

Start crop Point



End crop Point

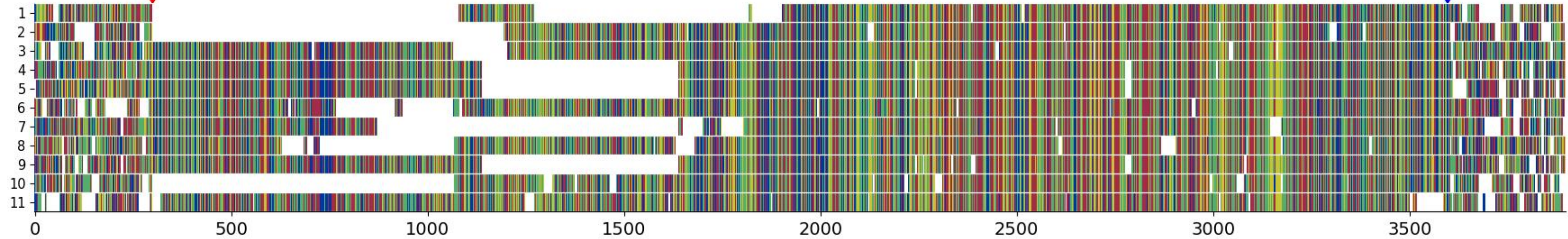


MSA length = 3298



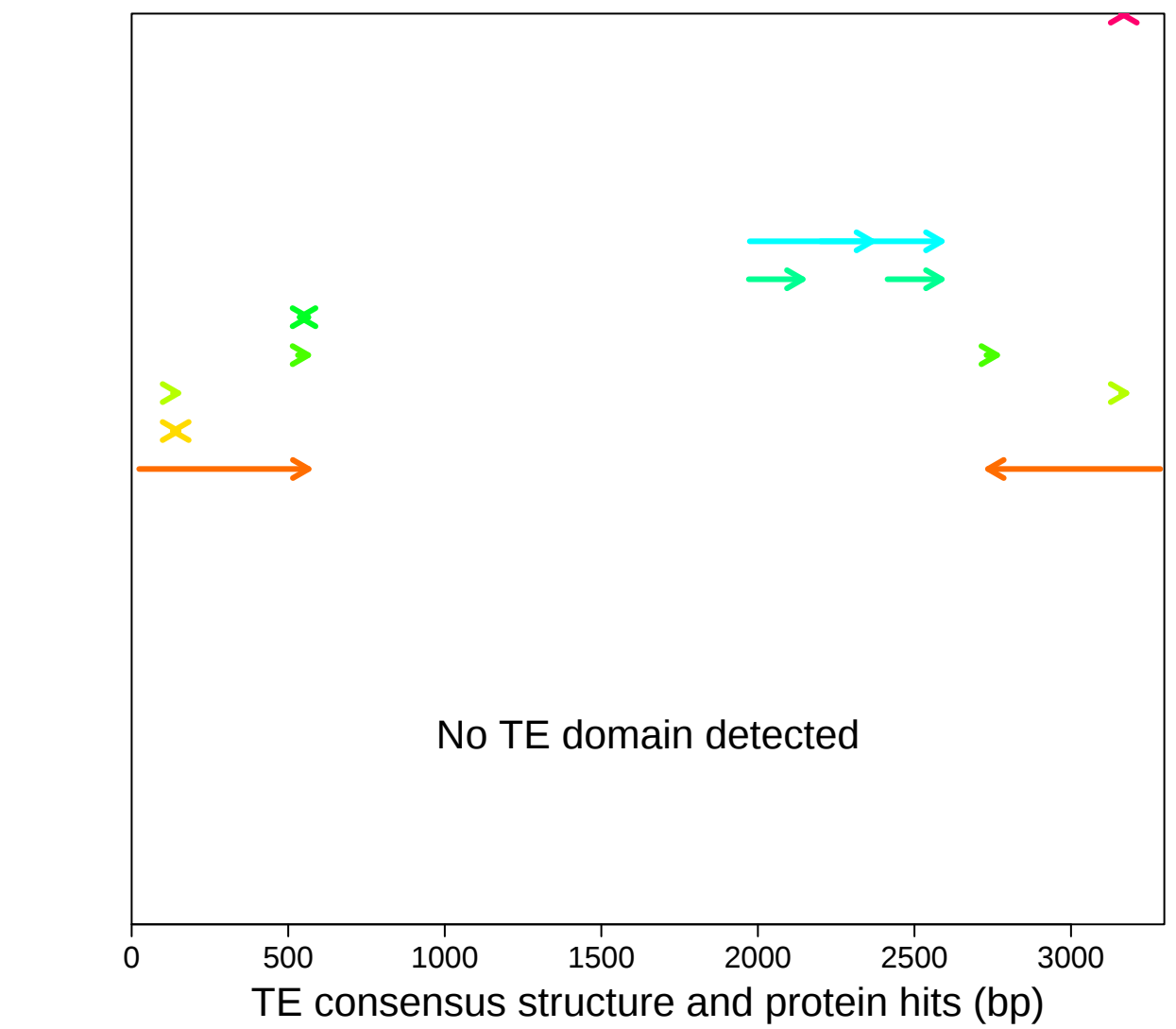
