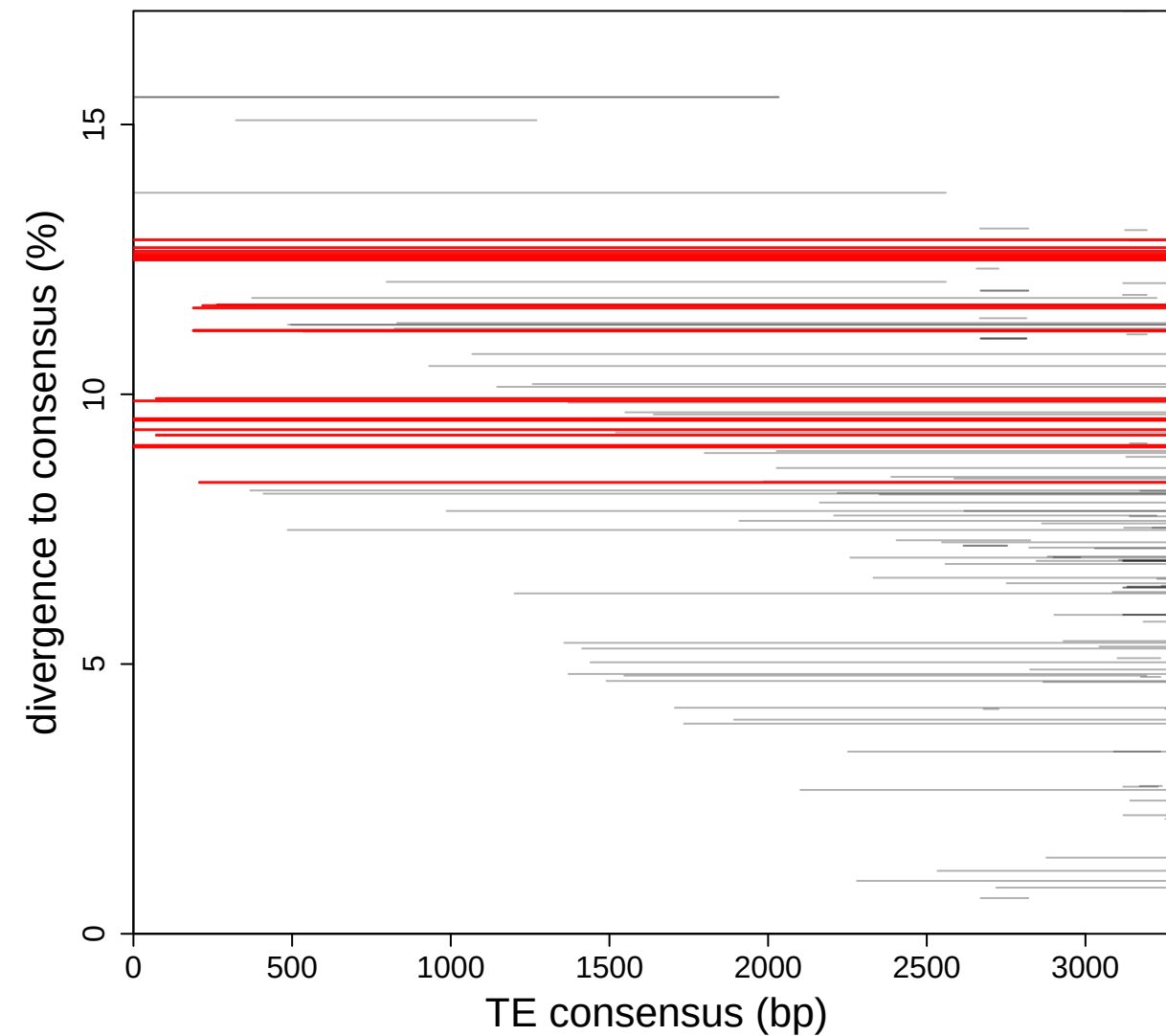
asta.b.bed_uf.bed_g_3.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_c
size: 6296bp; fragments: 620; full length: 0 (>=5666.4bp)

