

Start crop Point

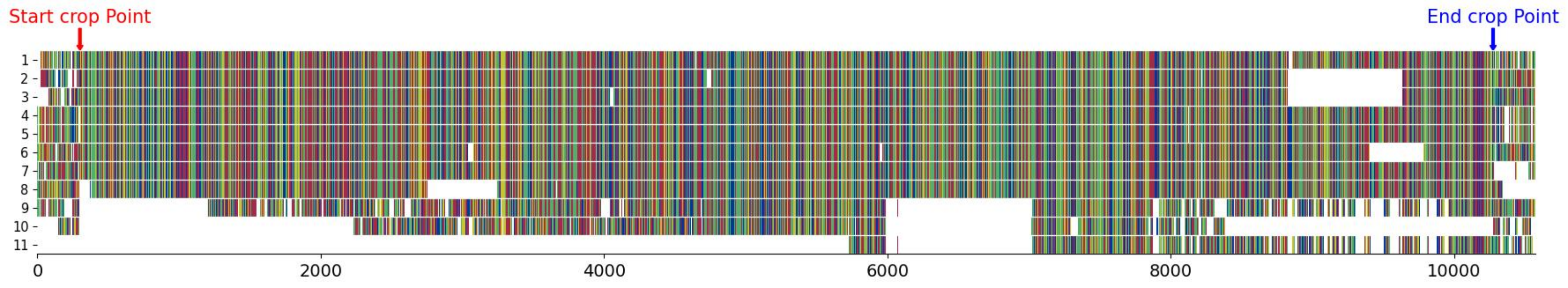


End crop Point



