

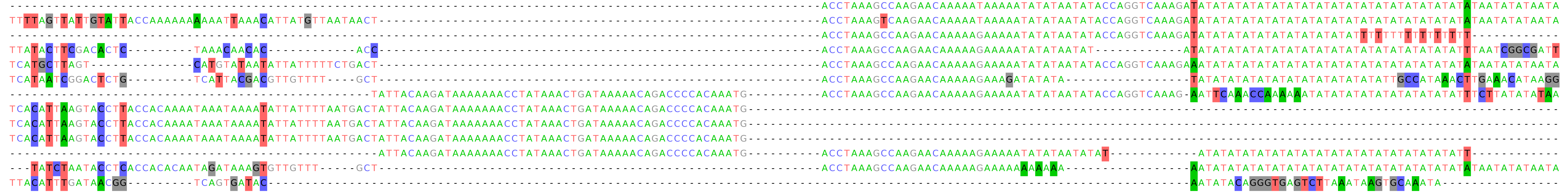
Start crop Point



End crop Point



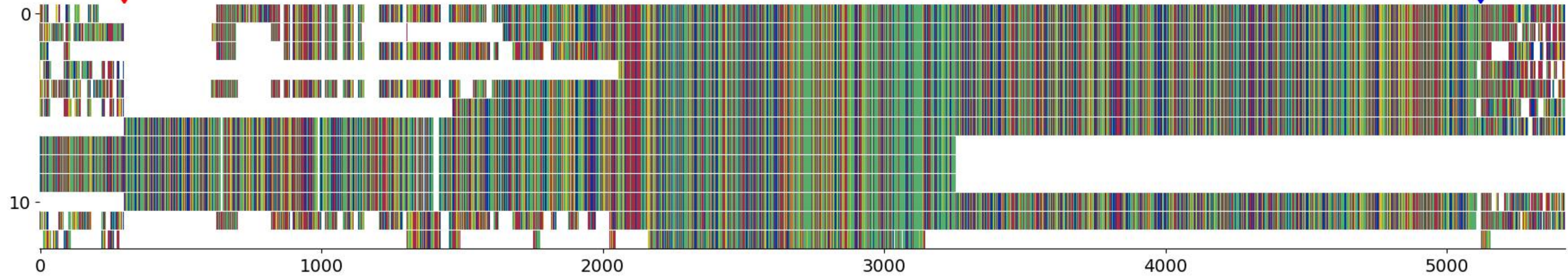