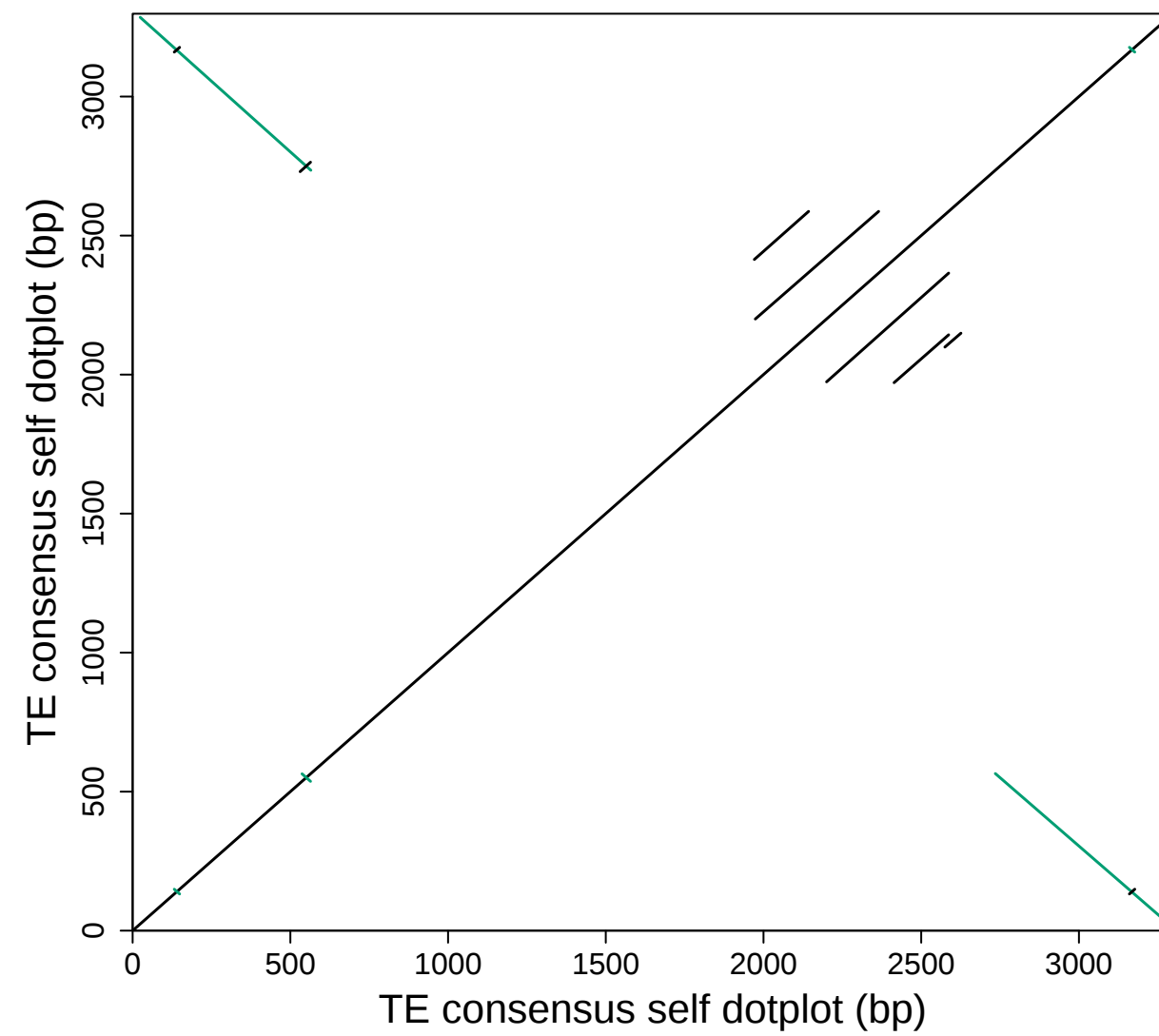
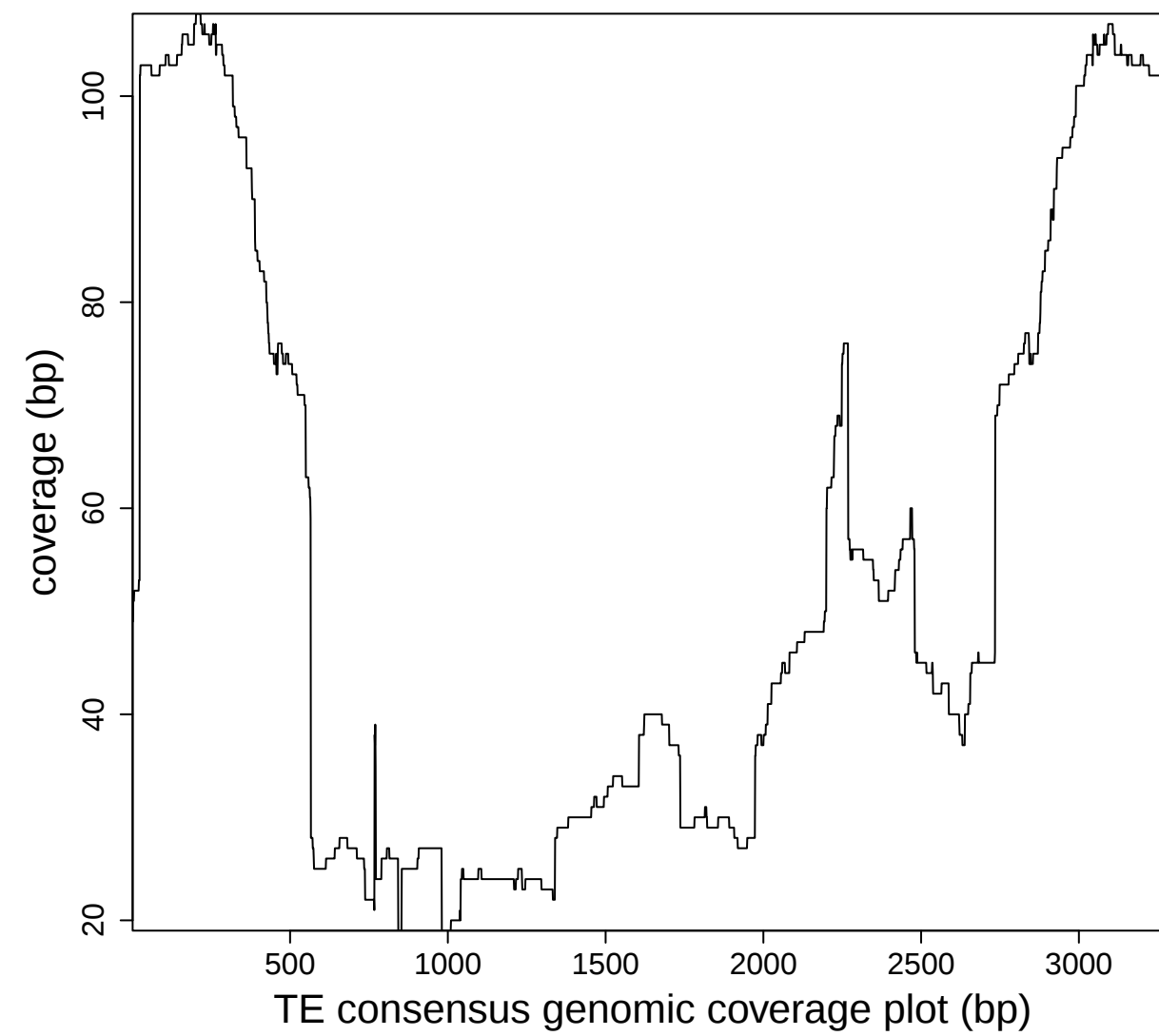
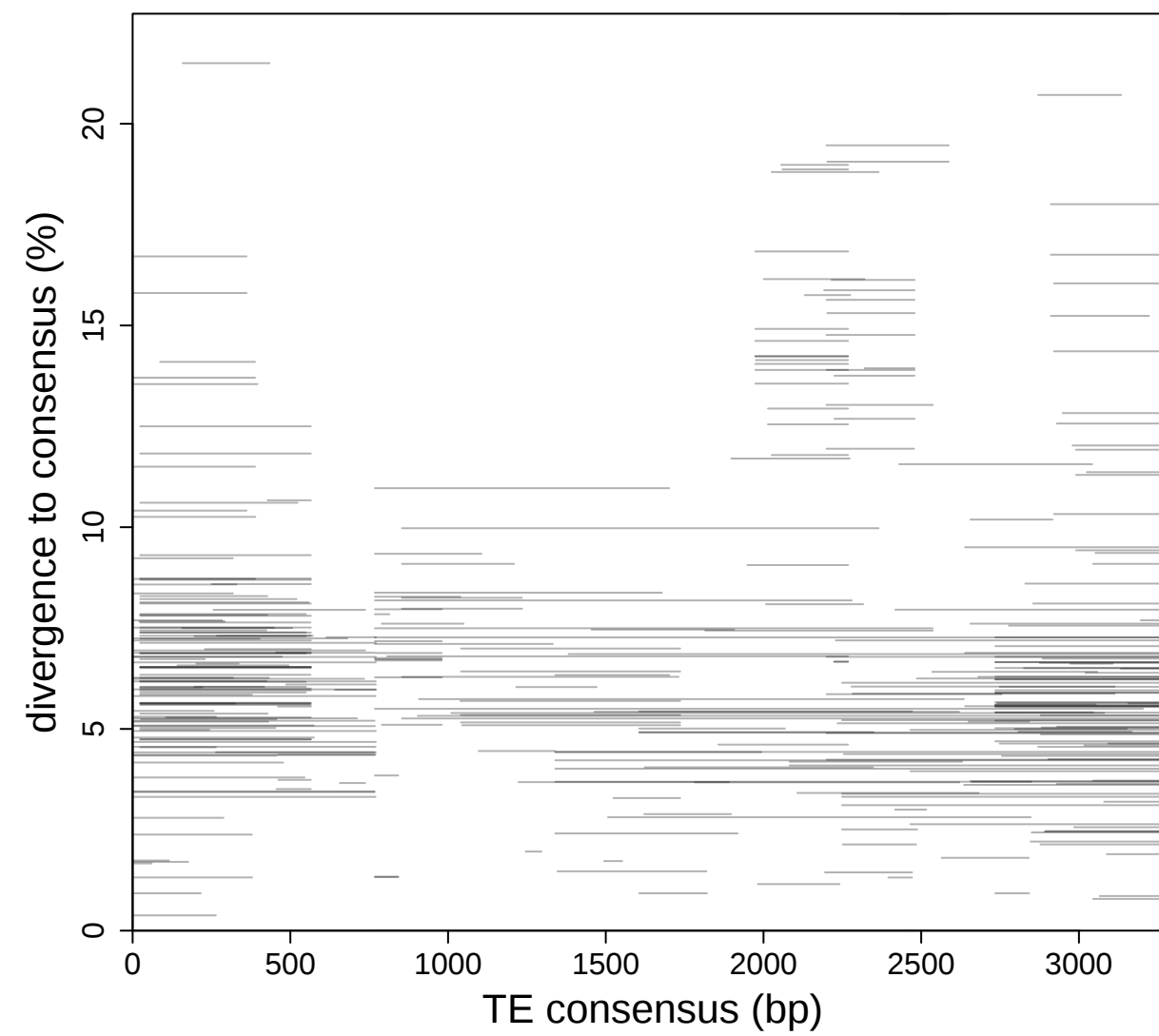
Start crop Point