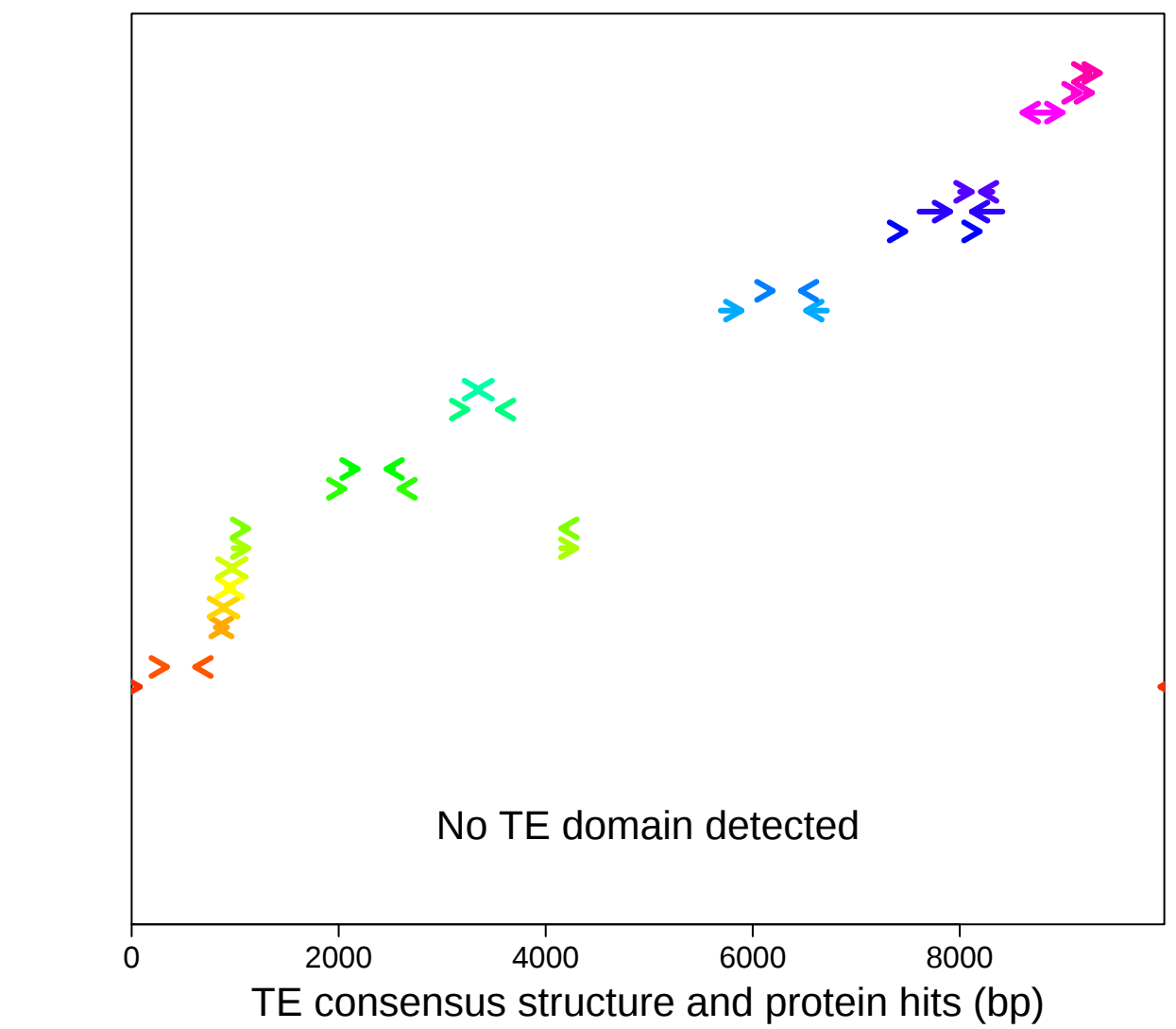
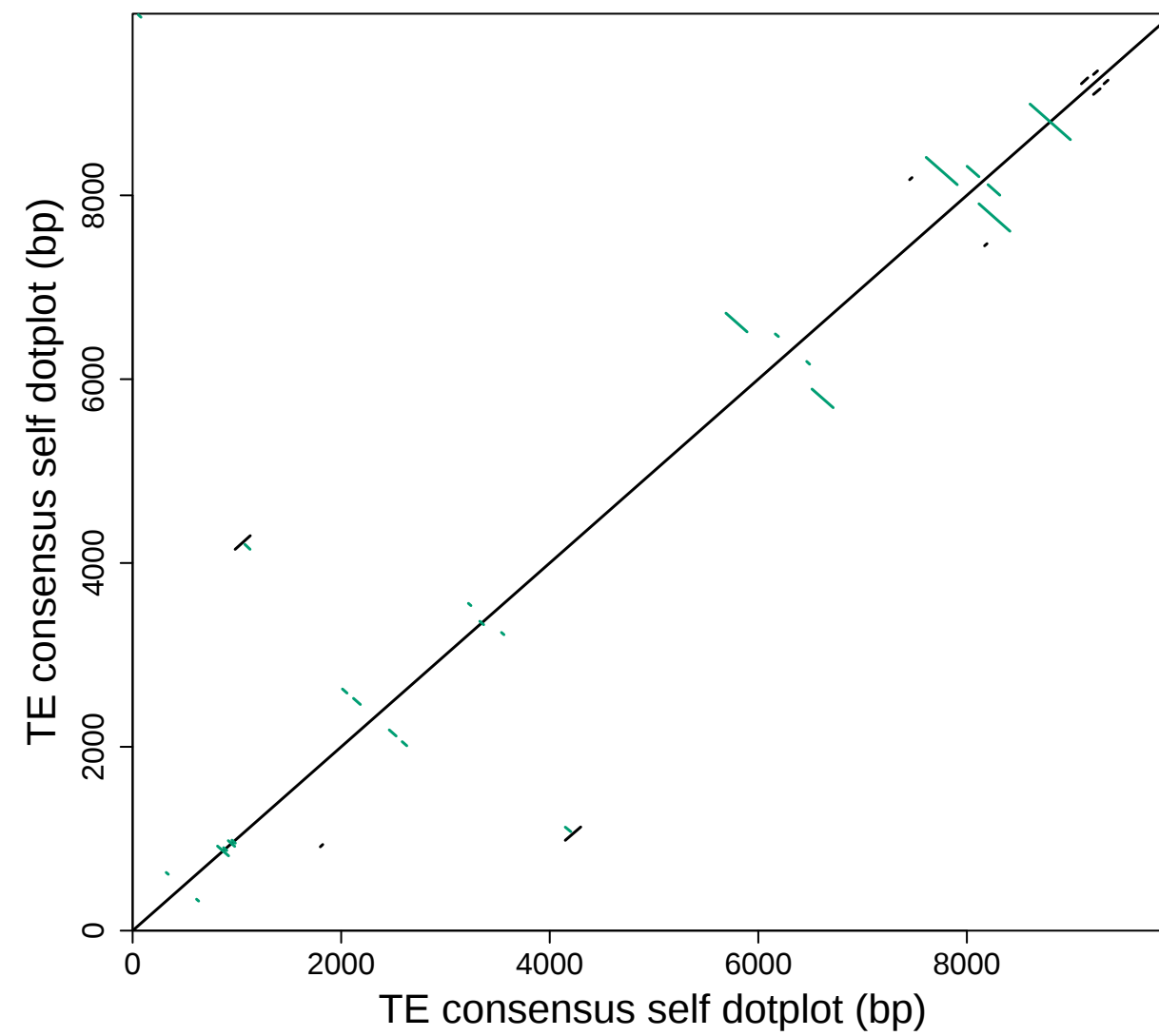