MSA length = 4822



Start crop Point

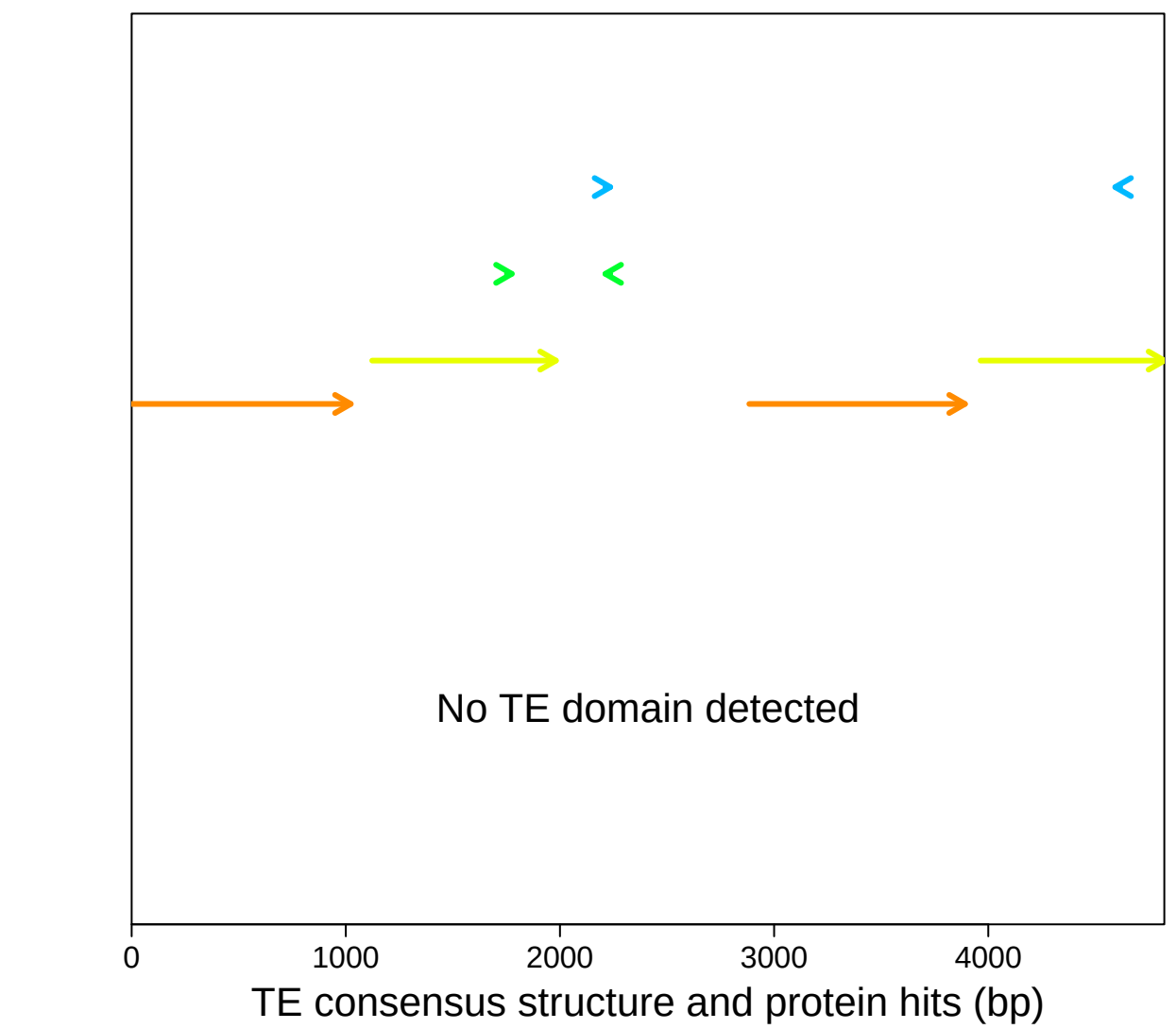