End crop Point



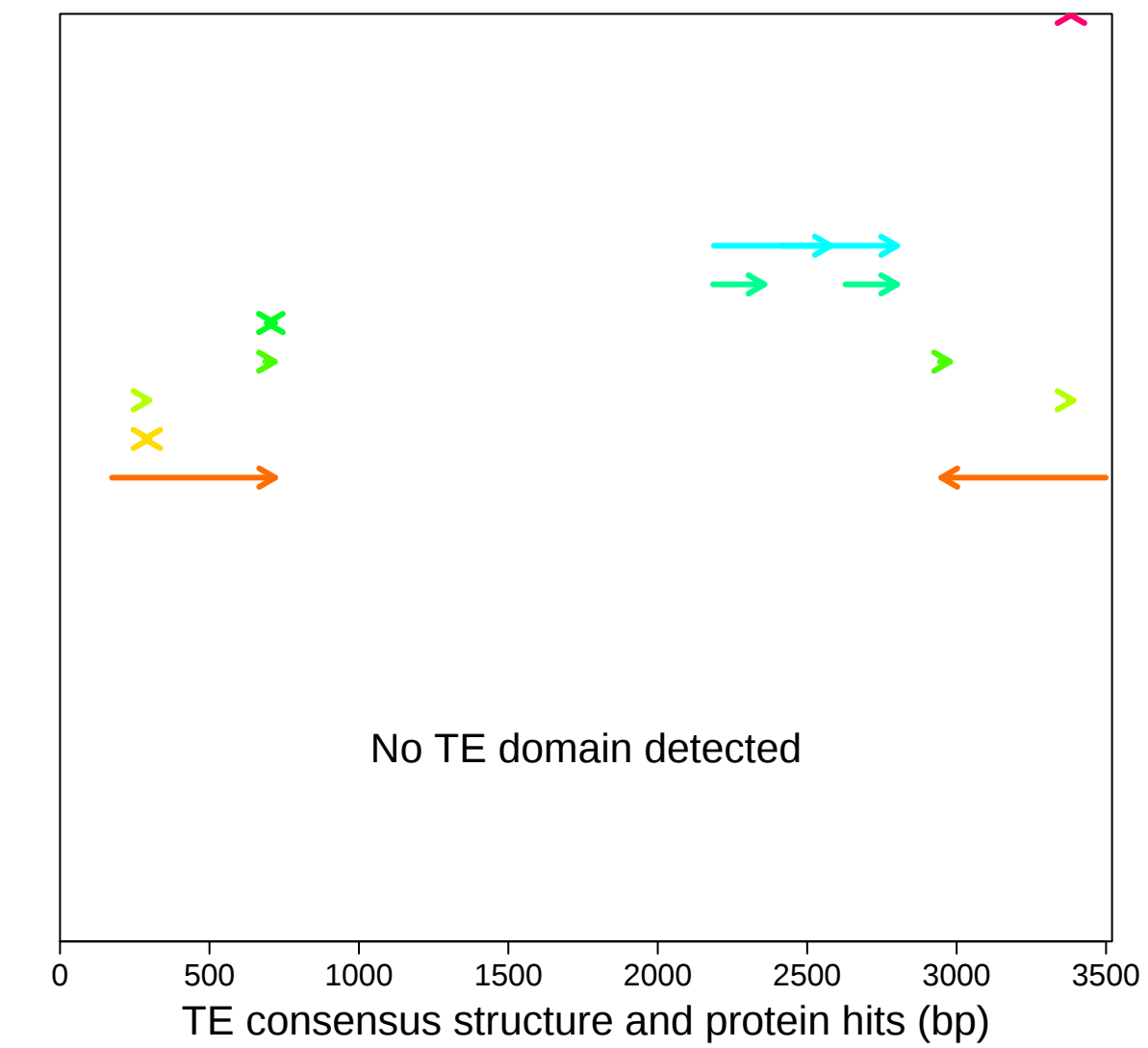
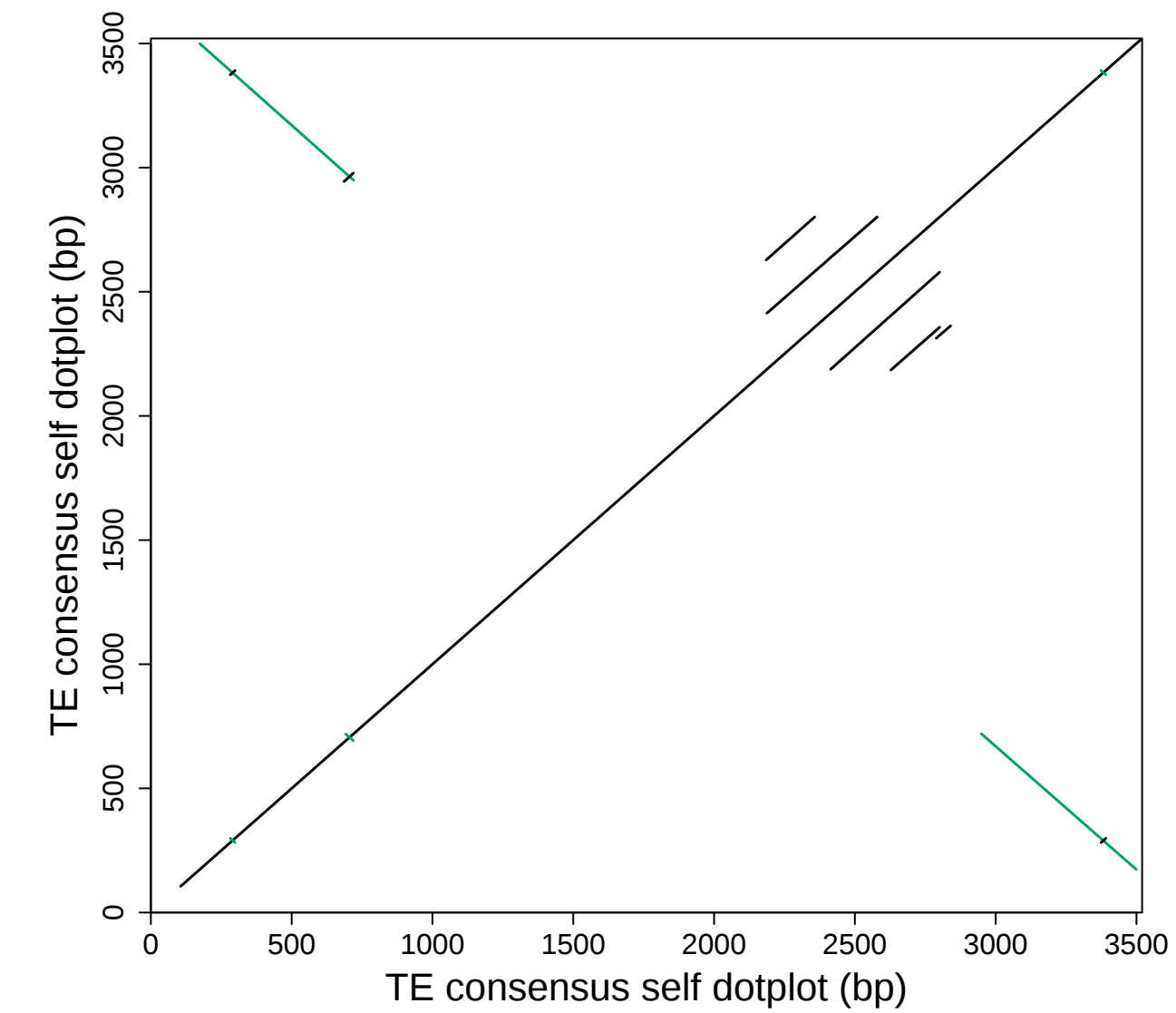
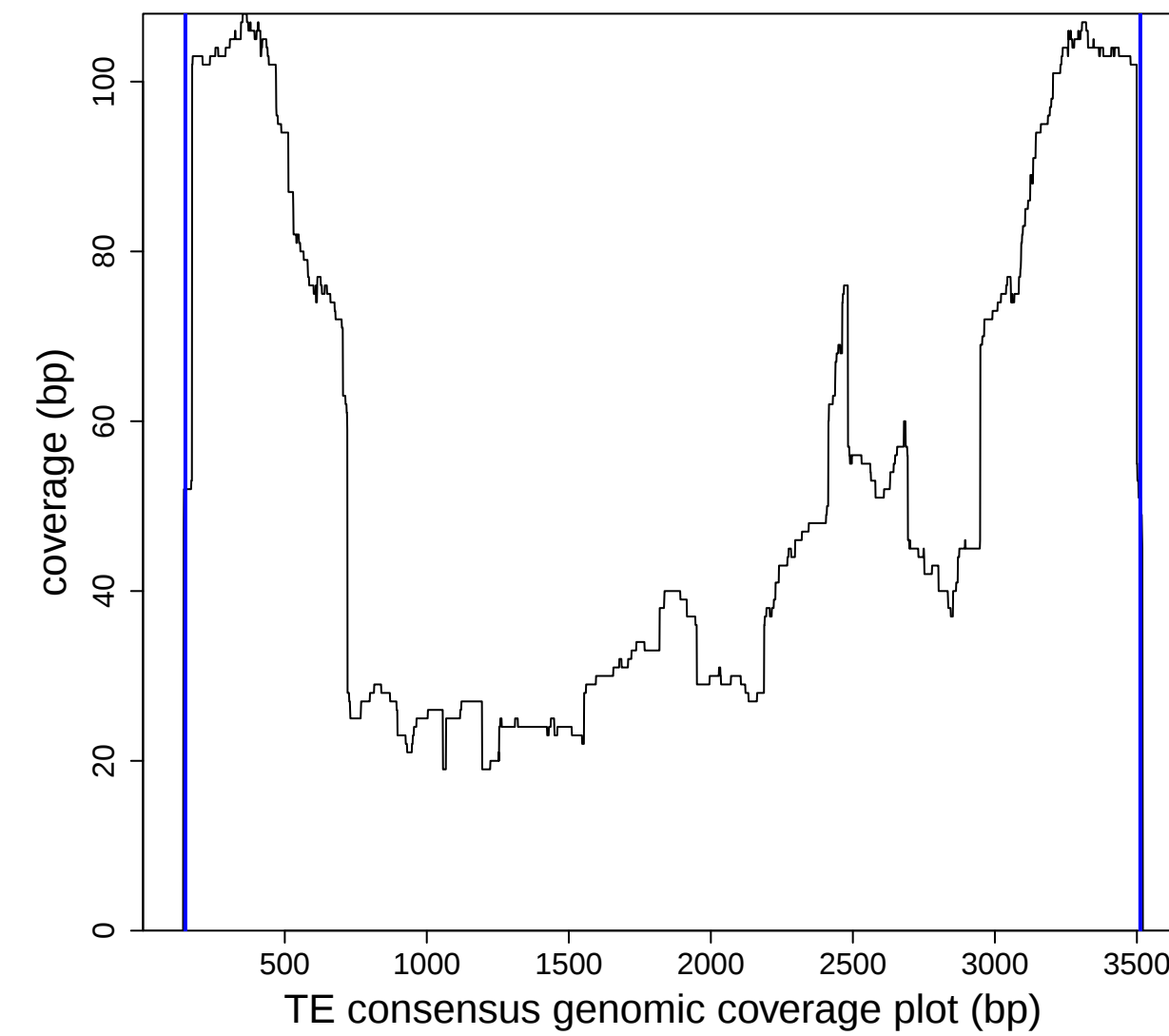