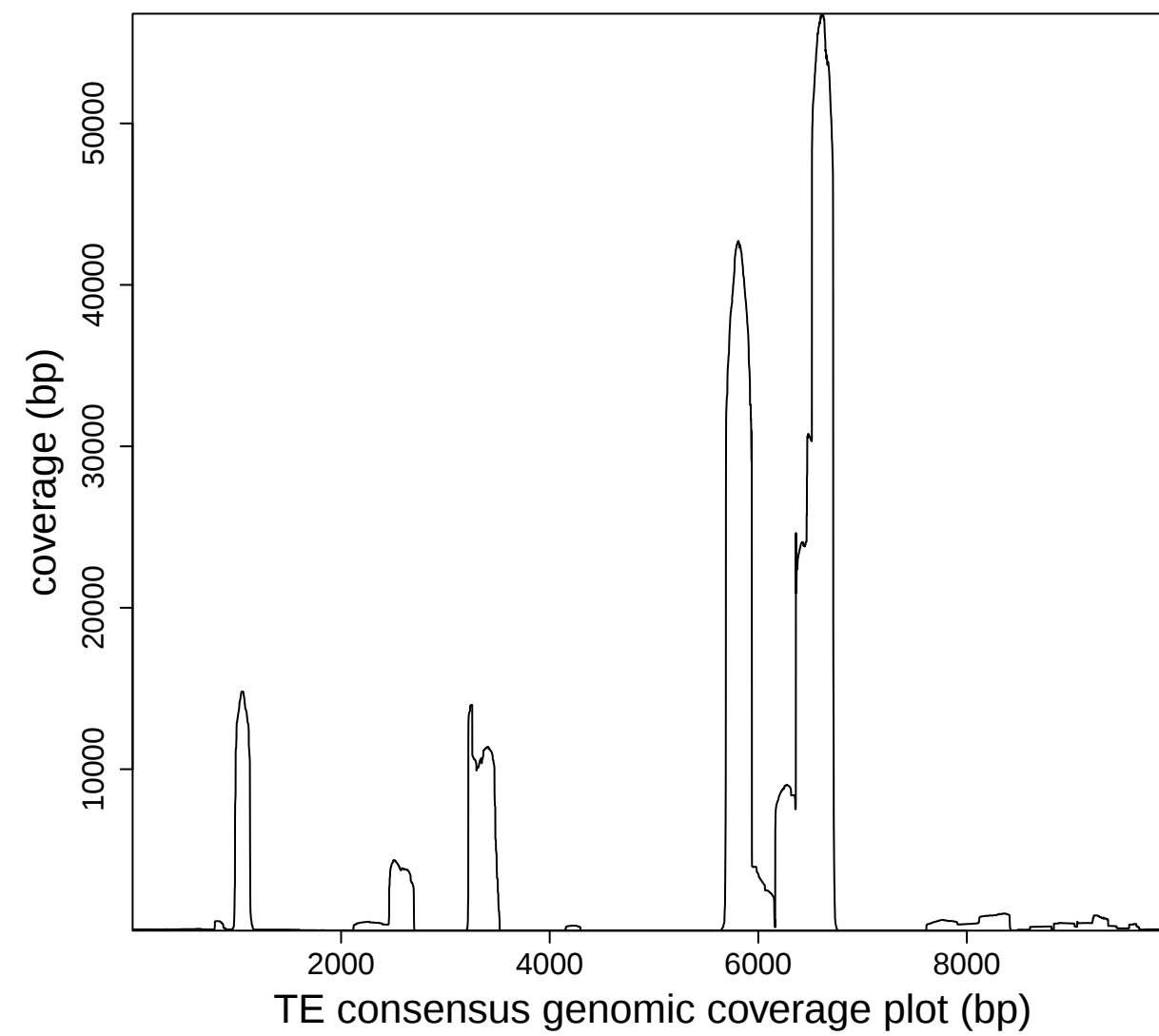
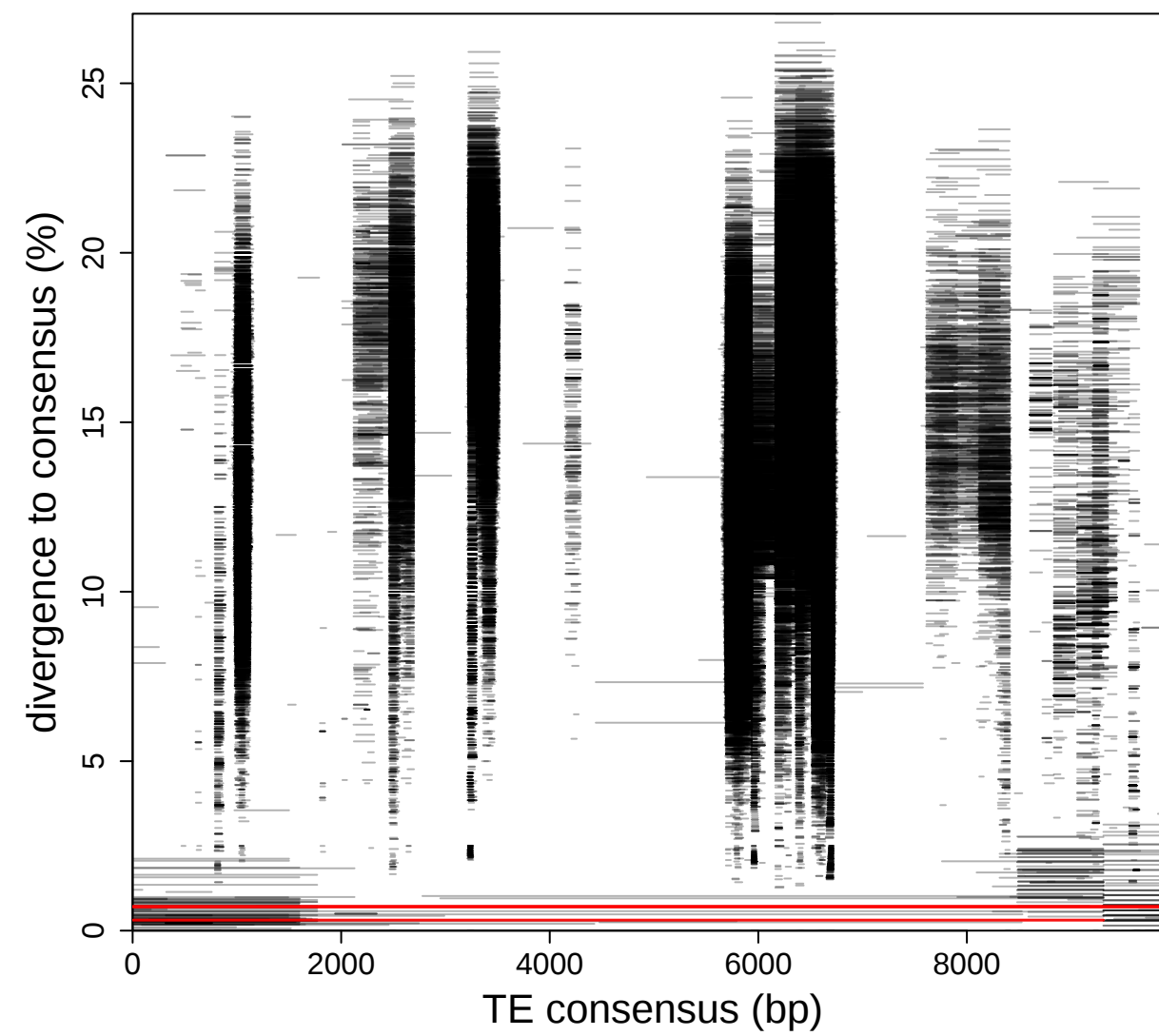
MSA length = 9976