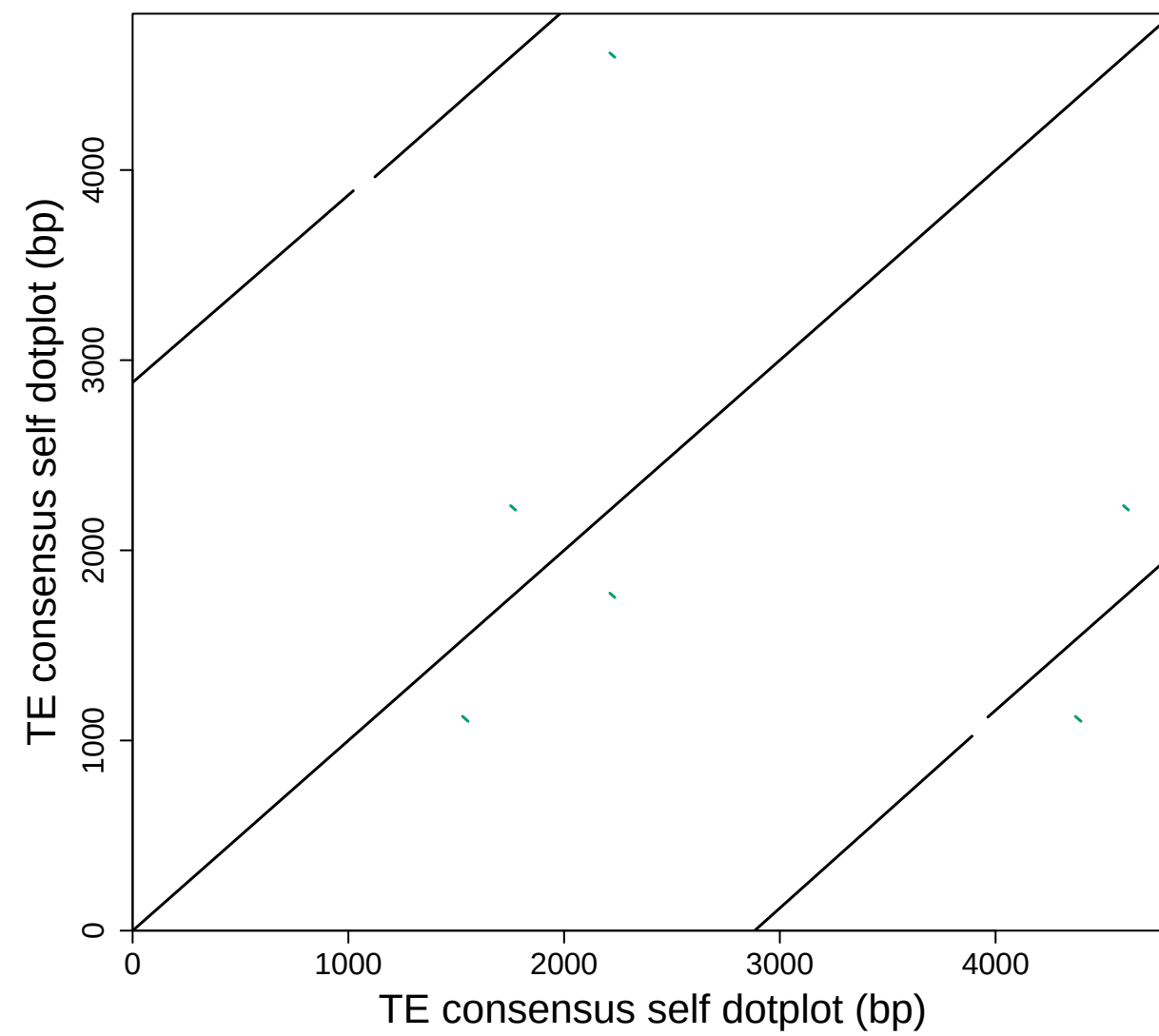
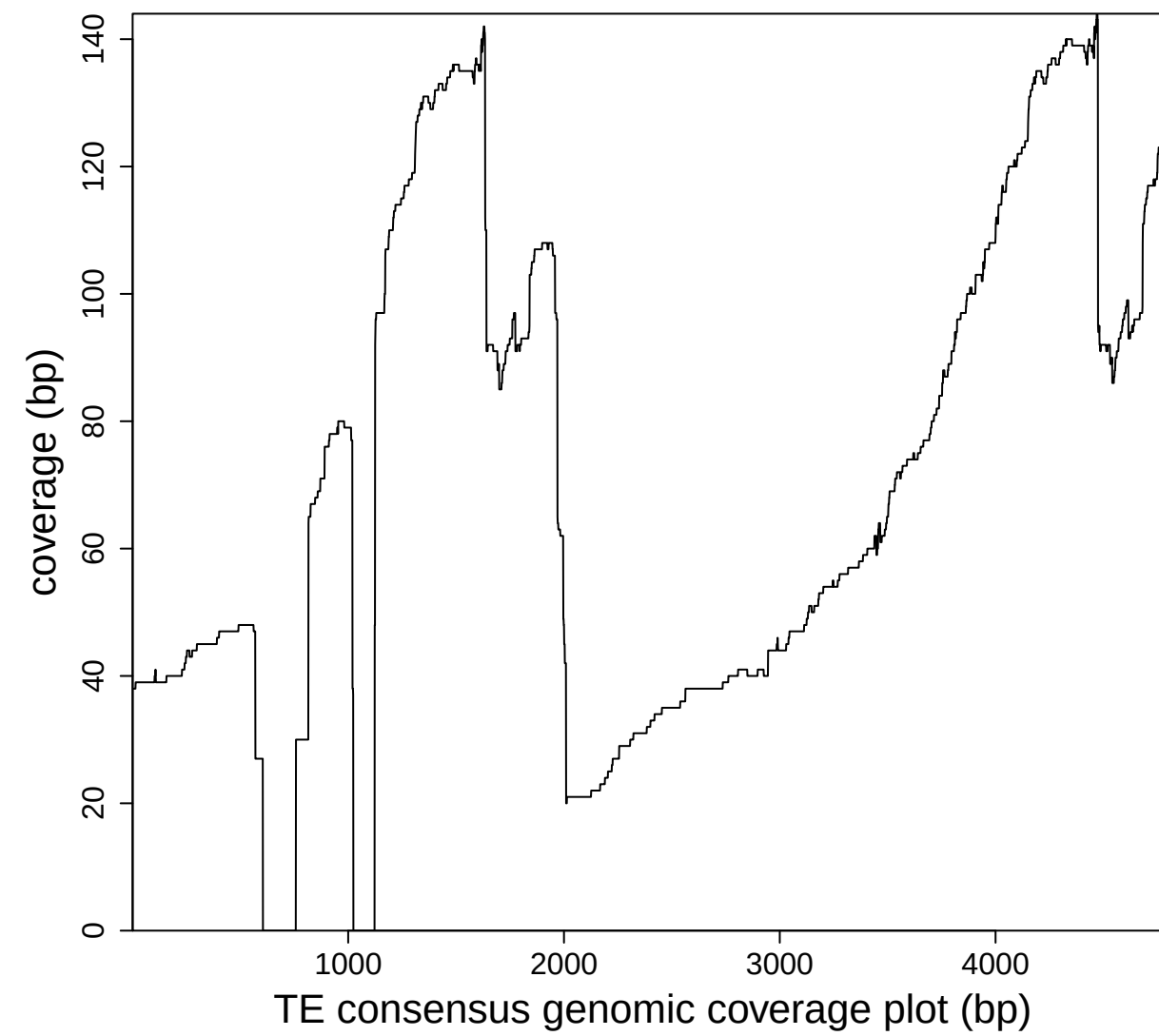
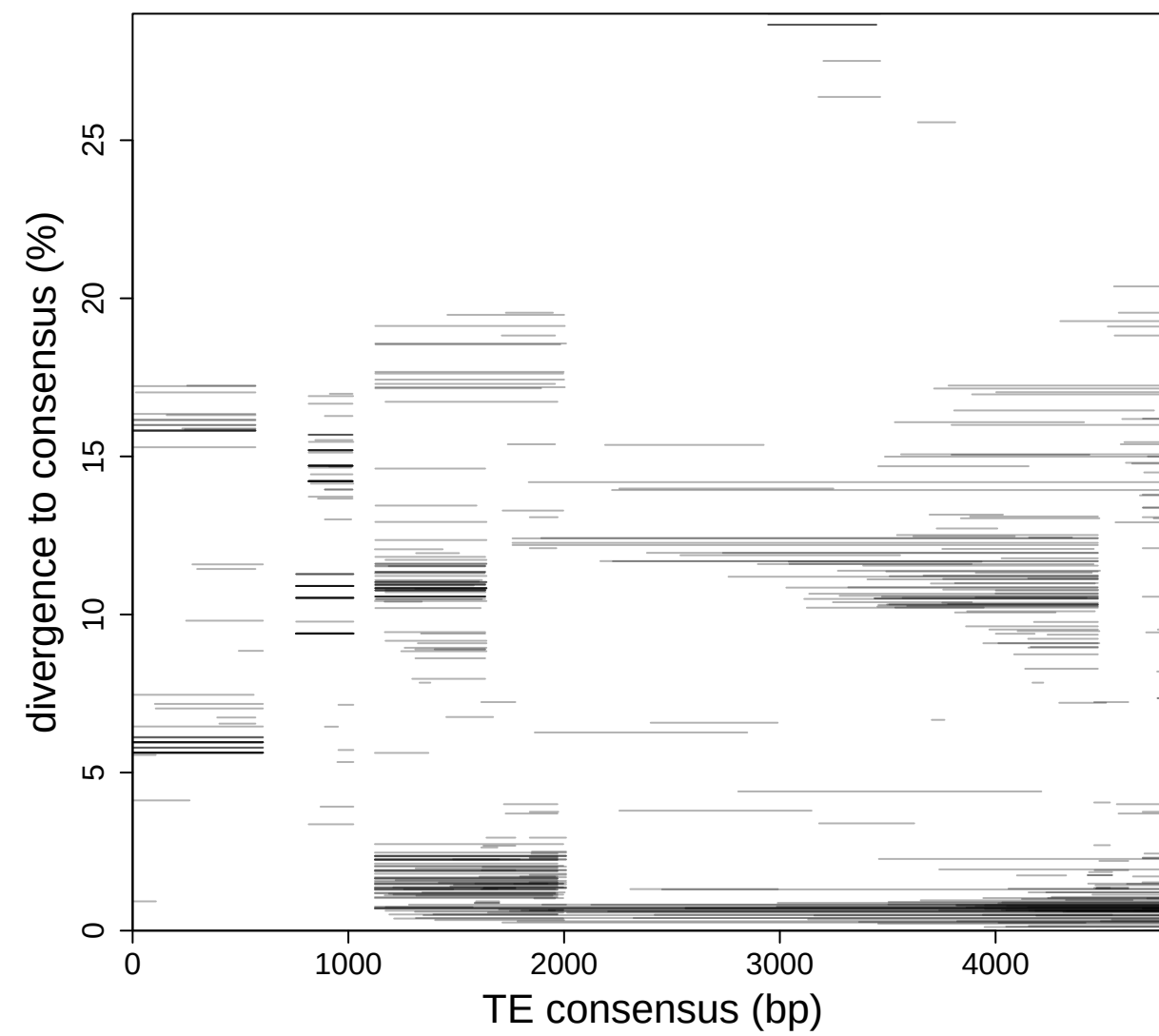