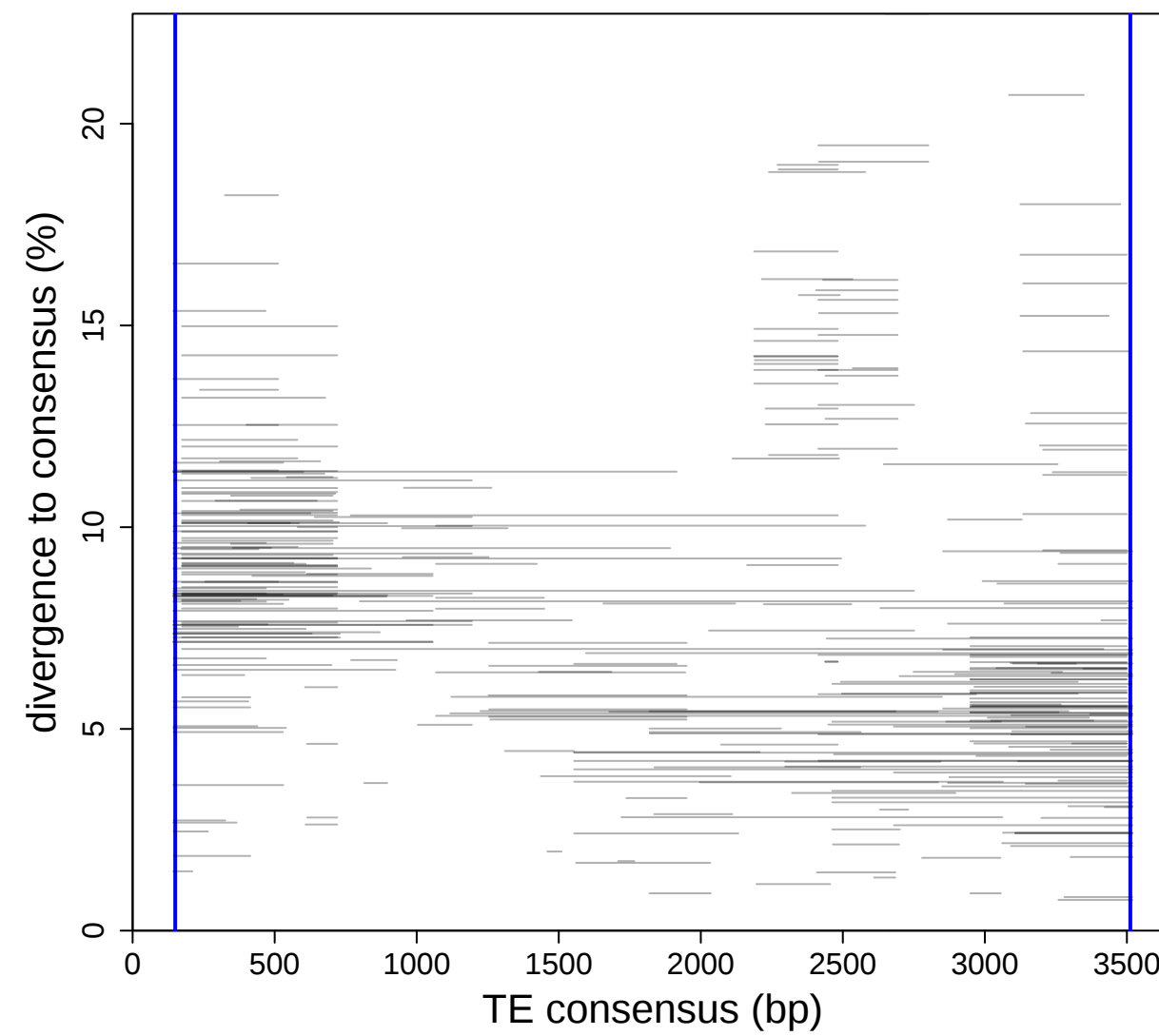
b.bed_uf.bed_g_2.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.fa
size: 3298bp; fragments: 351; full length: 0 (>=2968.2bp)

