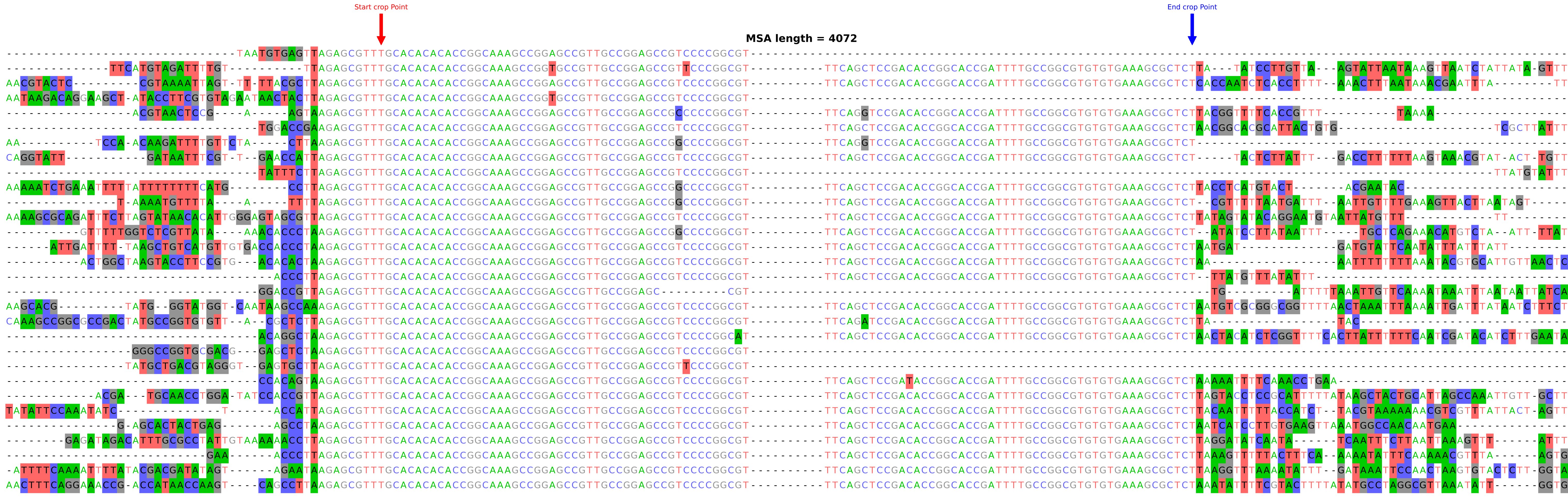
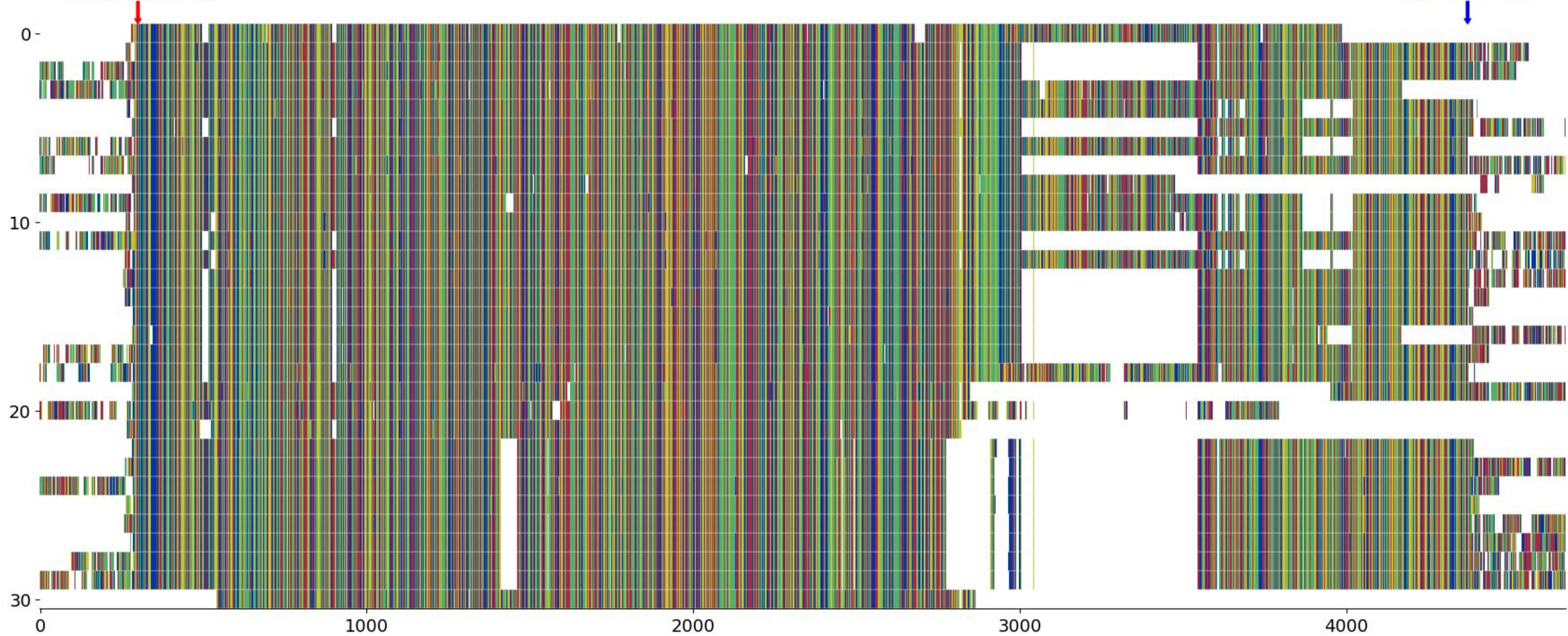