End crop Point



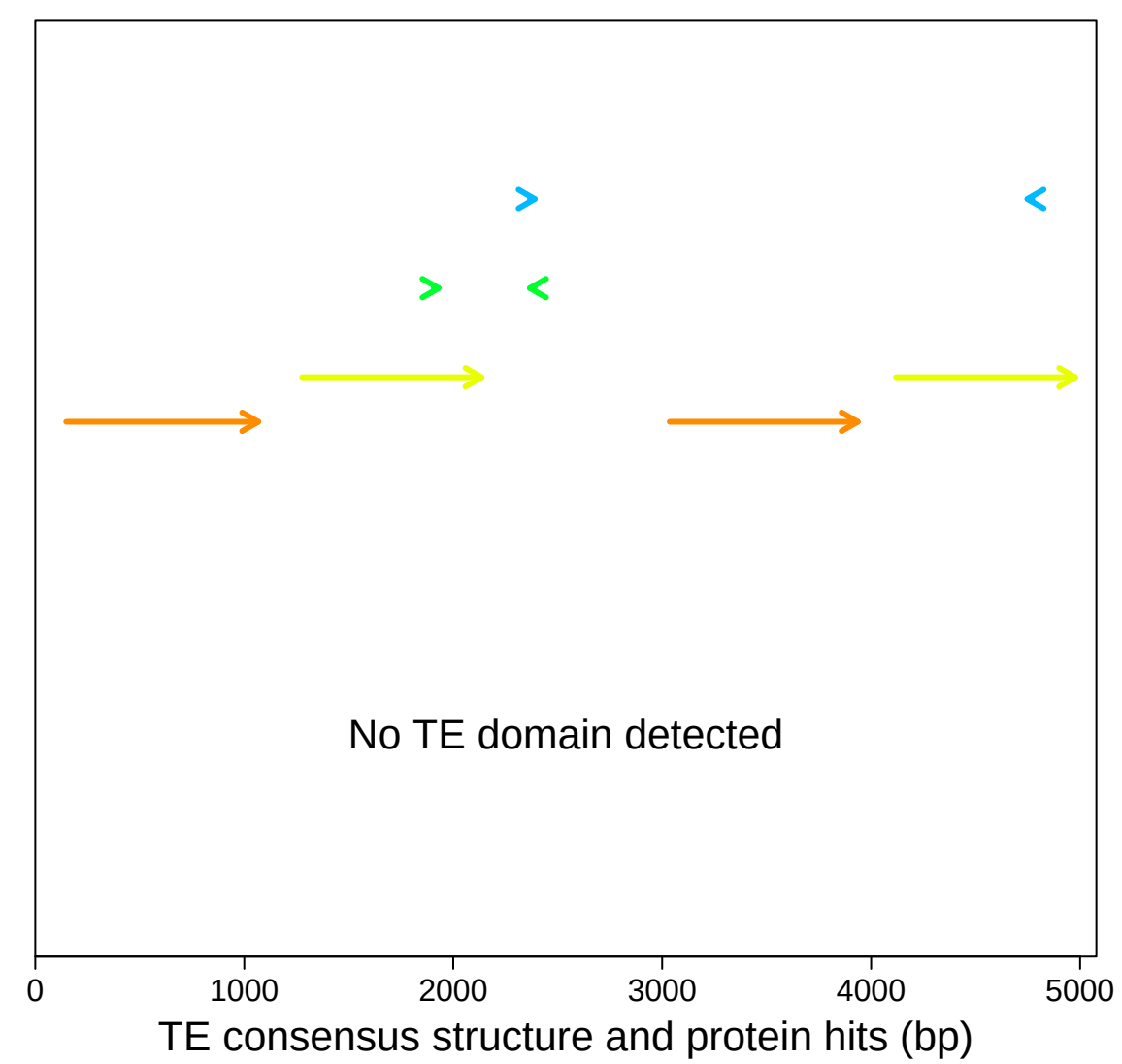
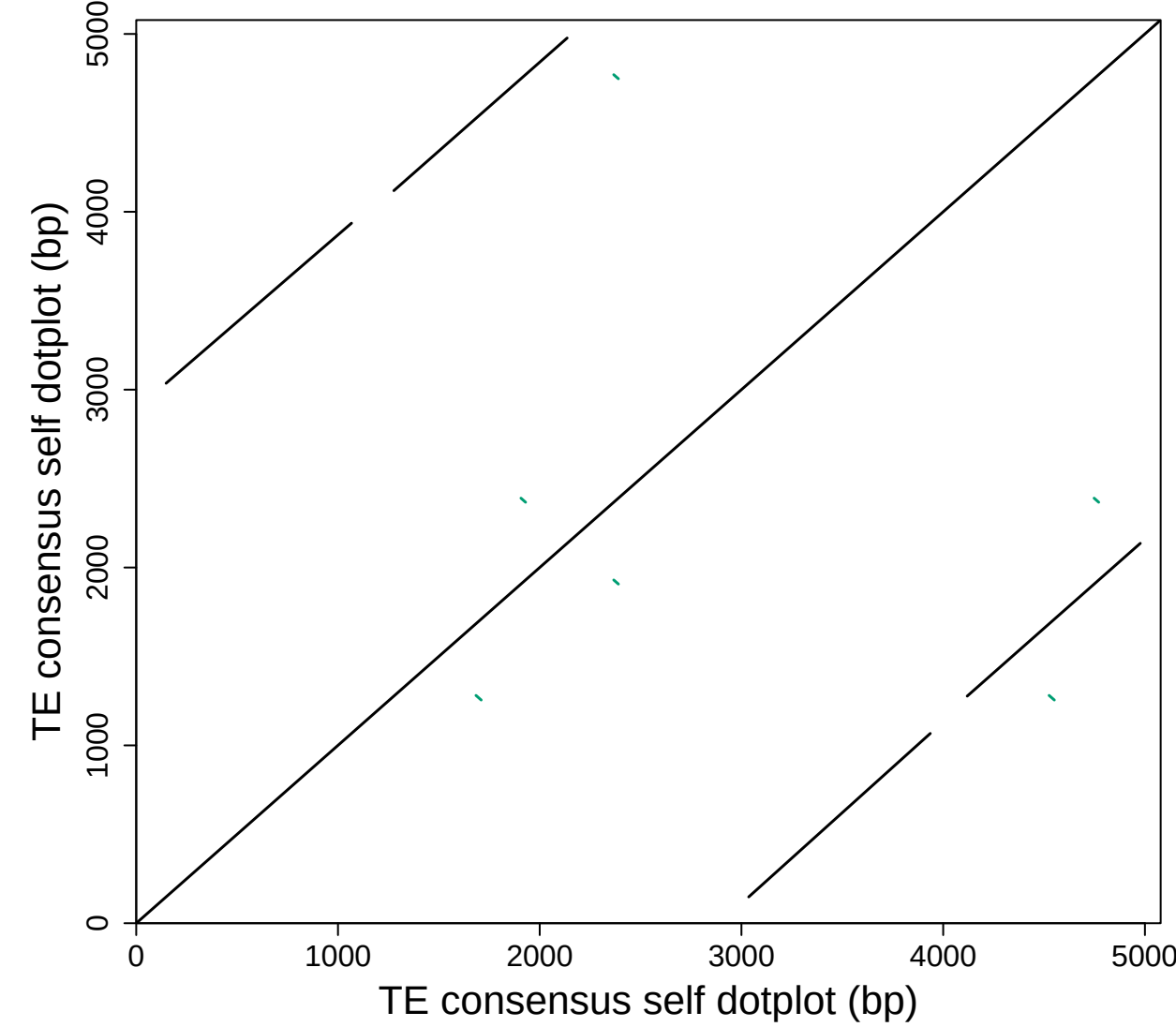
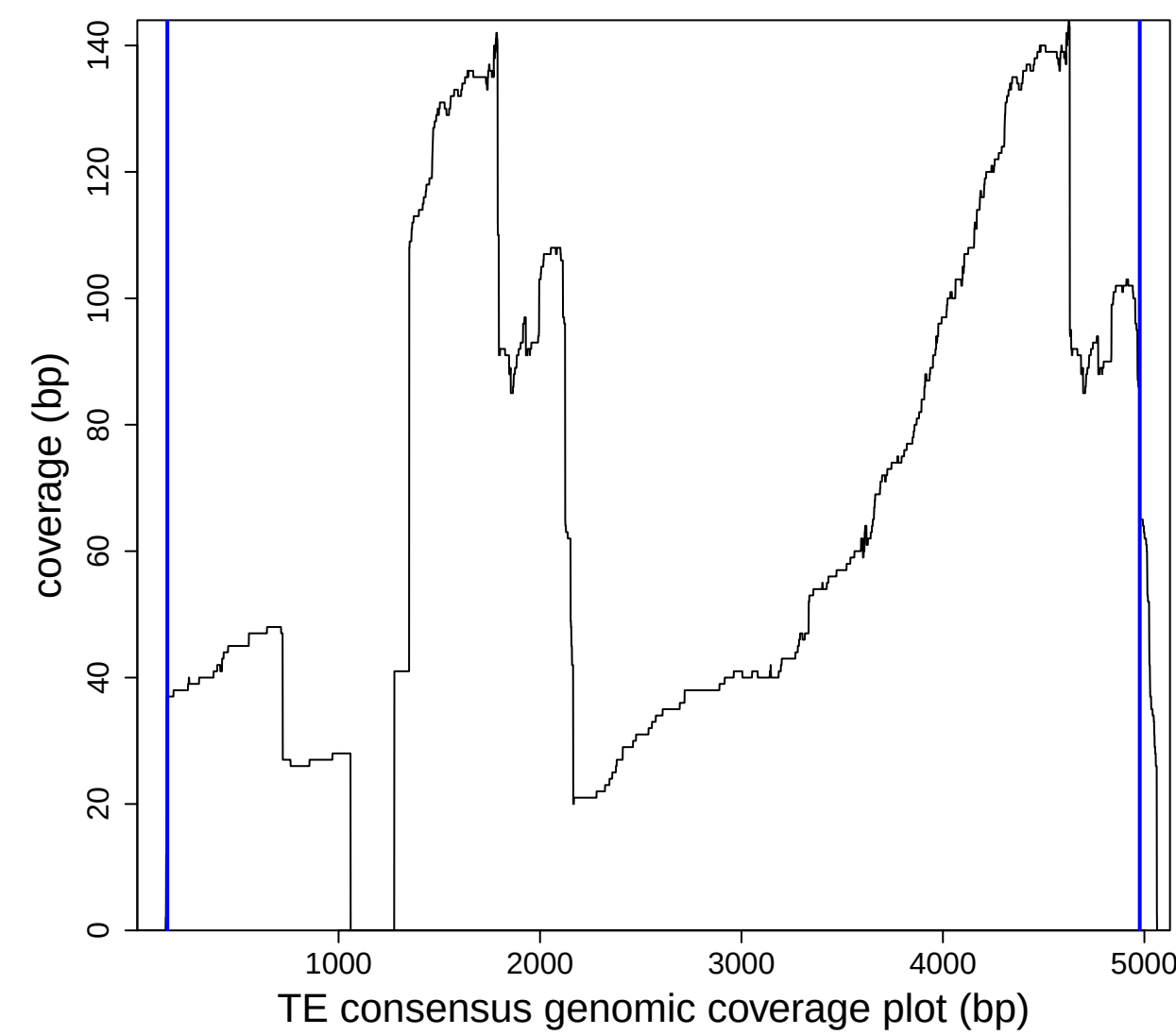