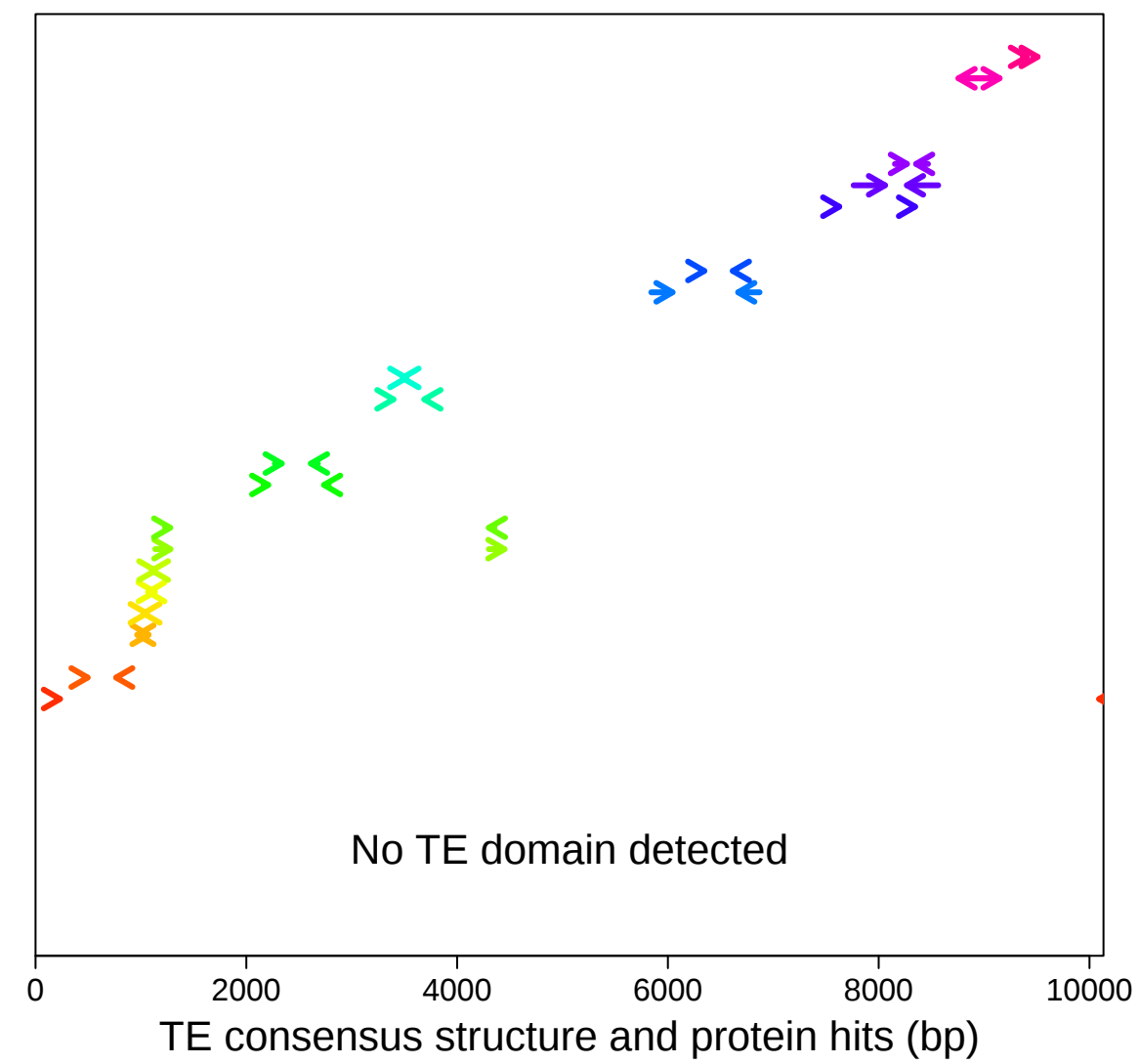
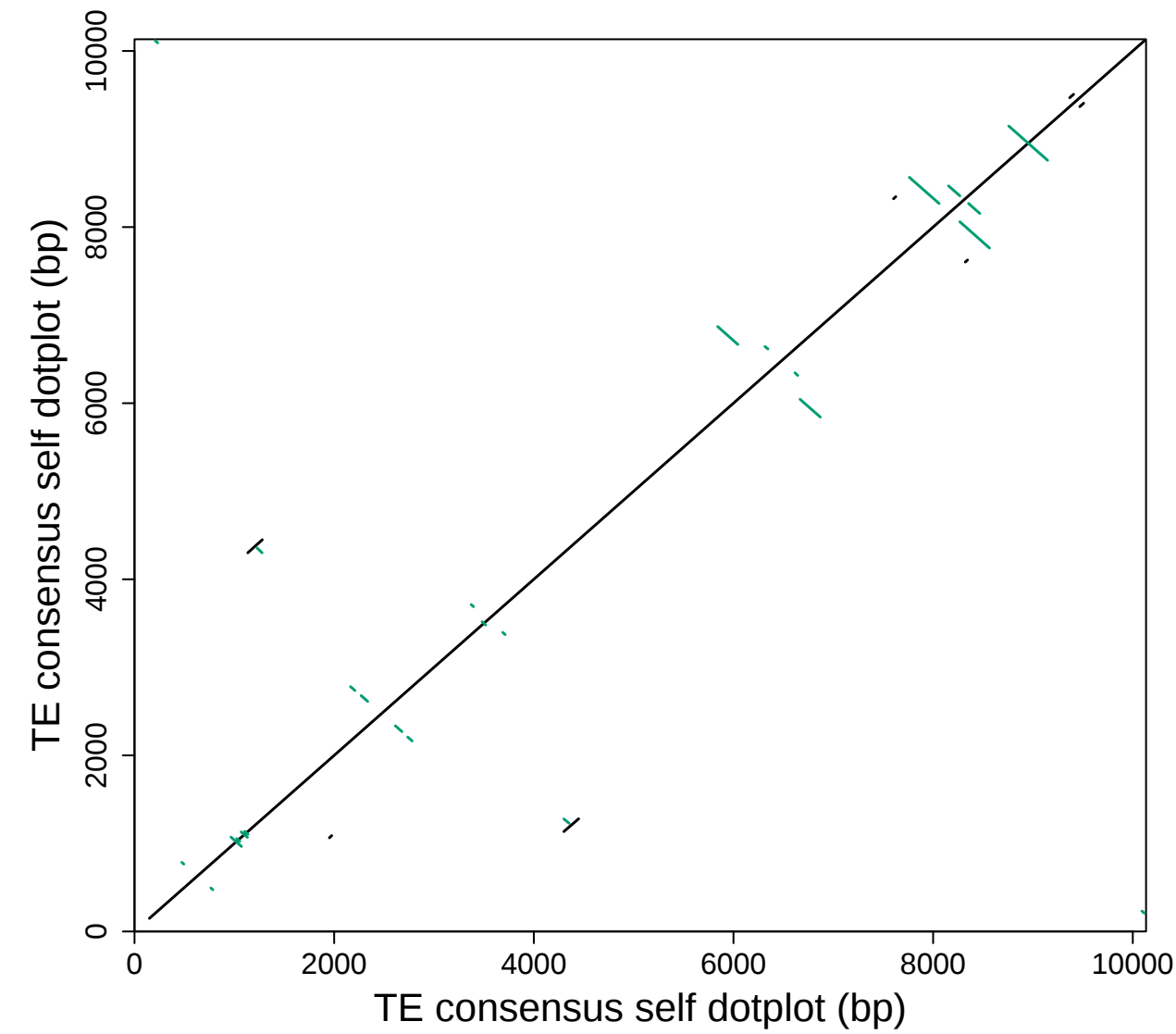
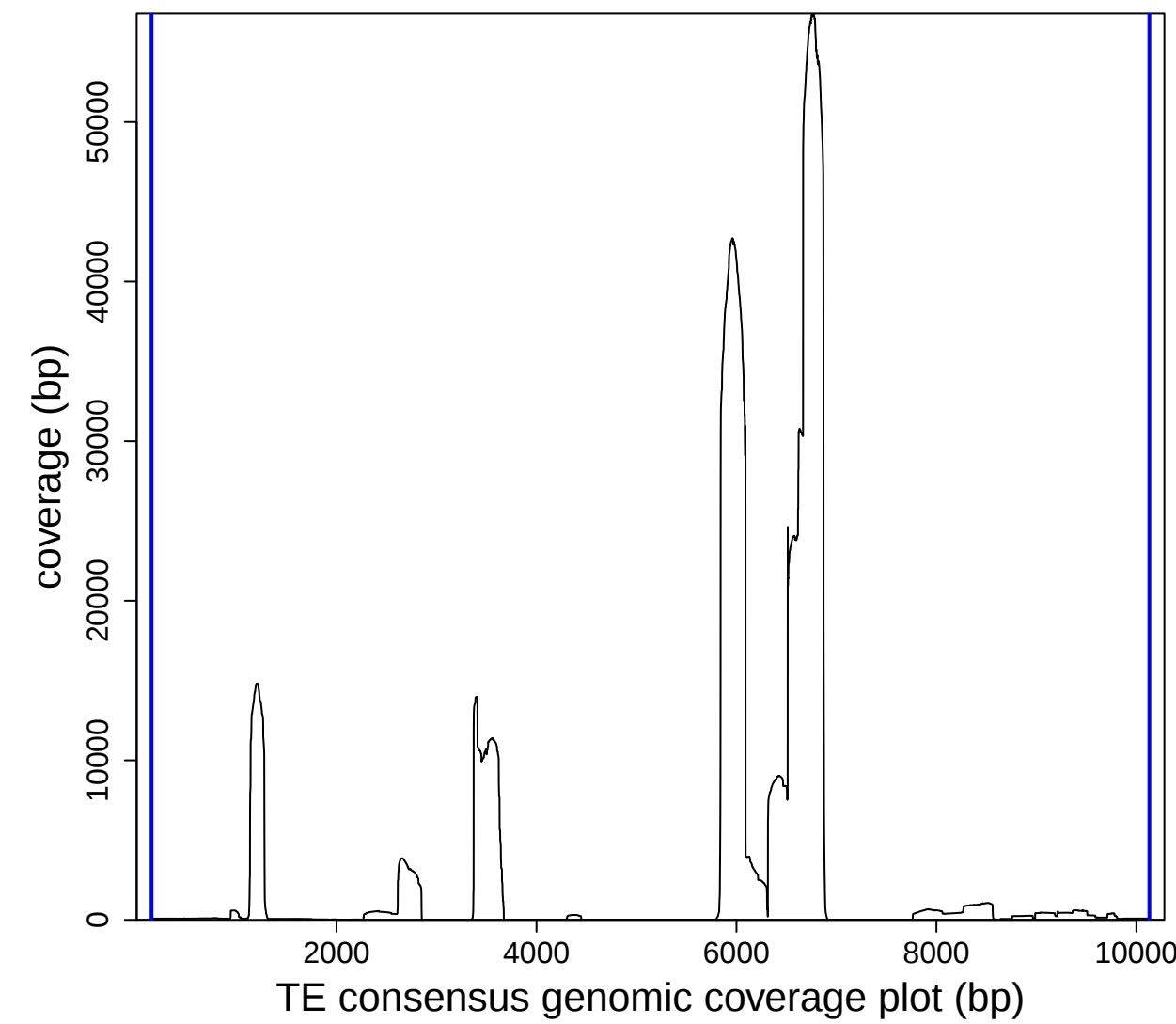
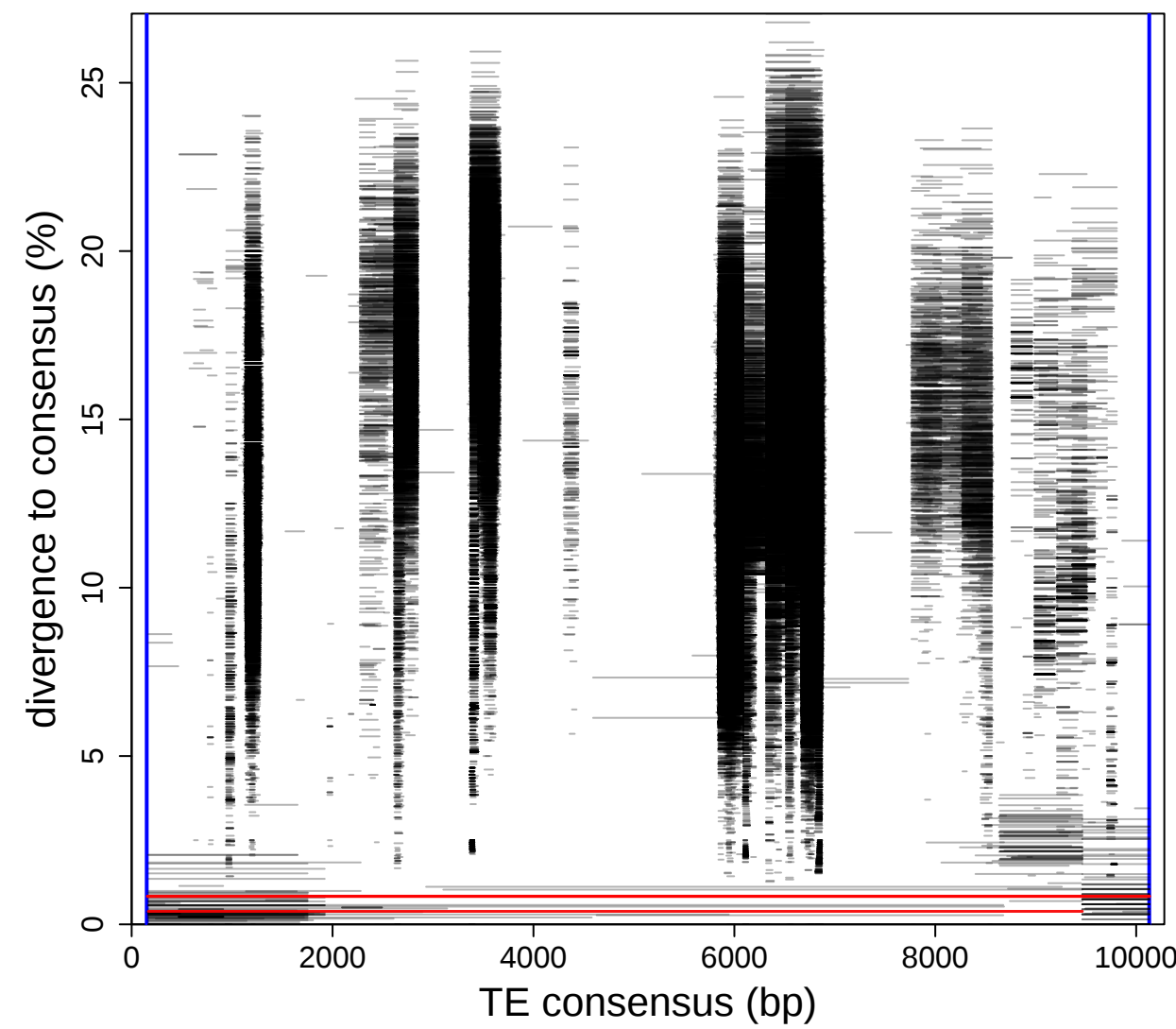
b.bed_uf.bed_g_2.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.fa
size: 9976bp; fragments: 159254; full length: 3 (>=8978.4bp)