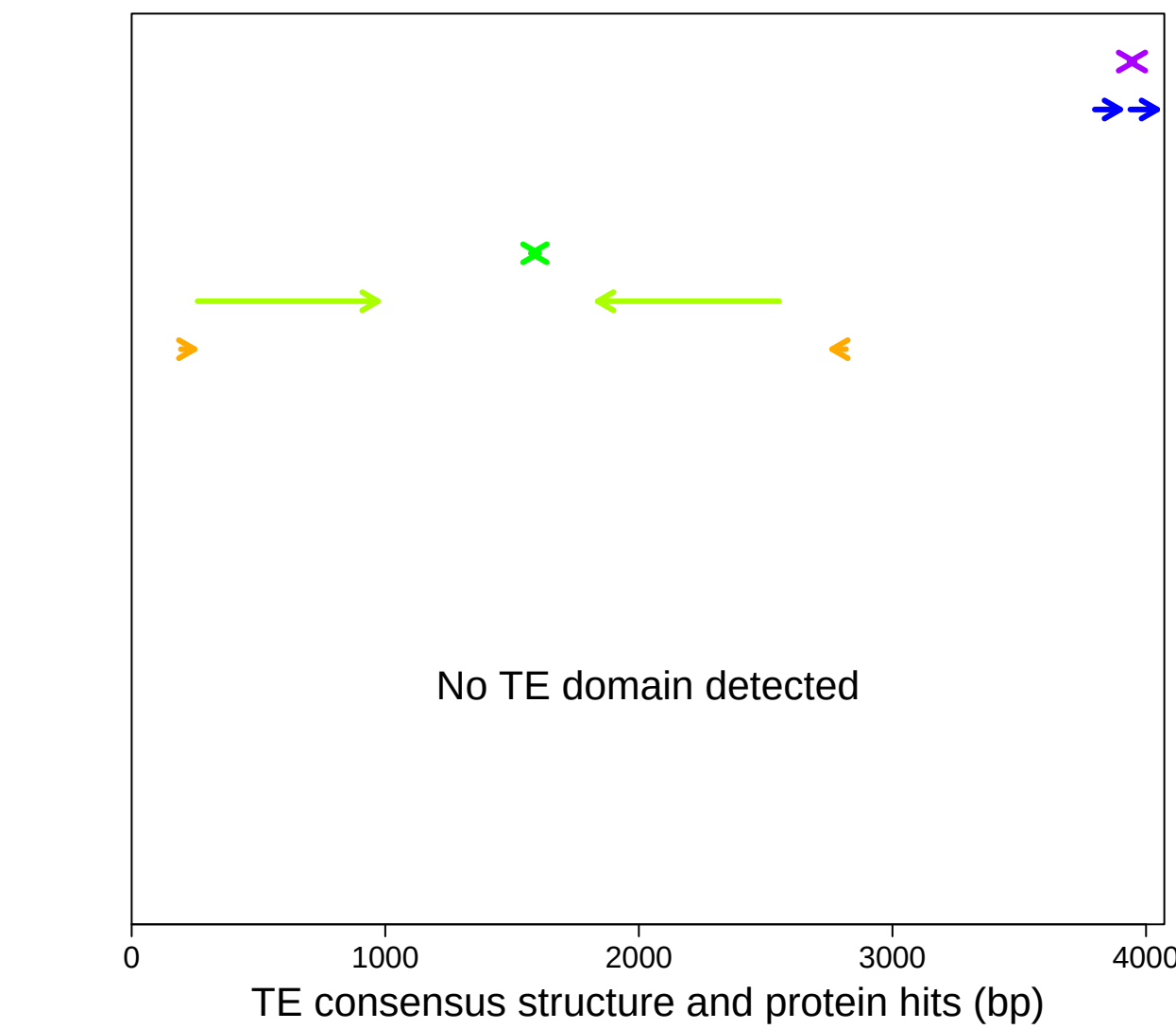
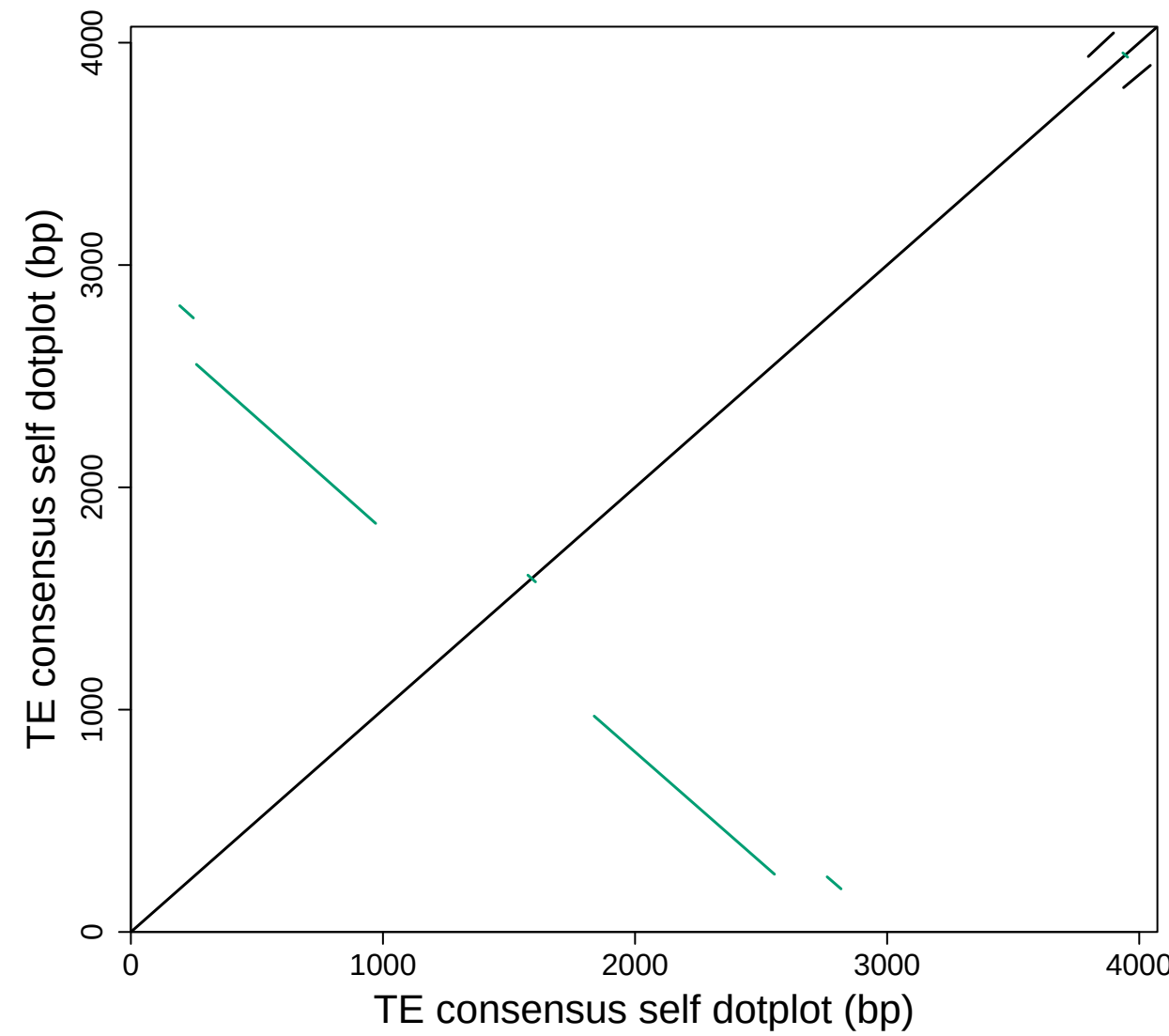