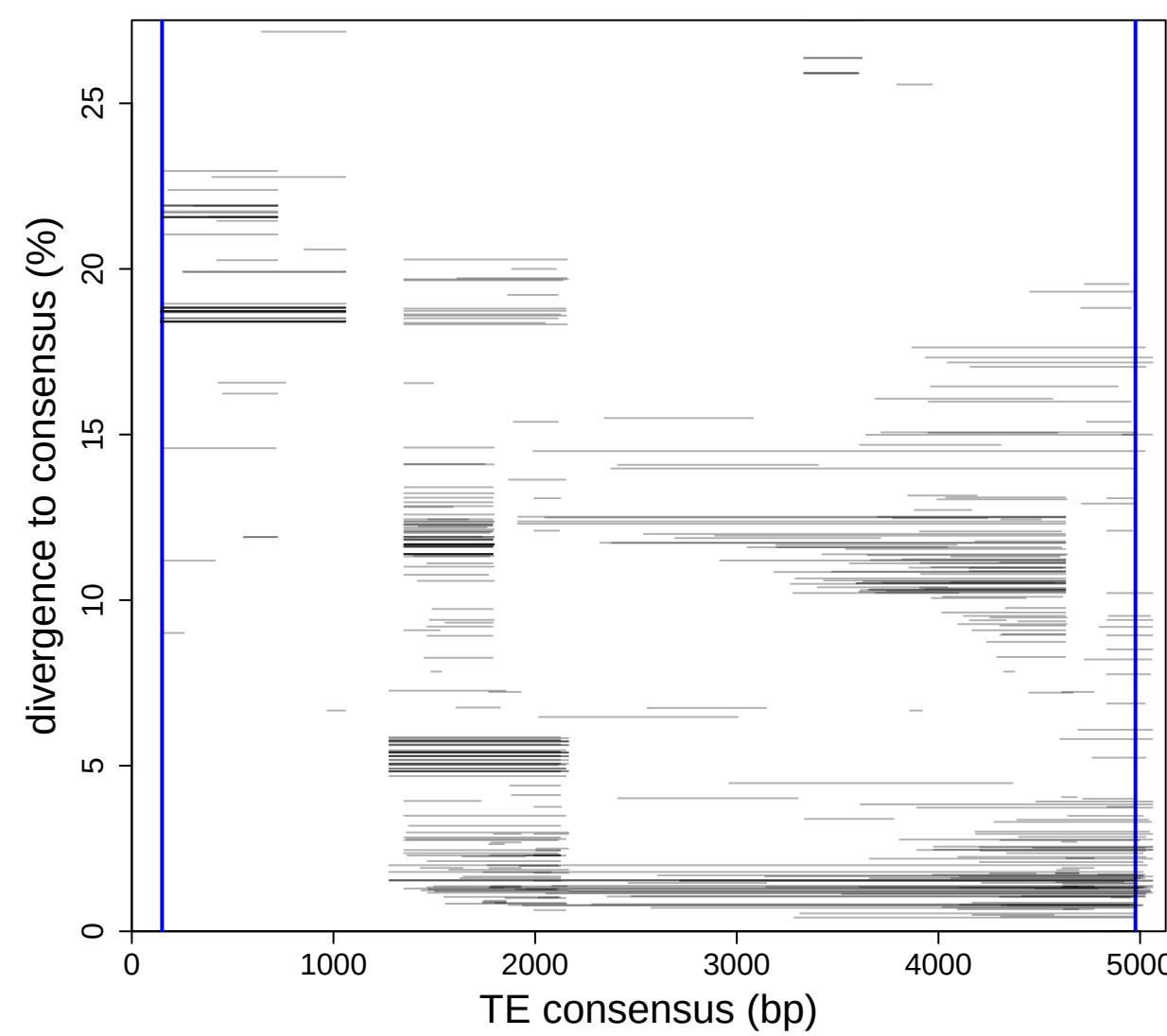
bed uf.bed g 2.bed fm 1.bed 0 0 bcln.fa aln.fa cl.fa gs.fa ce
size: 4822bp; fragments: 535; full length: 0 (>=4339.8bp)