After TEtrimmer 9976 bp

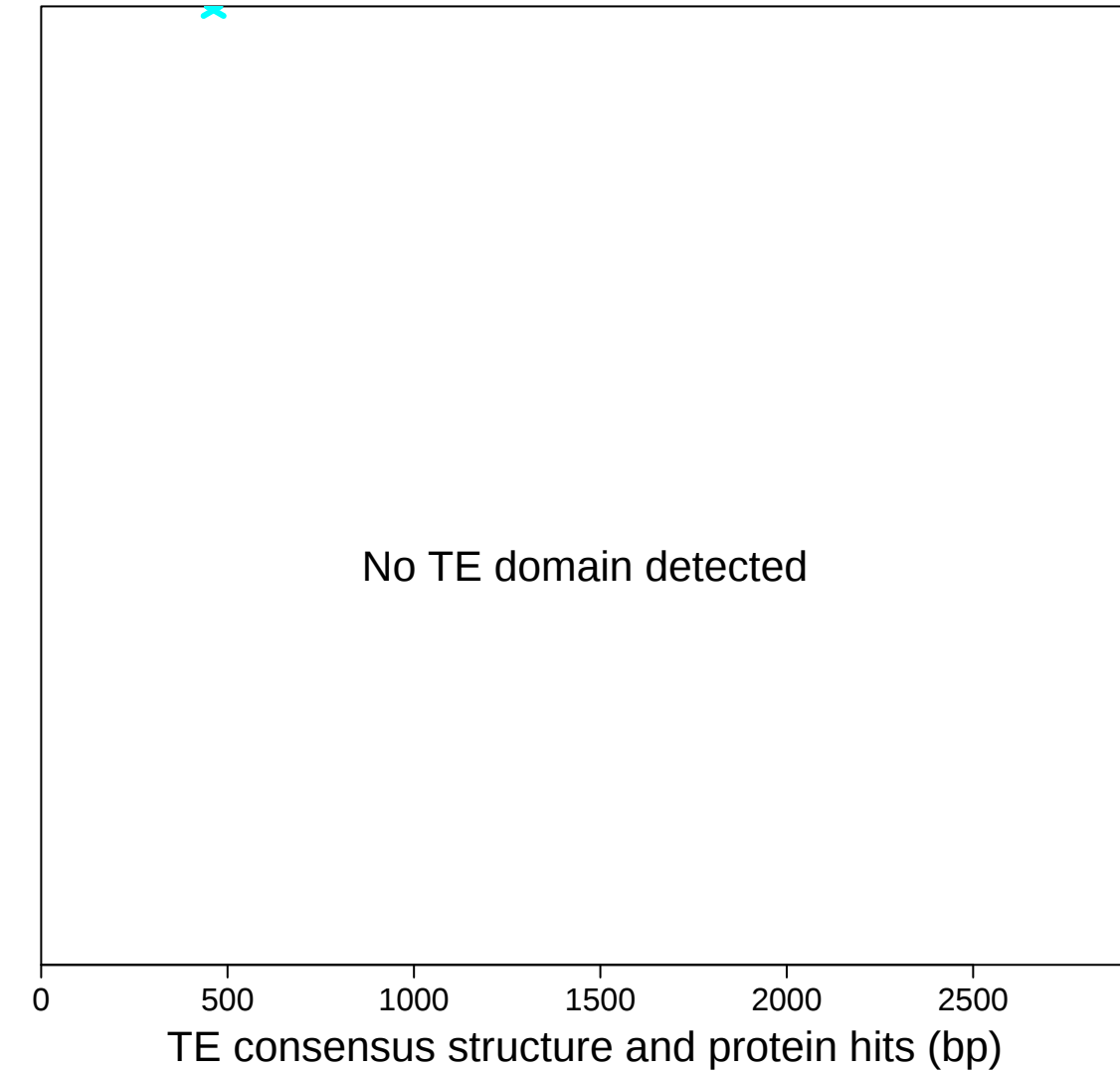
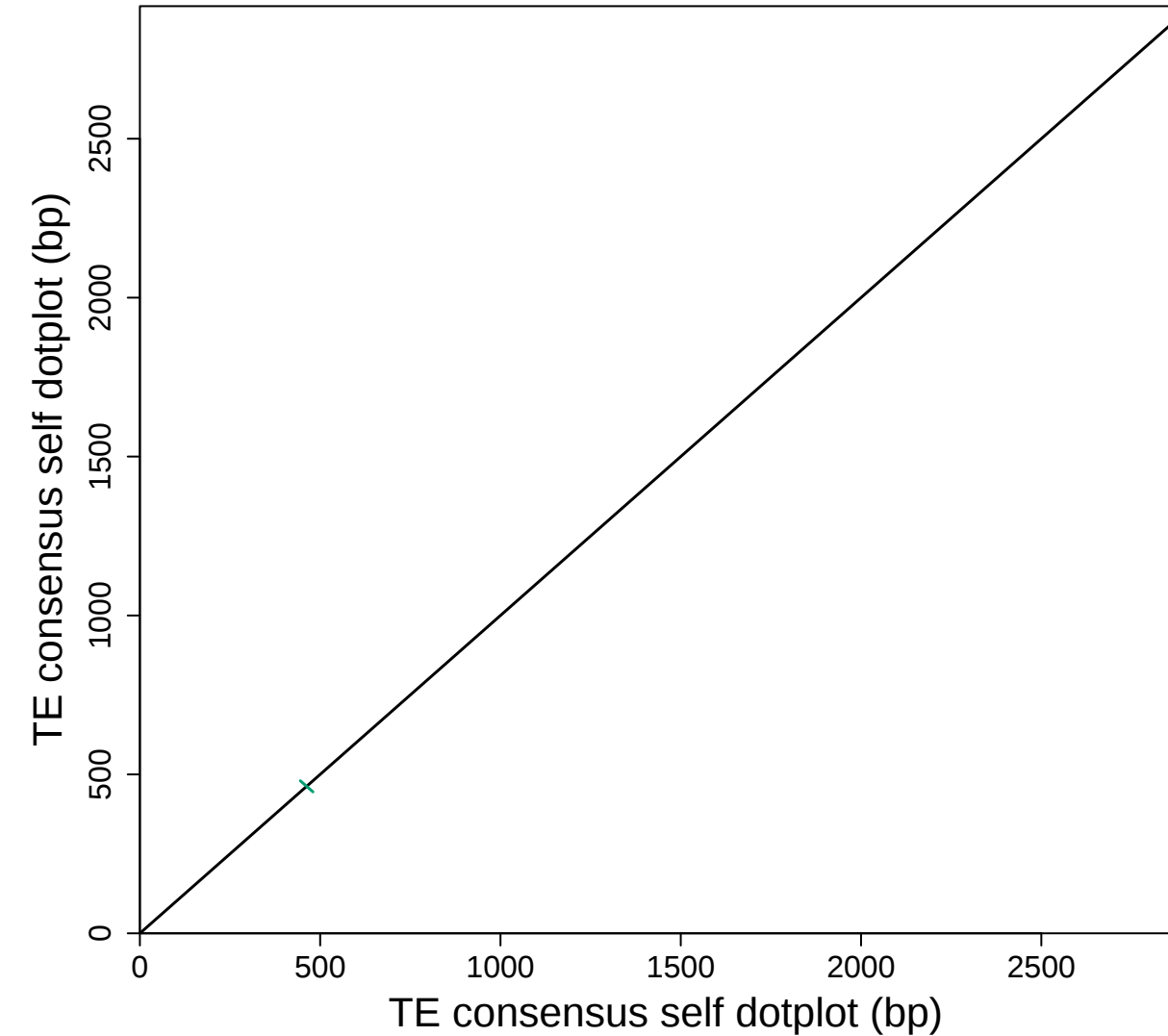