After TEtrimmer 3298 bp



2.fasta.b.bed_uf.bed_g_2.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa
size: 3662bp; fragments: 332; full length: 0 (>=3295.8bp)

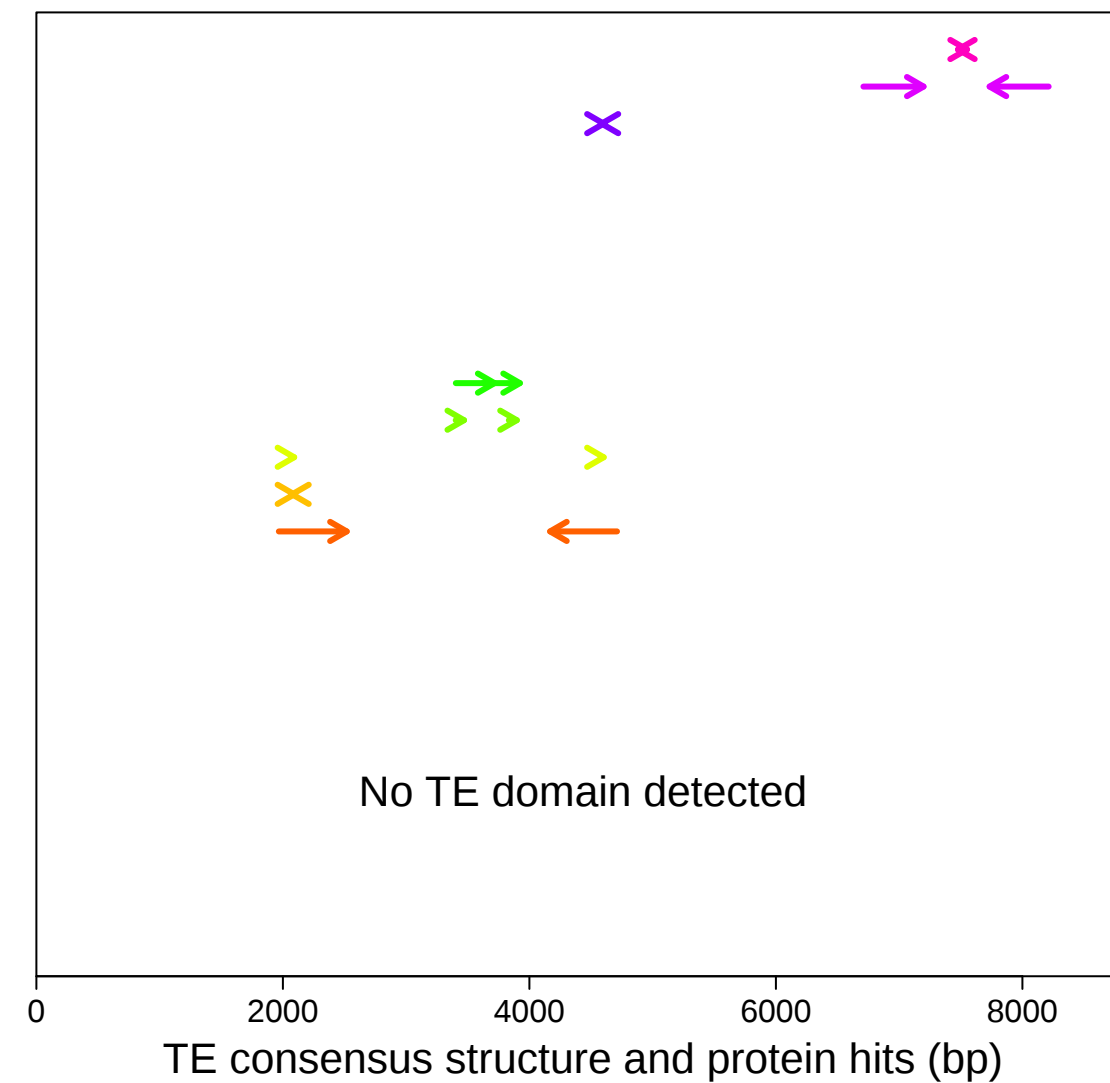