After TEtrimmer Extended plot Blue lines are boundaries

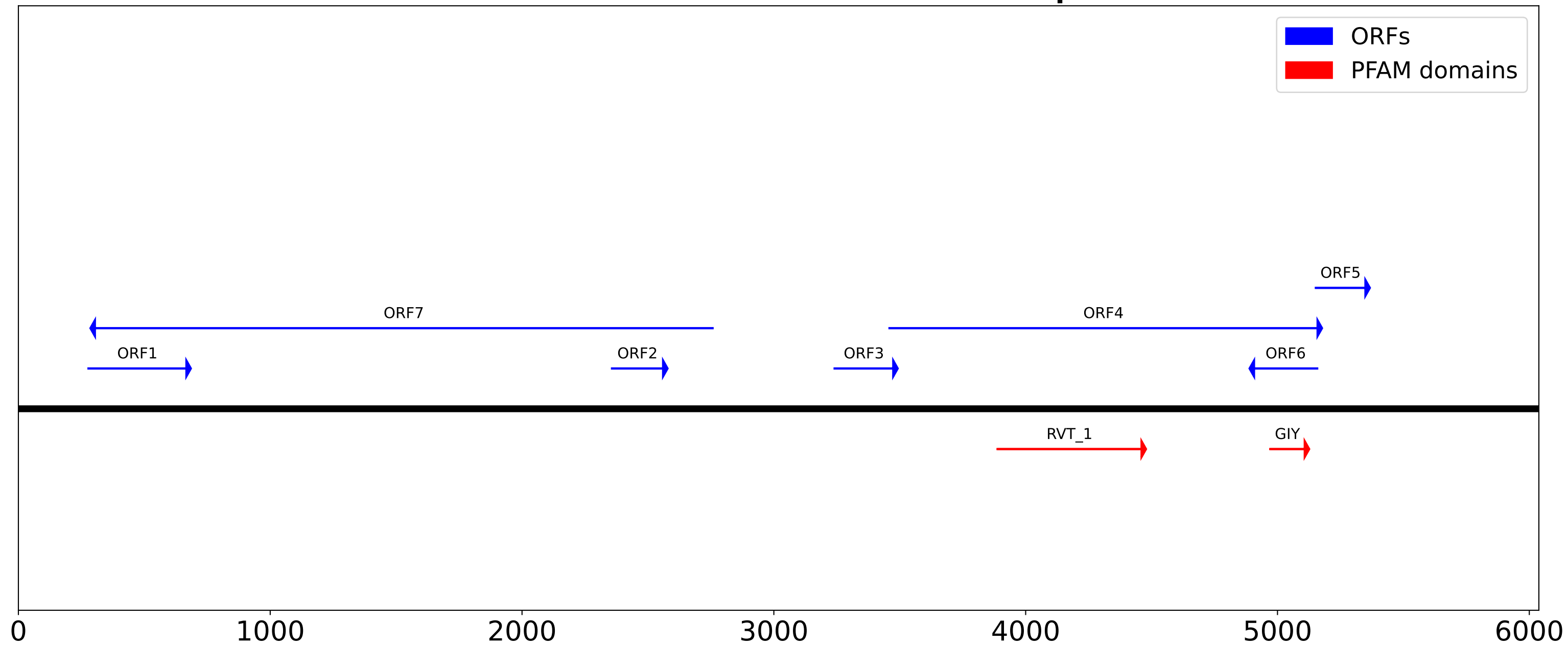


TE: rnd_4_family_4674
size: 3298bp; fragments: 140; full length: 24 (≥ 2968.2 bp)