After TEtrimmer 4822 bp

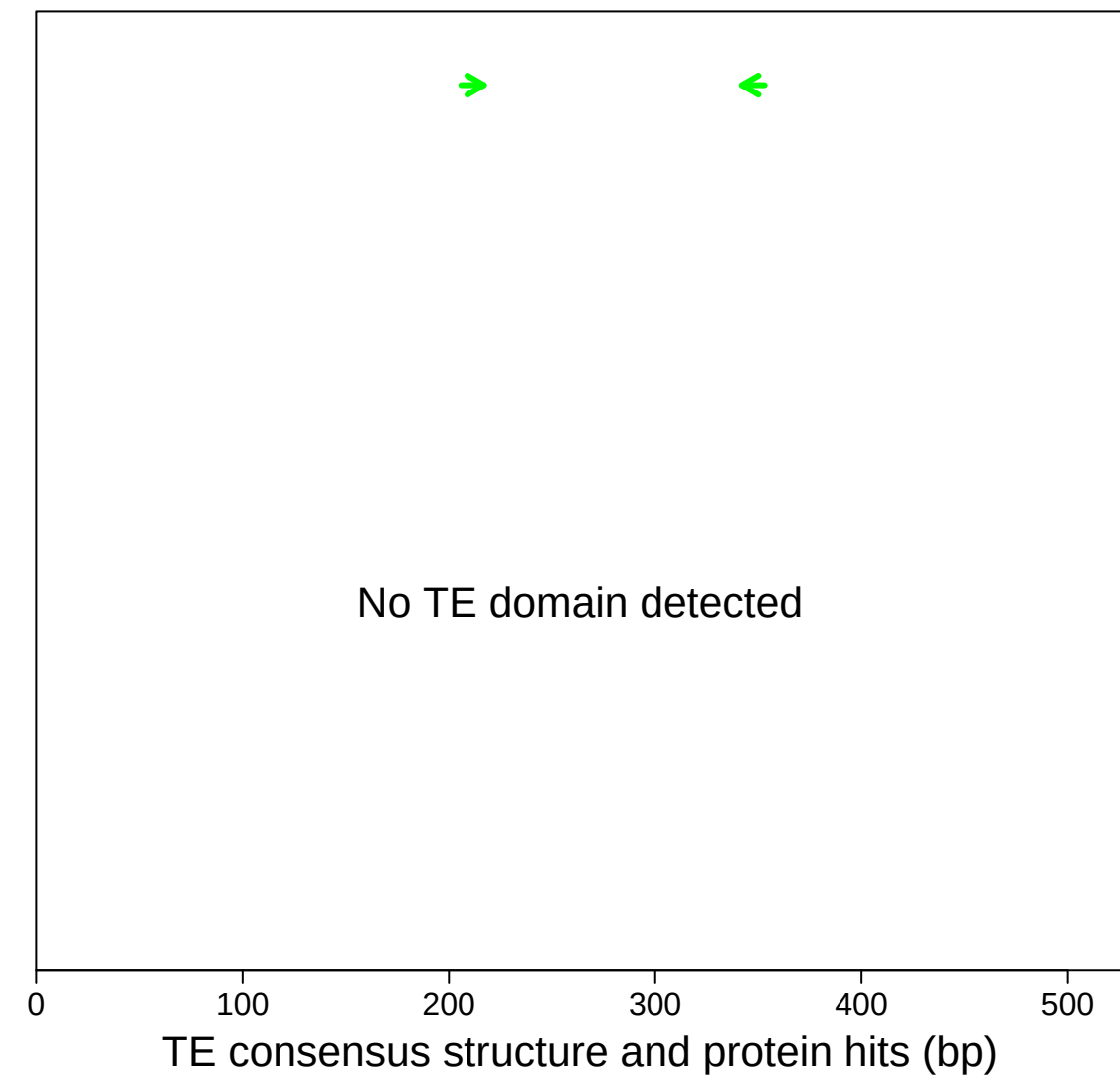
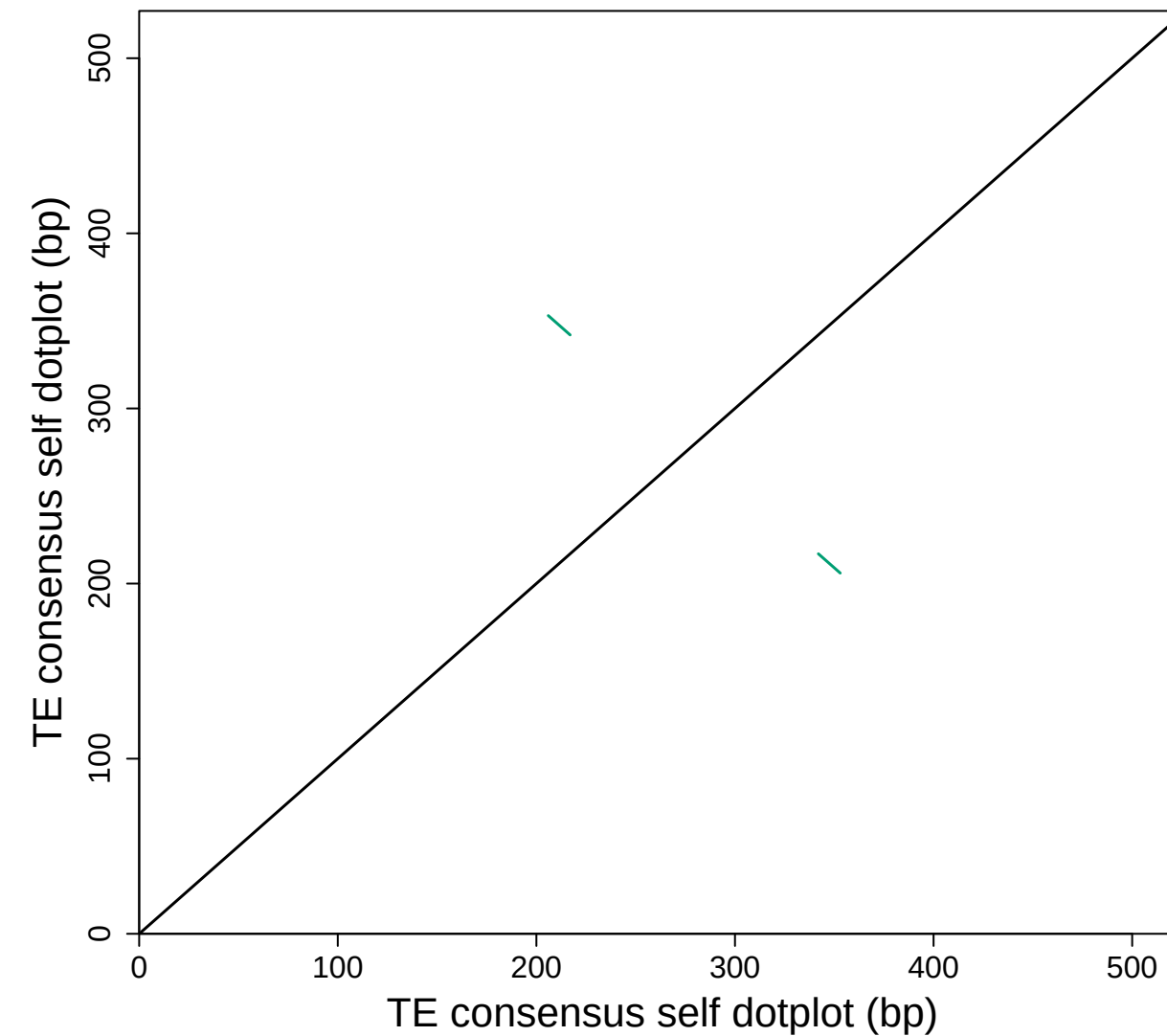