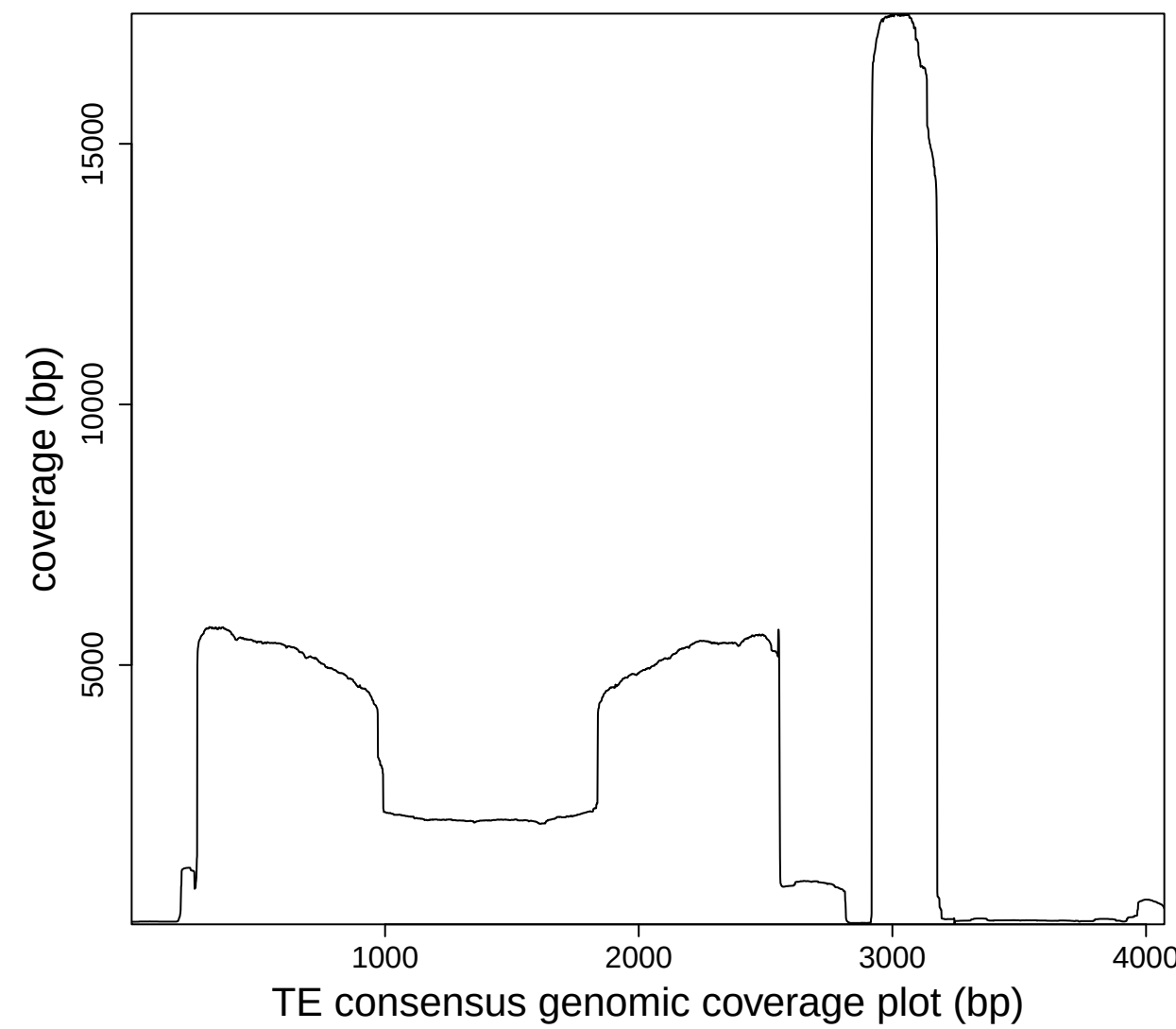
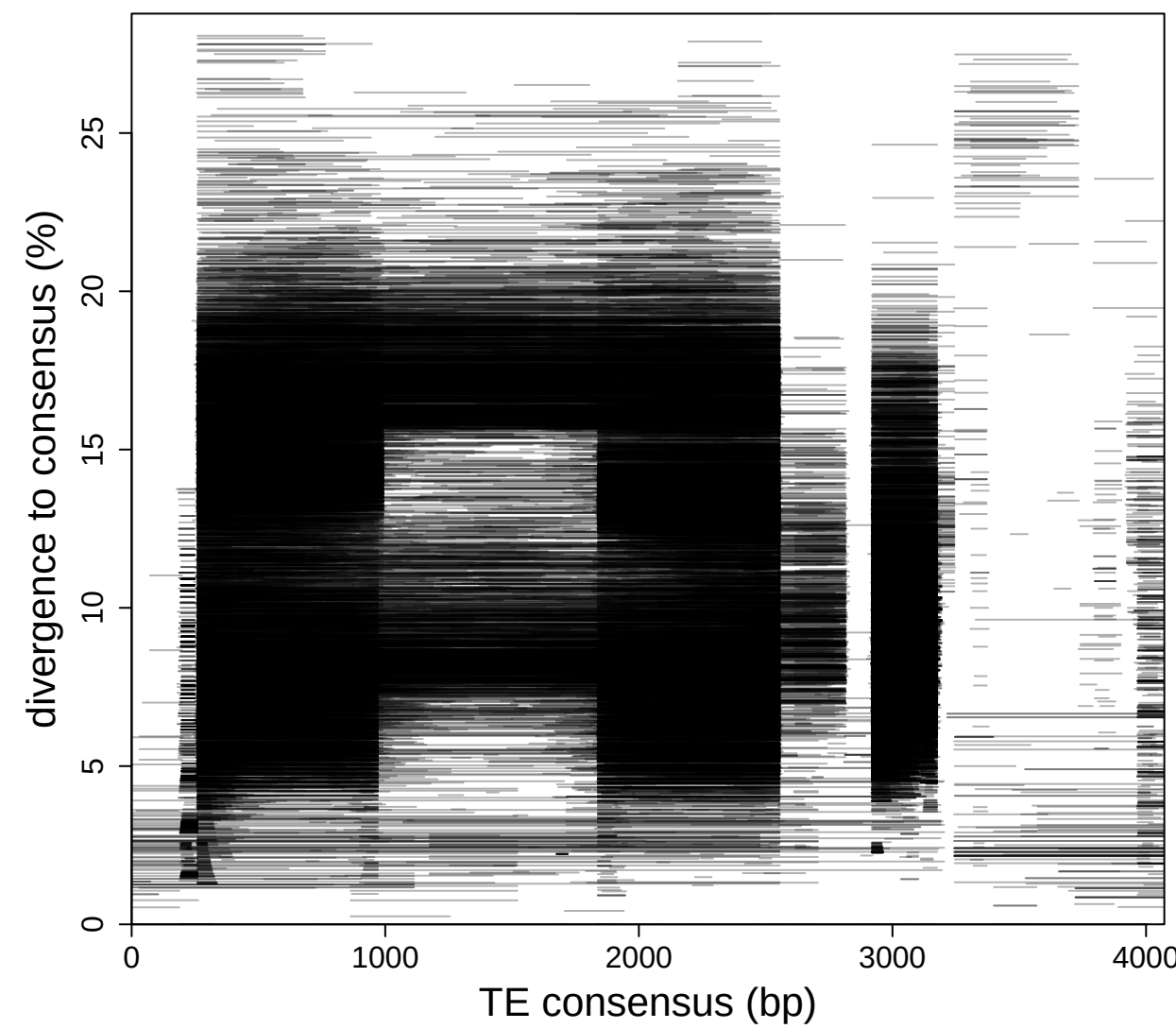