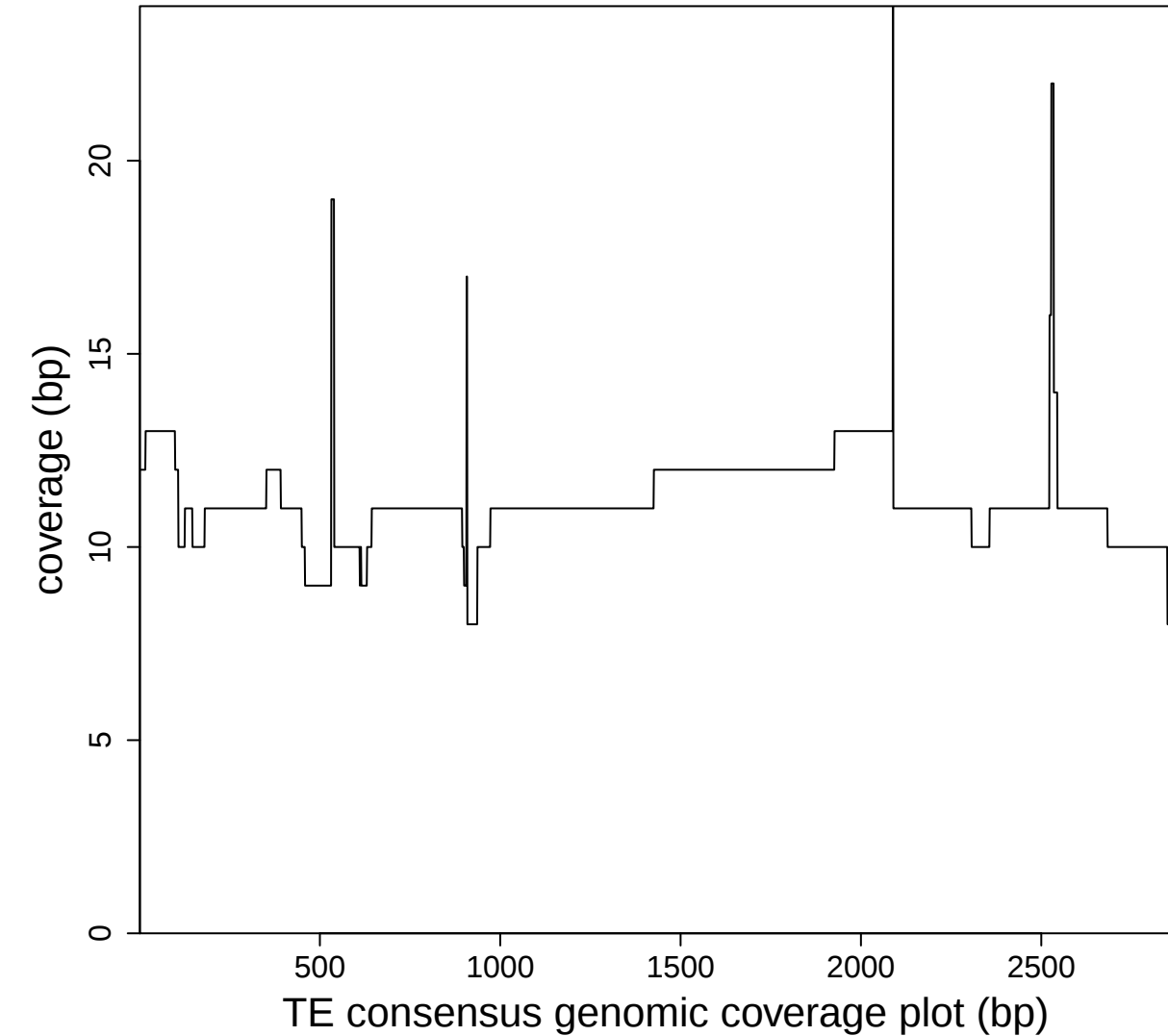
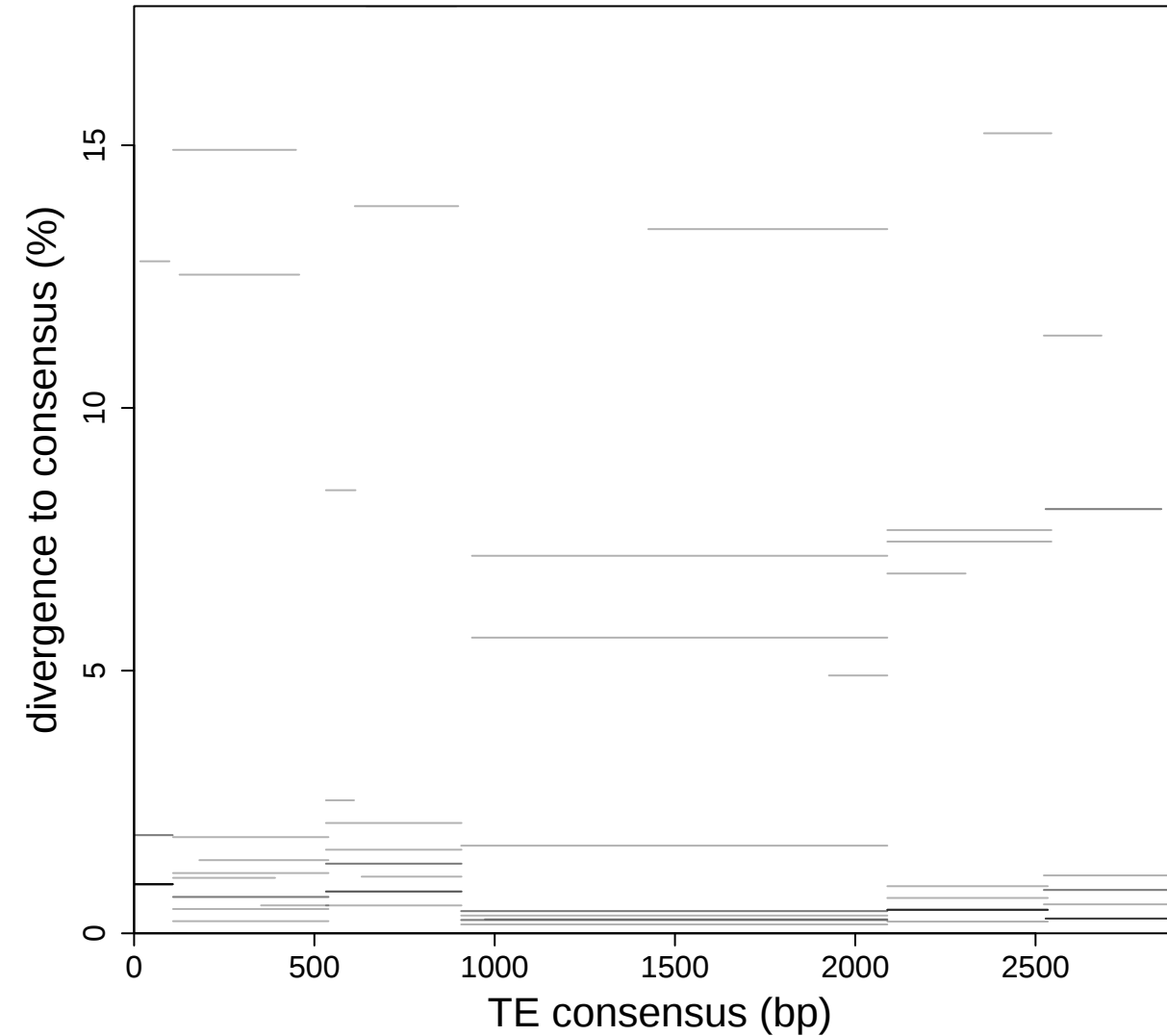