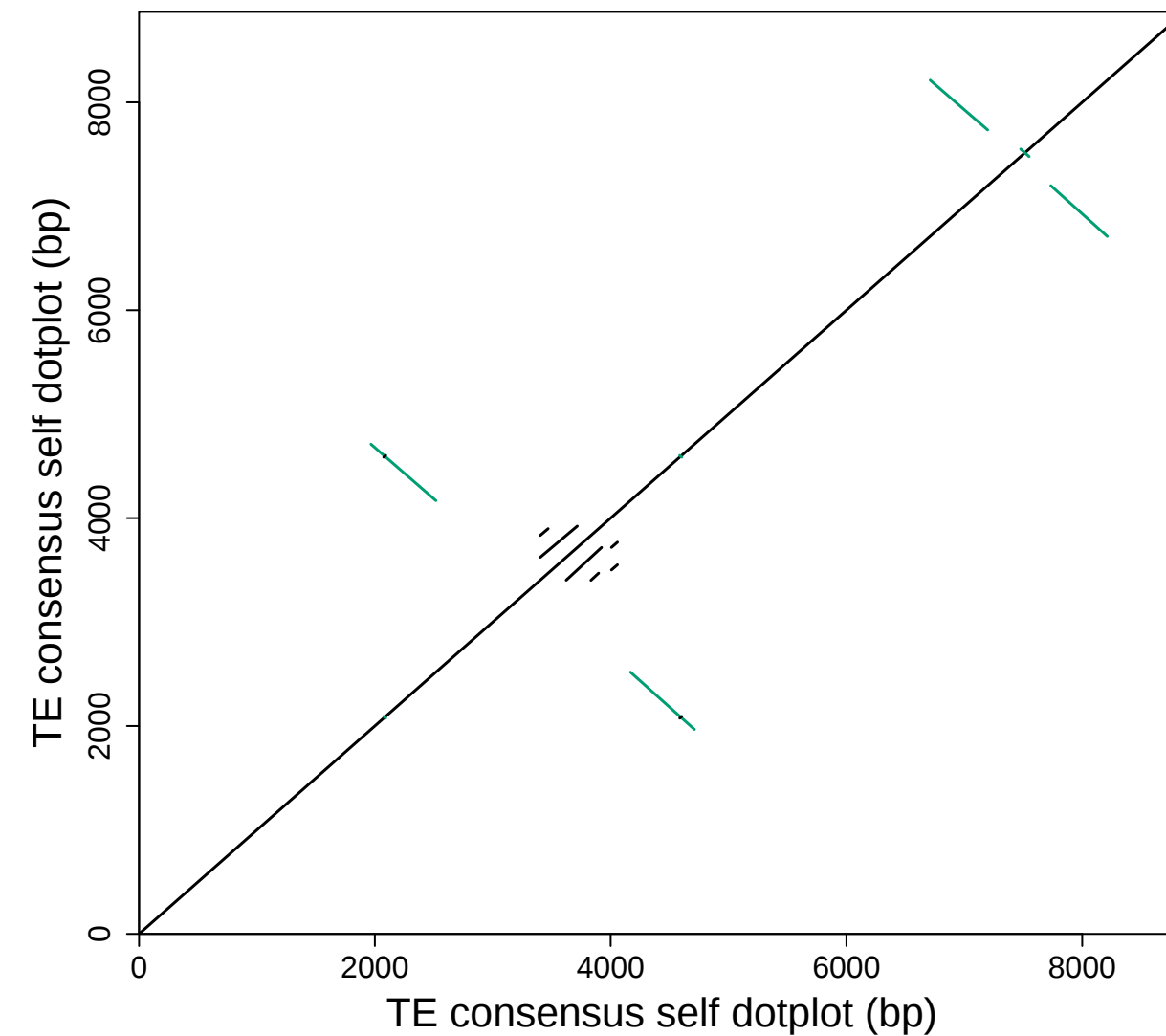
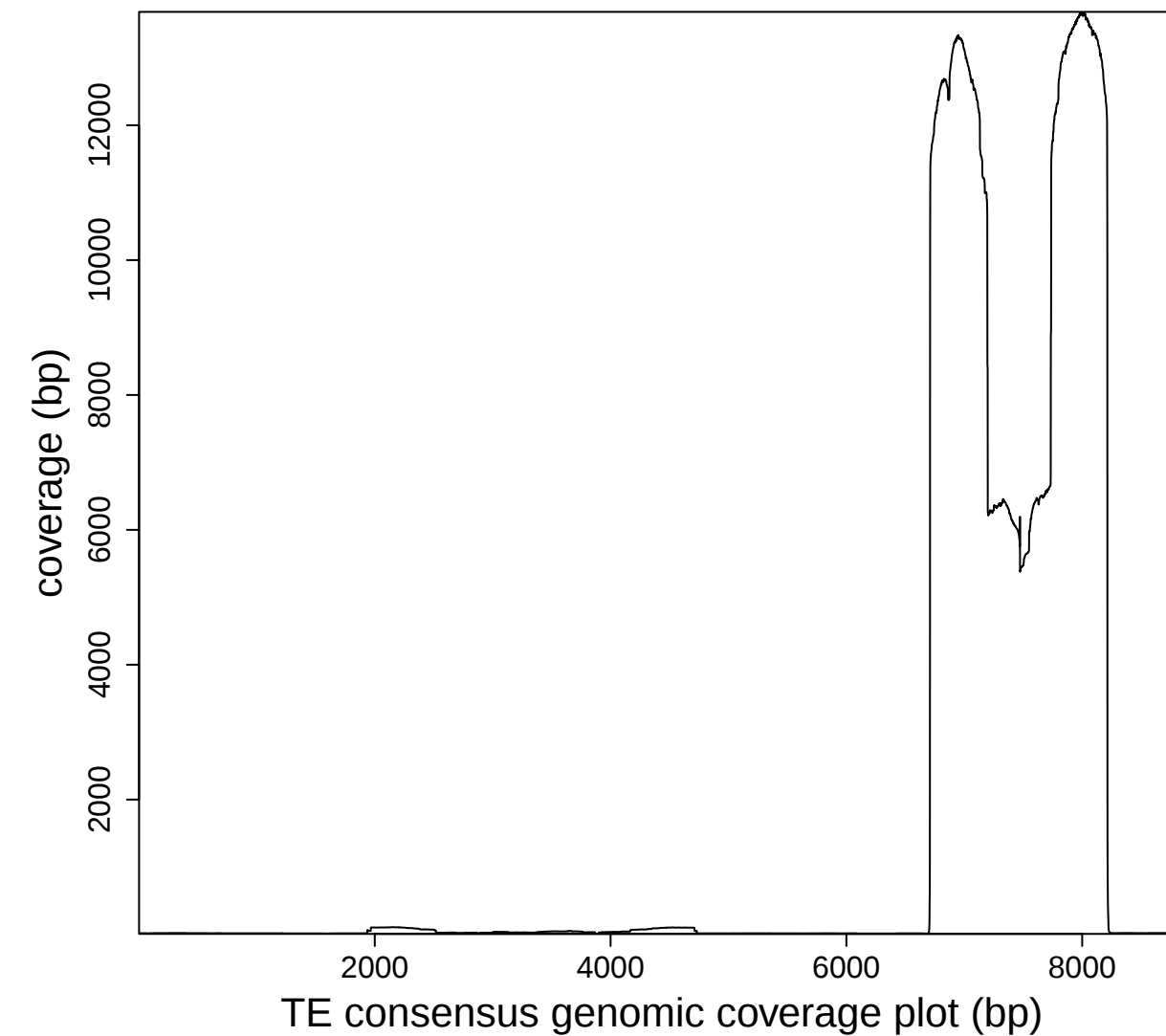
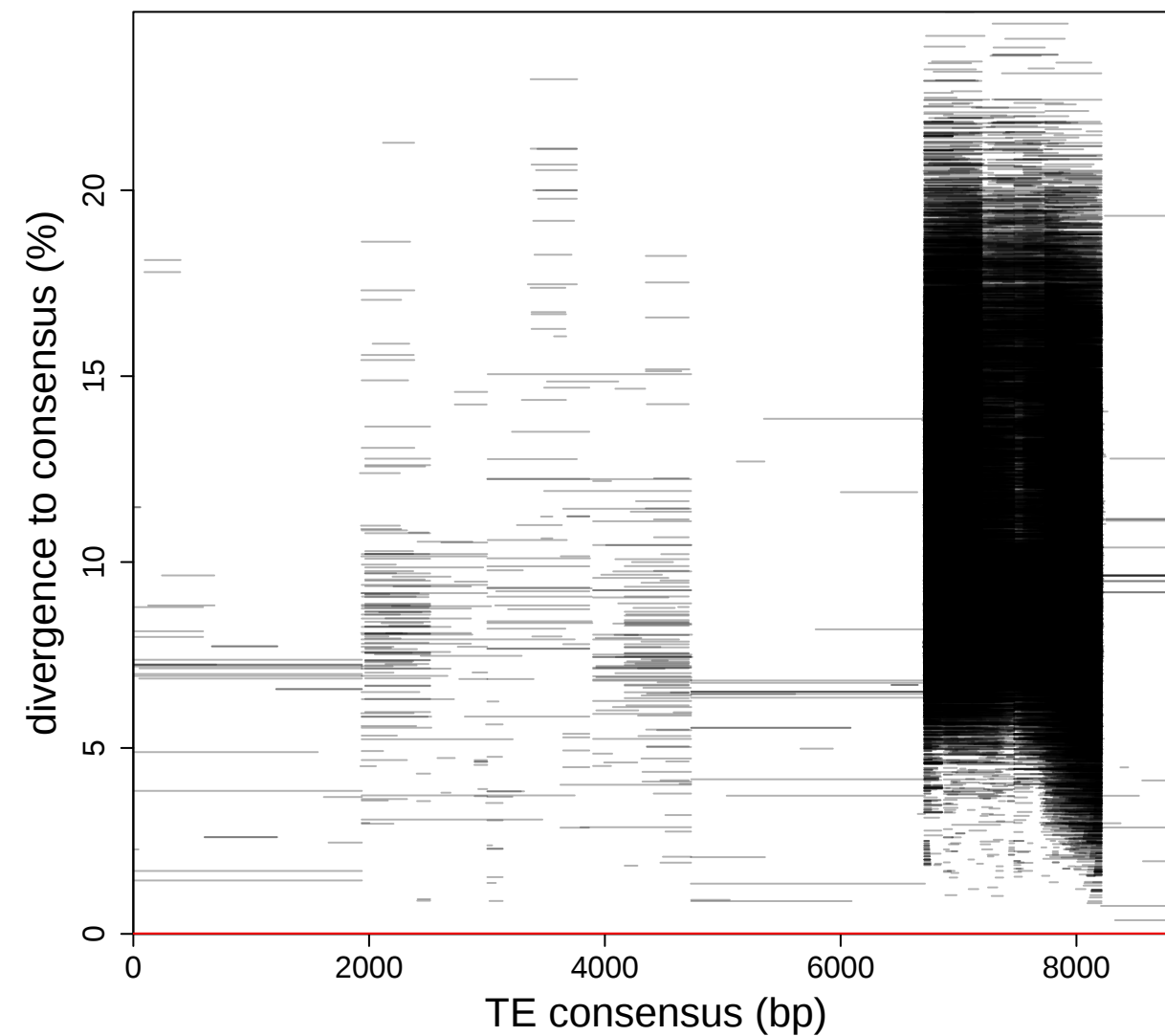
After TEtrimmer Extended plot Blue lines are boundaries