Start crop Point

End crop Point



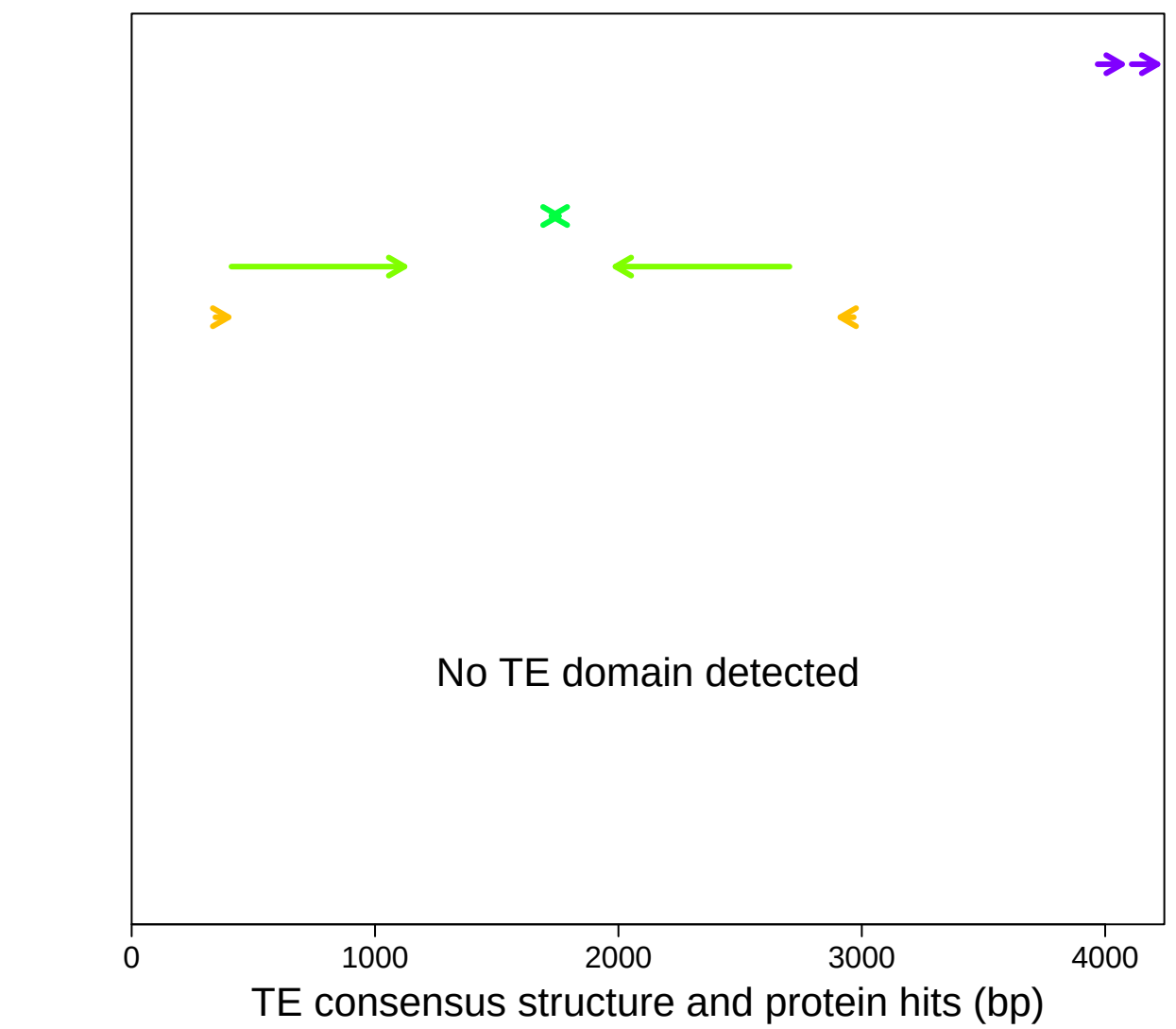