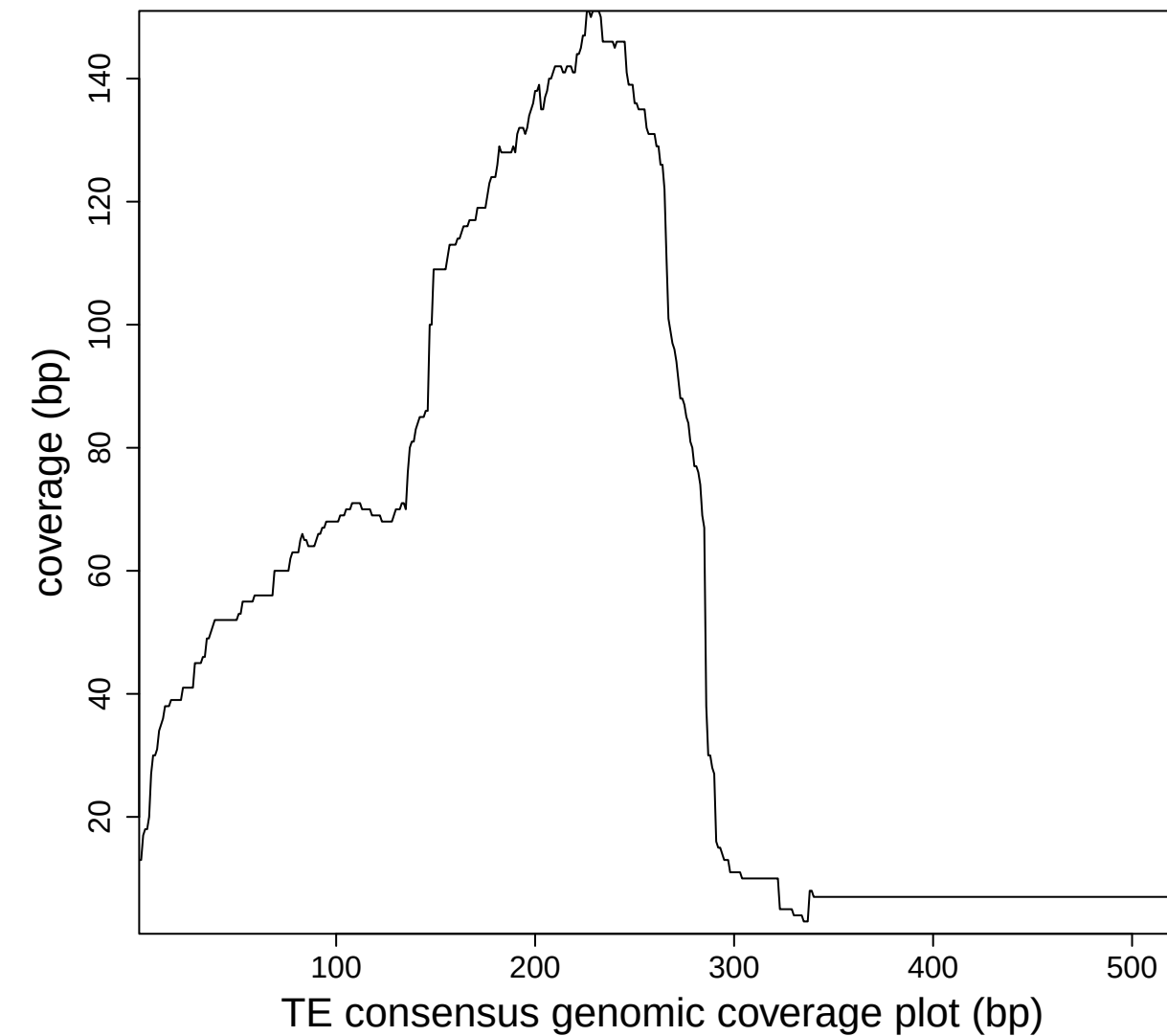
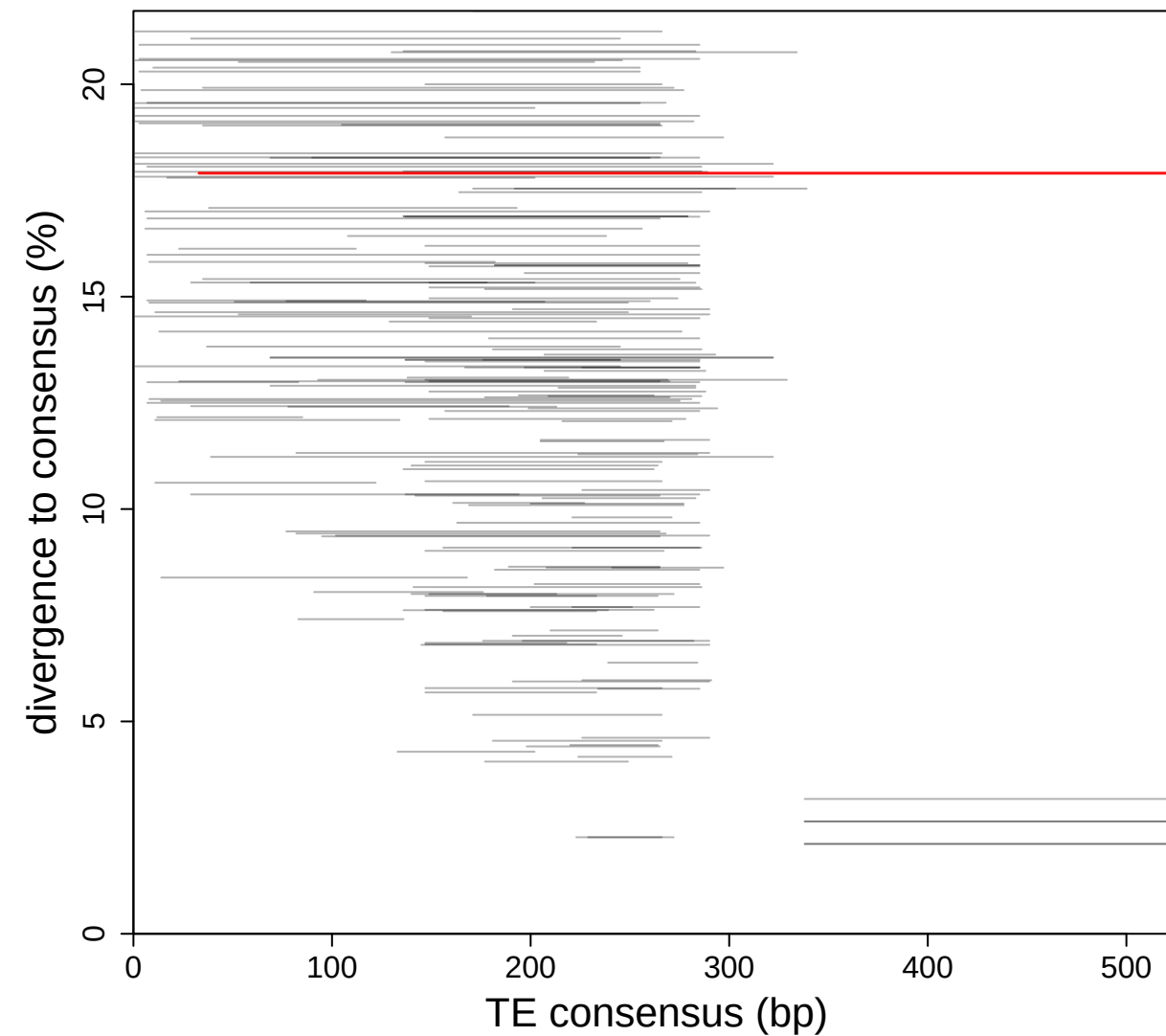