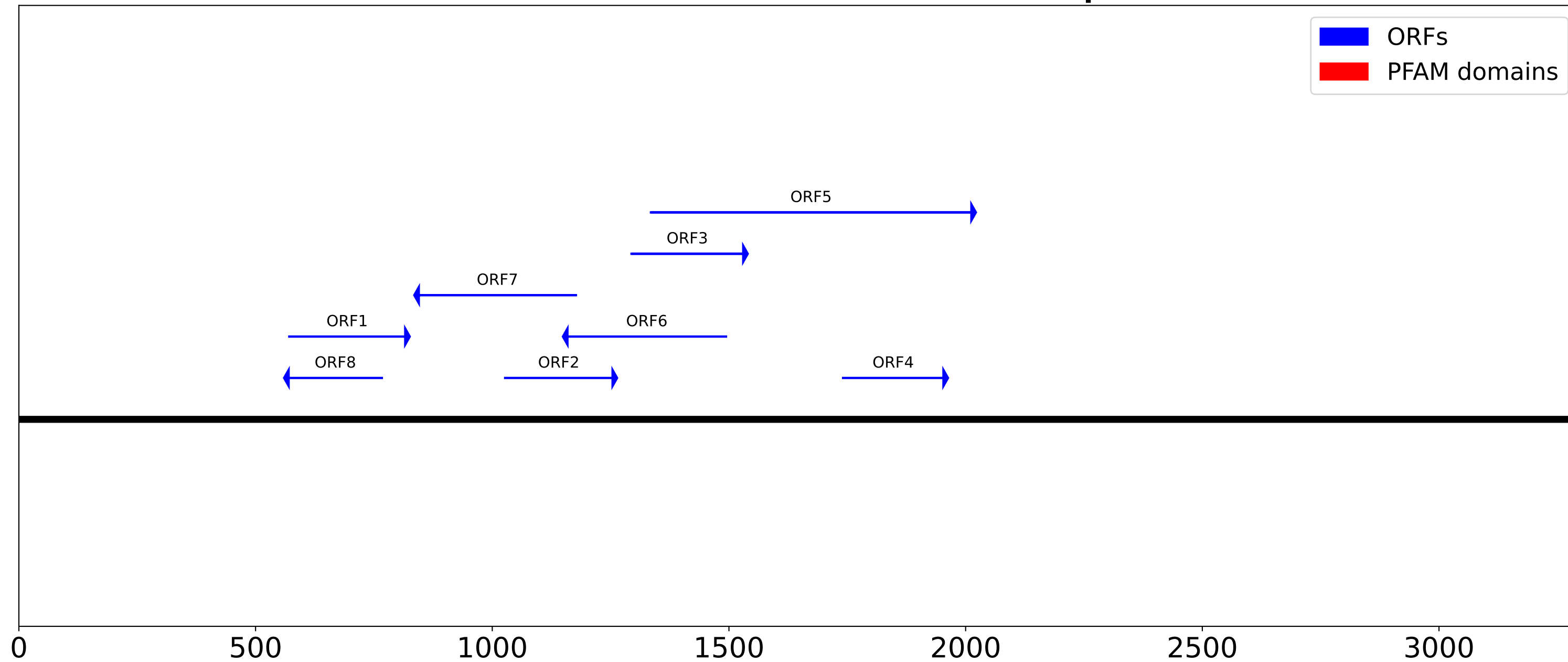
Before TEtrimmer 8870 bp

TE: ltr_1_family_622

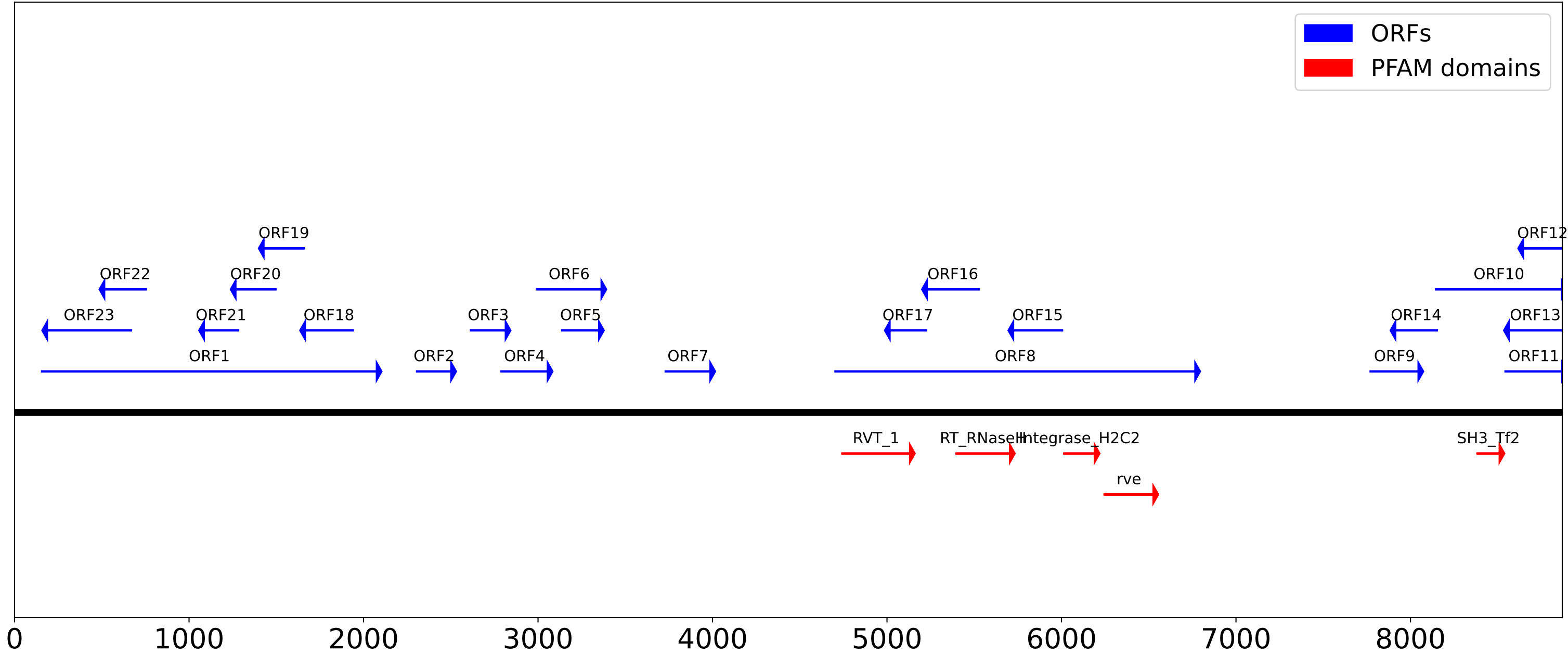
size: 8870bp; fragments: 29341; full length: 1 (>=7983bp)