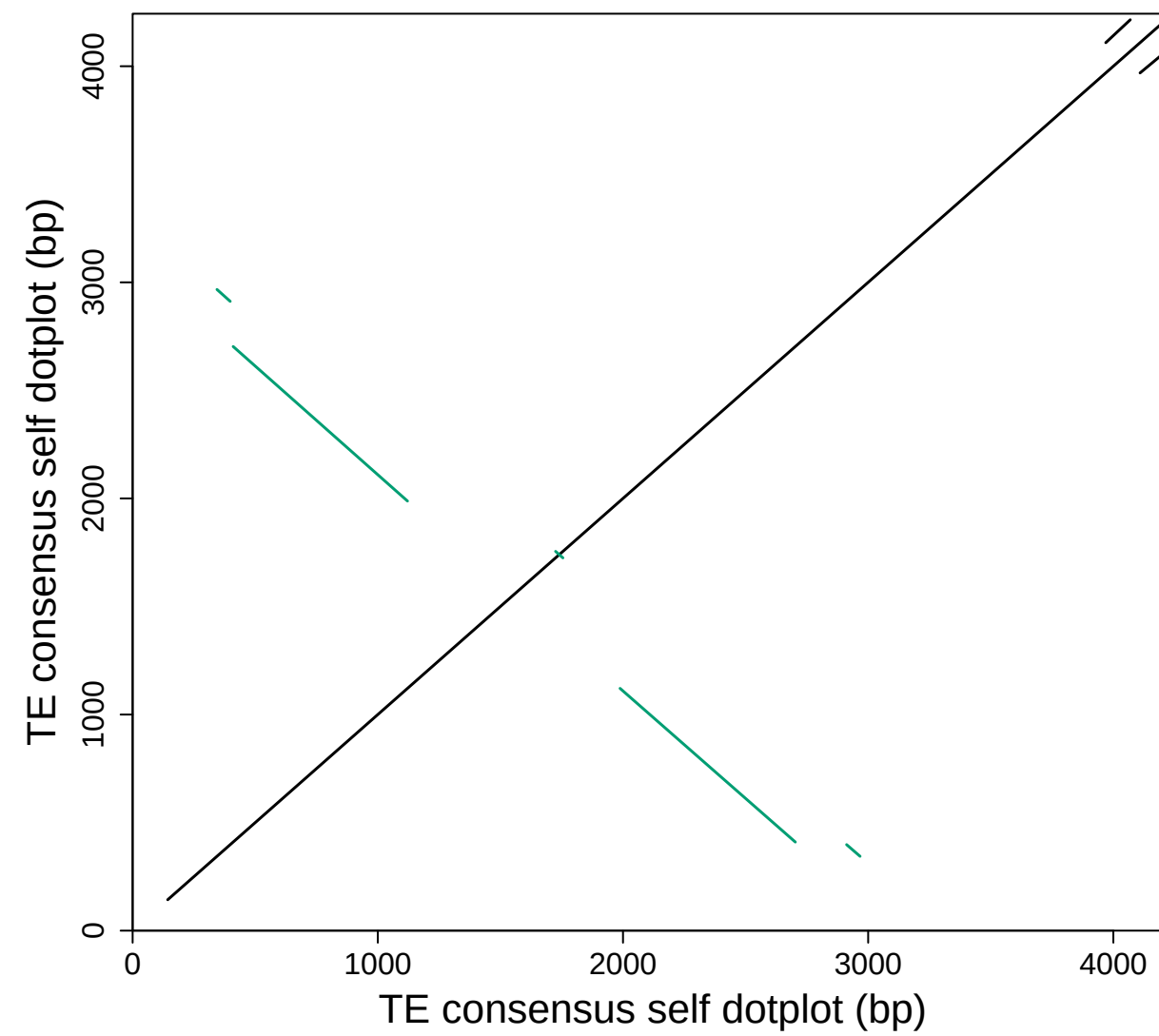
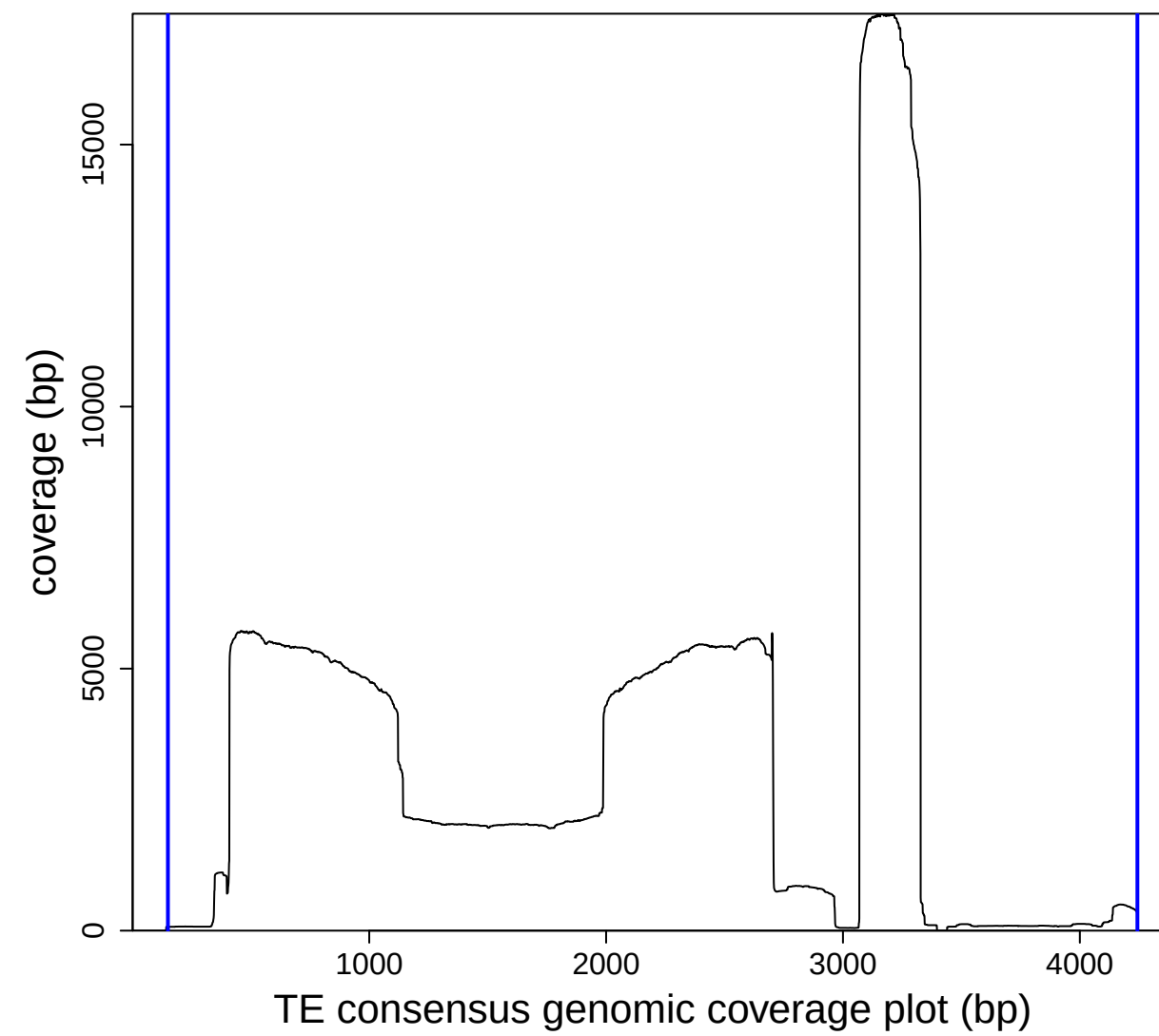