.fasta.b.bed_uf.bed_g_2.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa
size: 10280bp; fragments: 158050; full length: 3 (>=9252bp)

After TEtrimmer Extended plot Blue lines are boundaries

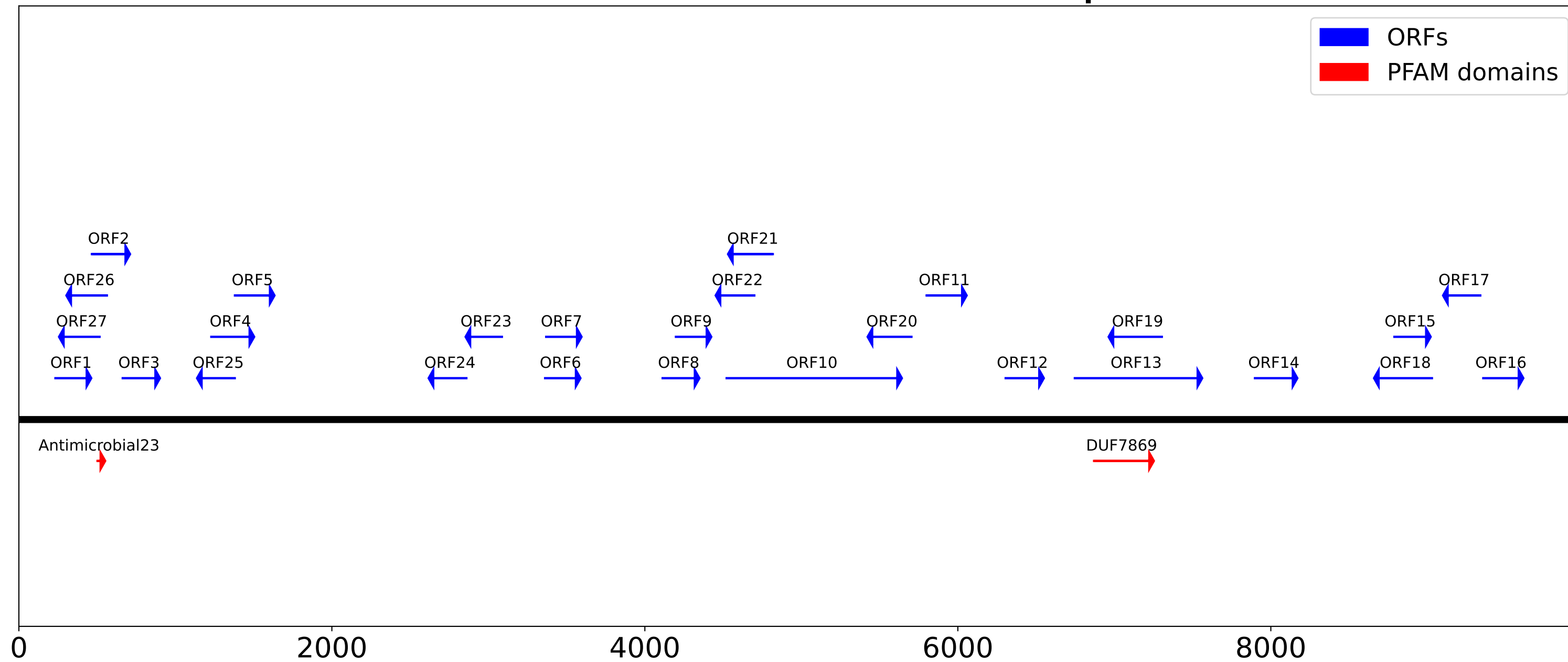


TE: ltr_1_family_936
size: 2917bp; fragments: 75; full length: 0 (≥ 2625.3 bp)

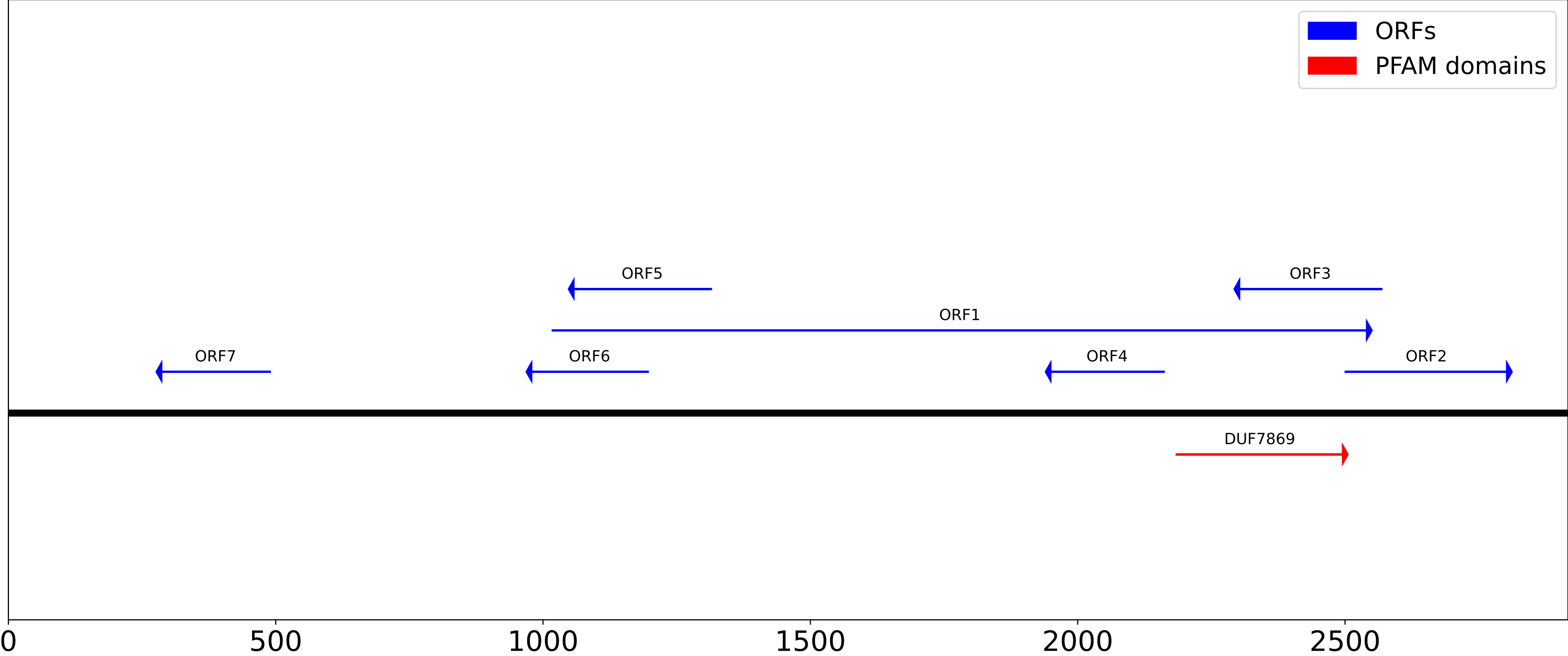


Before TEtrimmer 2917 bp

After TEtrimmer ORF and PFAM domain plot



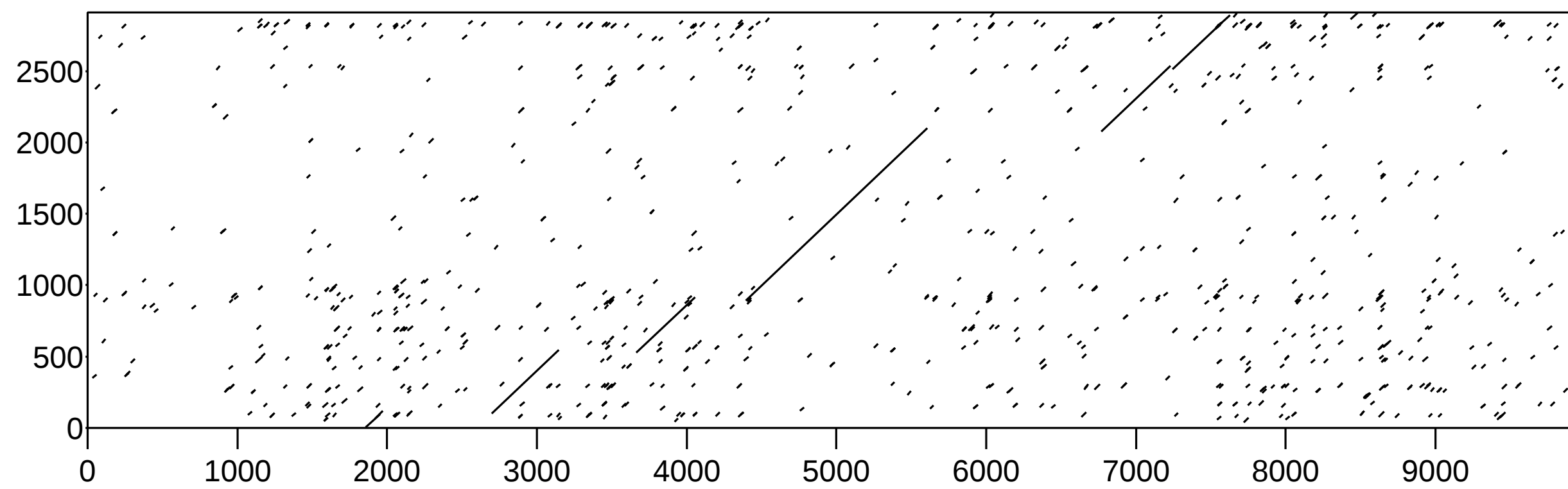
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)