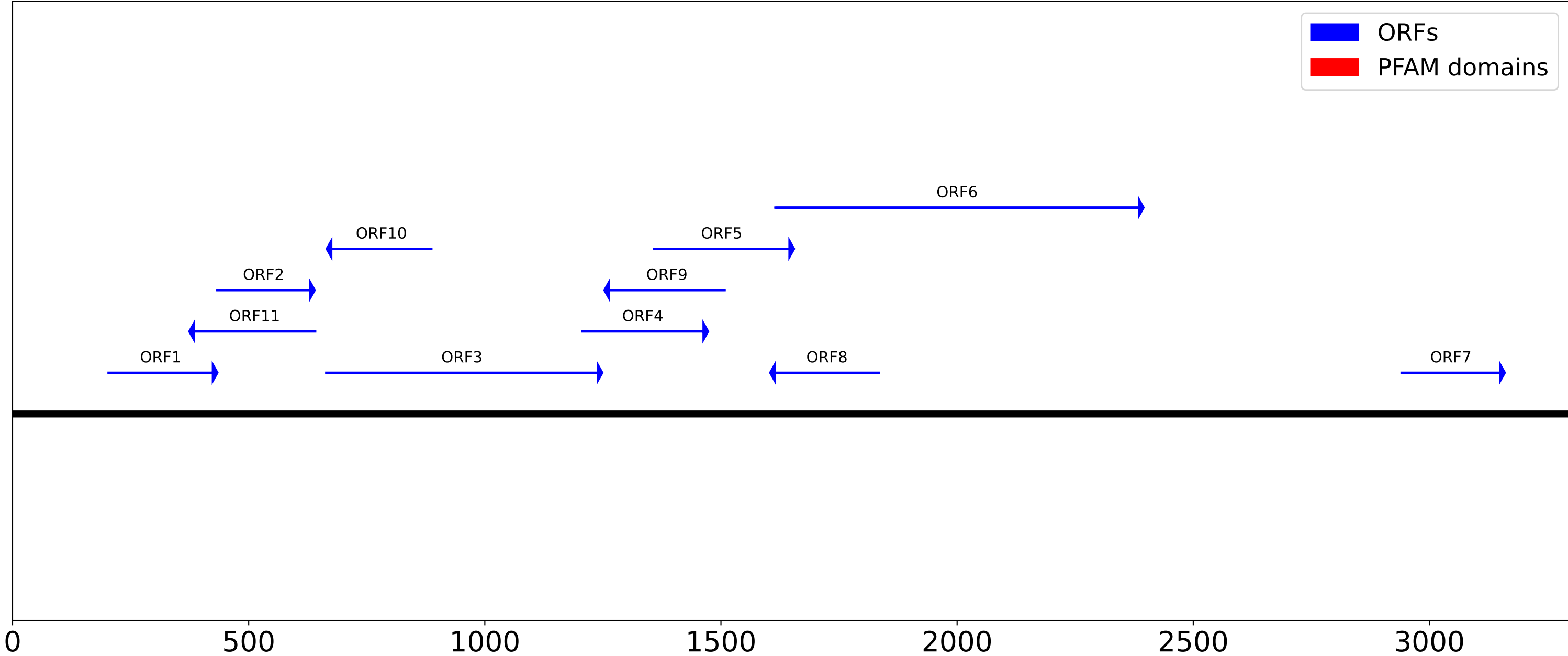
Before TEtrimmer 3298 bp



After TEtrimmer ORF and PFAM domain plot



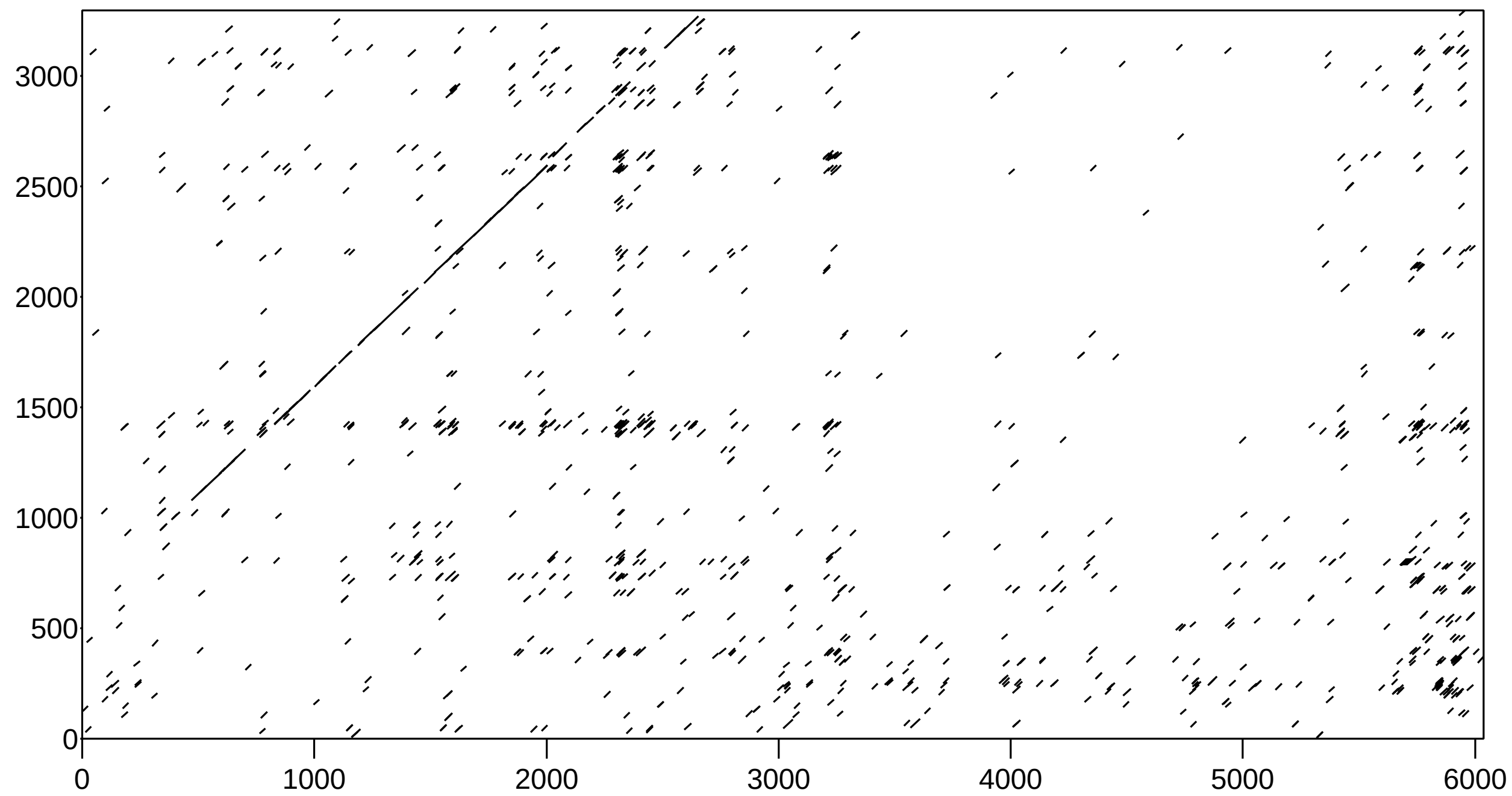
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)