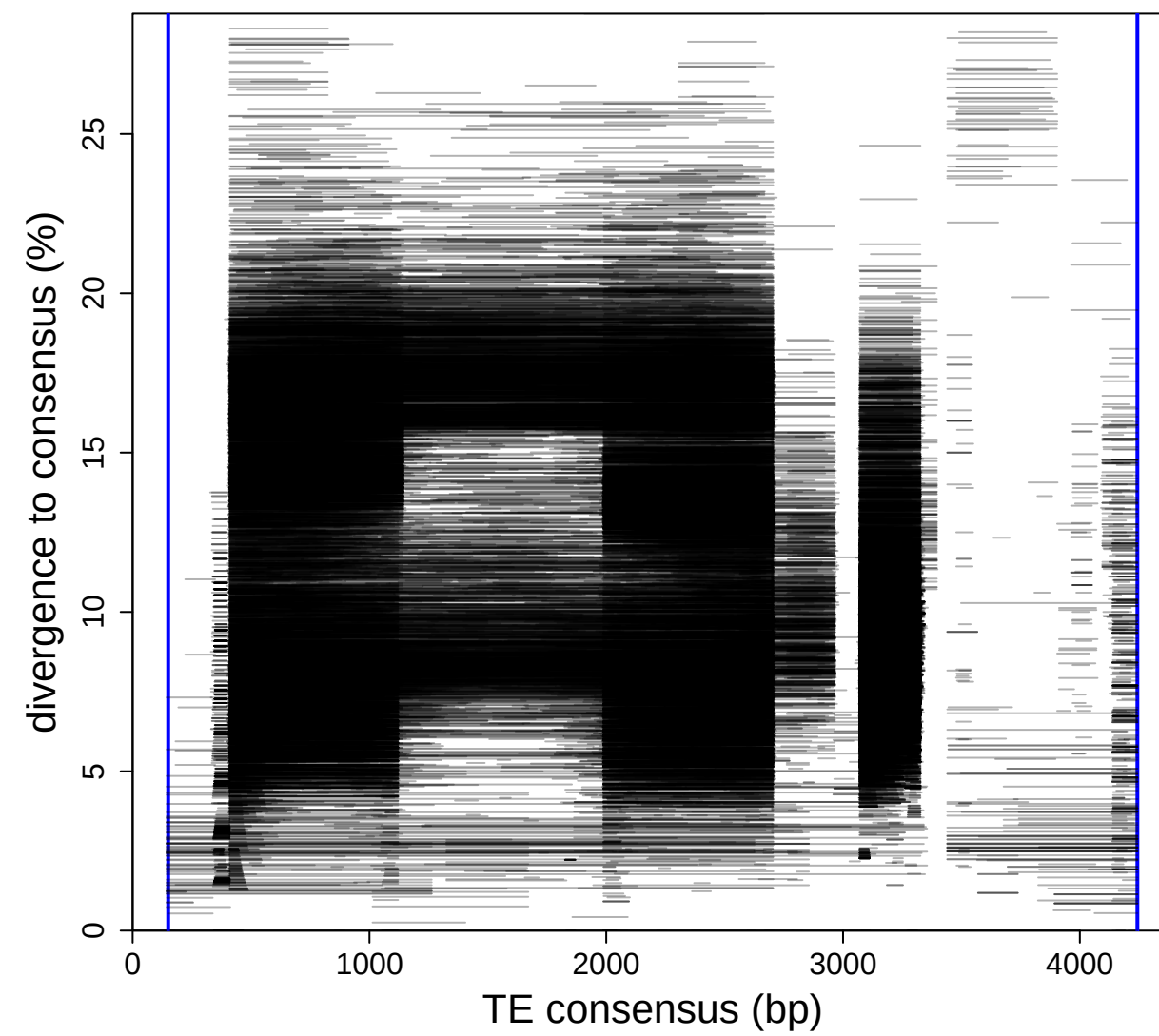
b.bed uf.bed g 1.bed fm 1.bed 