ily_656.fasta.b.bed_uf.bed_g_2.bed_fm_1.bed_0_0_bcln.fa_aln.f
size: 5127bp; fragments: 434; full length: 0 (>=4614.3bp)

After TEtrimmer Extended plot Blue lines are boundaries

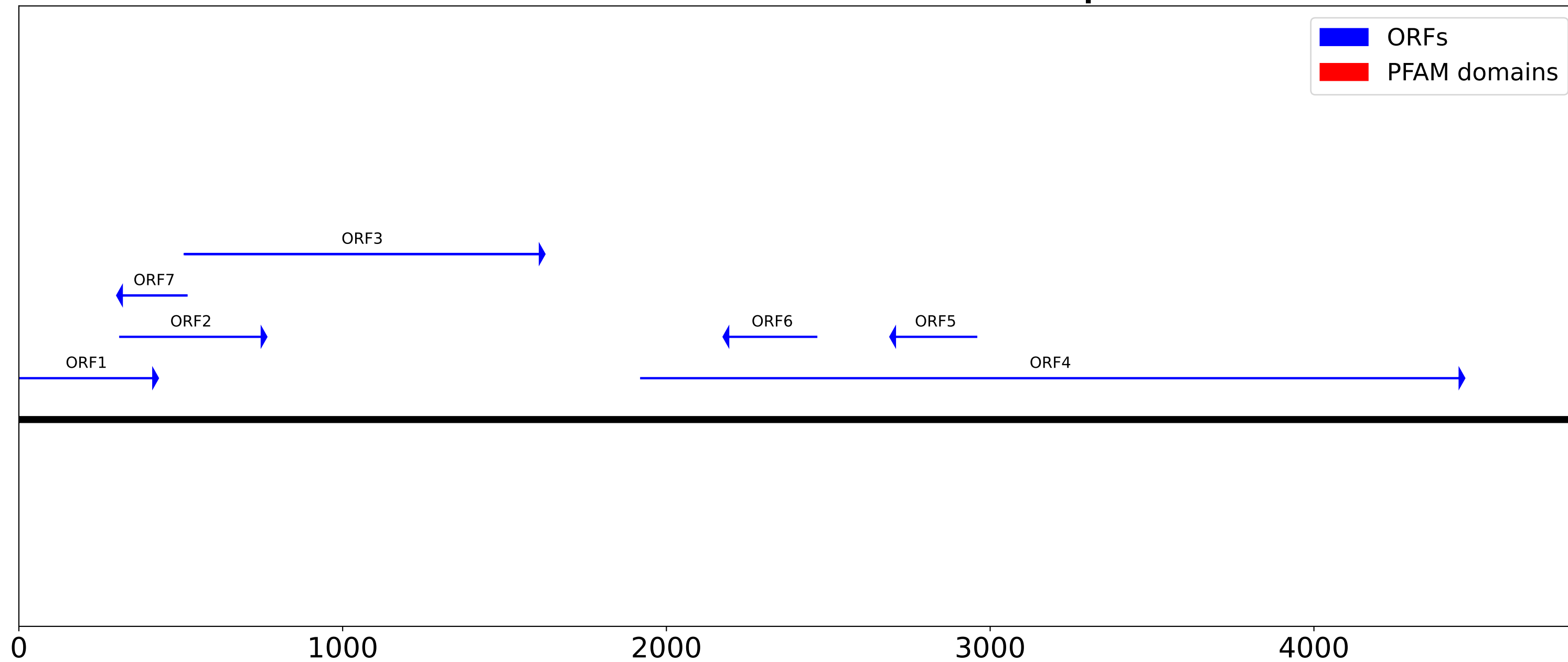


TE: rnd_4_family_656
size: 527bp; fragments: 184; full length: 1 (≥ 474.3 bp)

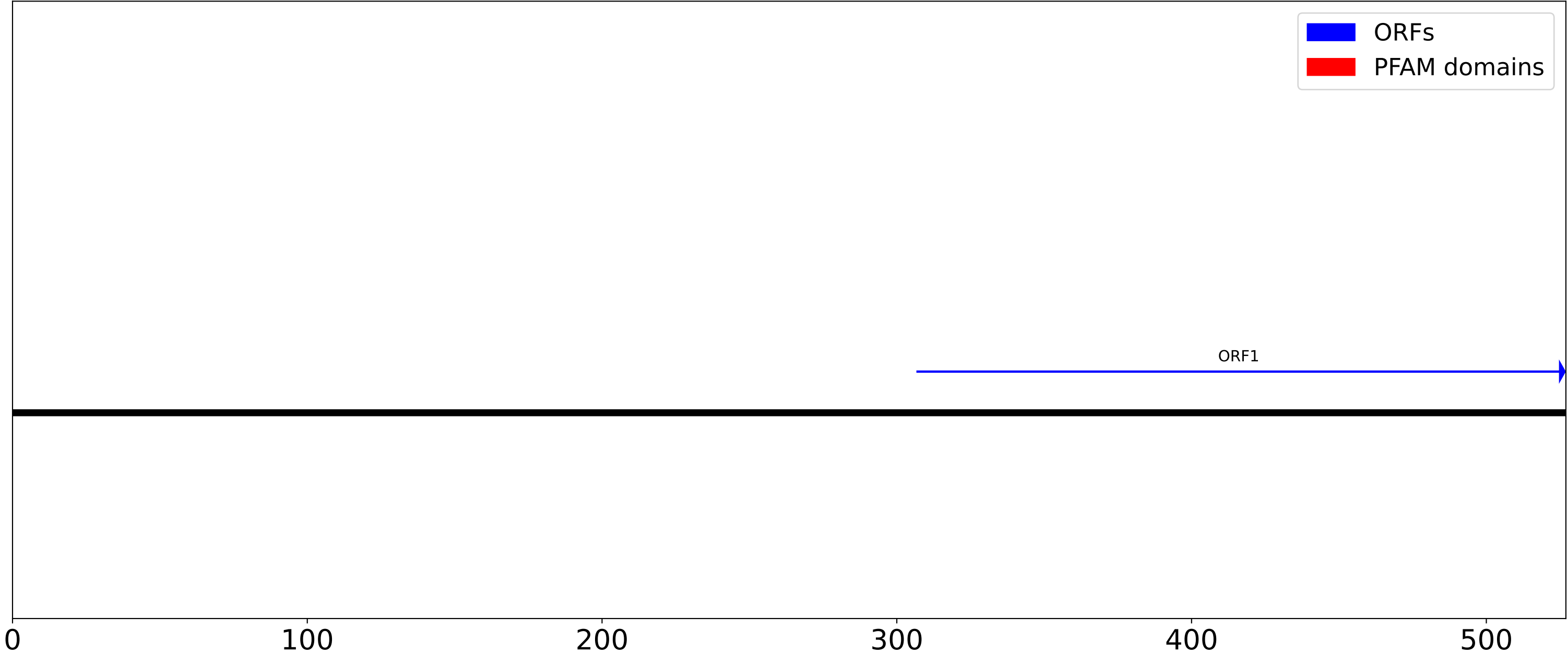


Before TEtrimmer 527 bp

After TEtrimmer ORF and PFAM domain plot



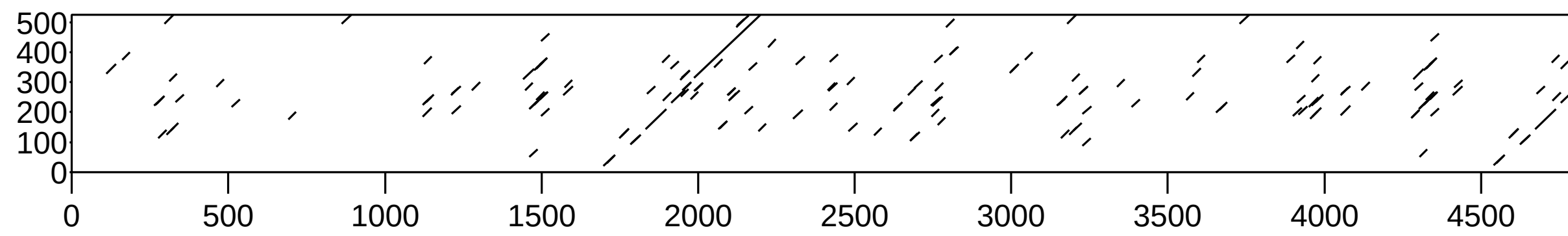
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)