After TEtrimmer ORF and PFAM domain plot

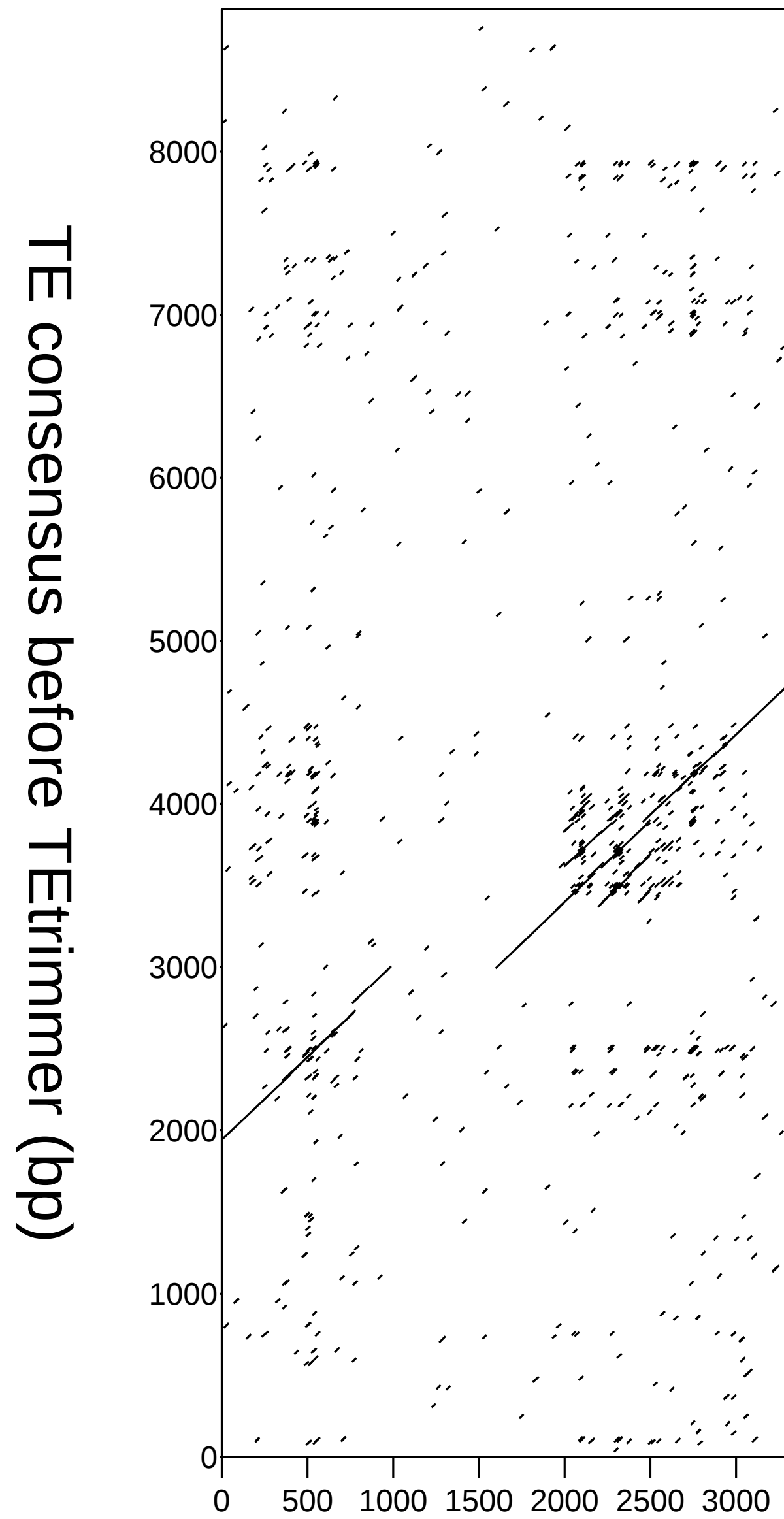


Before TEtrimmer ORF and PFAM domain plot



Dotplot

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



TE consensus after TETrimmer (bp)