0 0 bcln.fa aln.fa cl.fa gs.fa
size: 4072bp; fragments: 35083; full length: 0 (>=3664.8bp)

After TEtrimmer 4072 bp



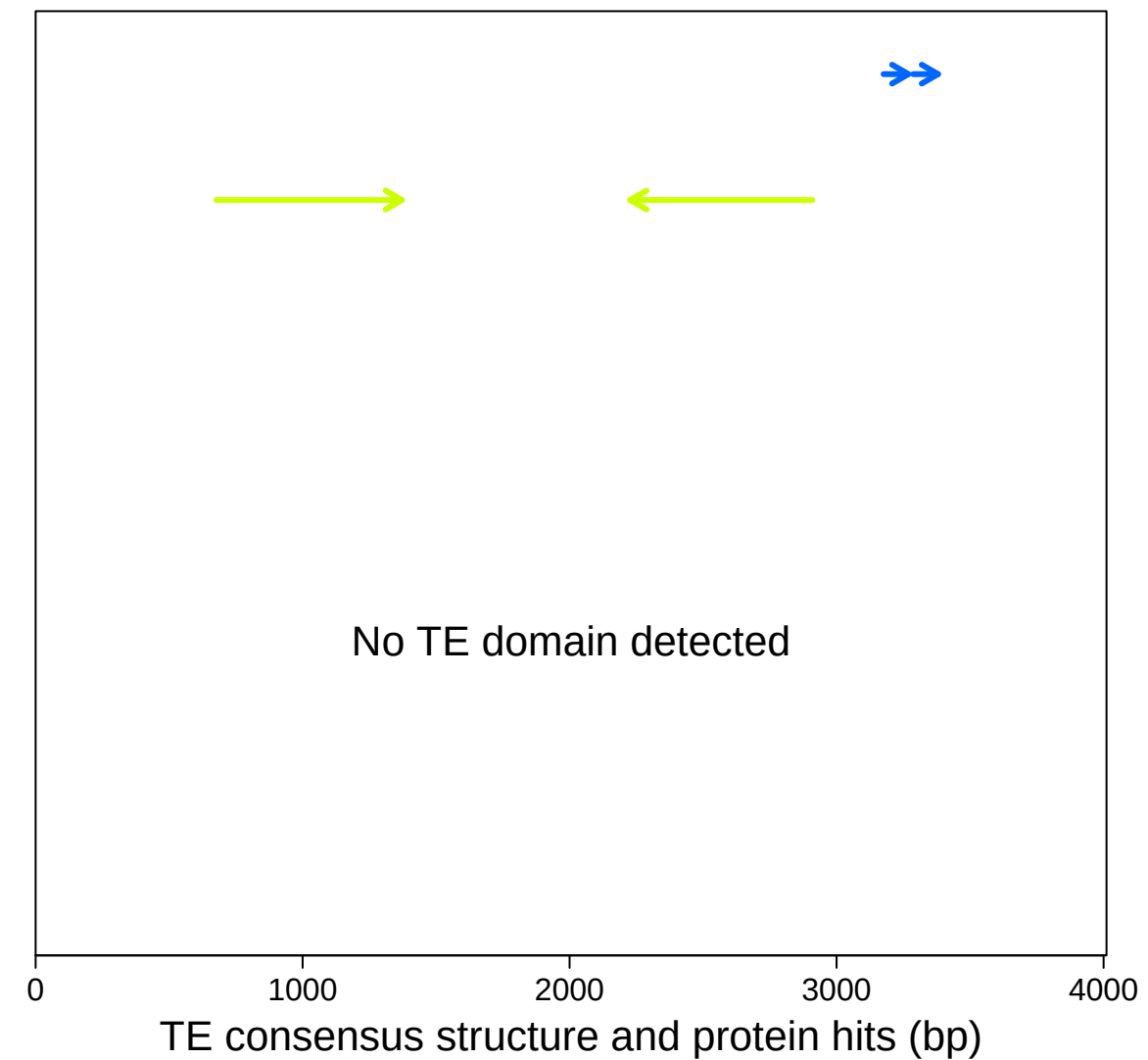
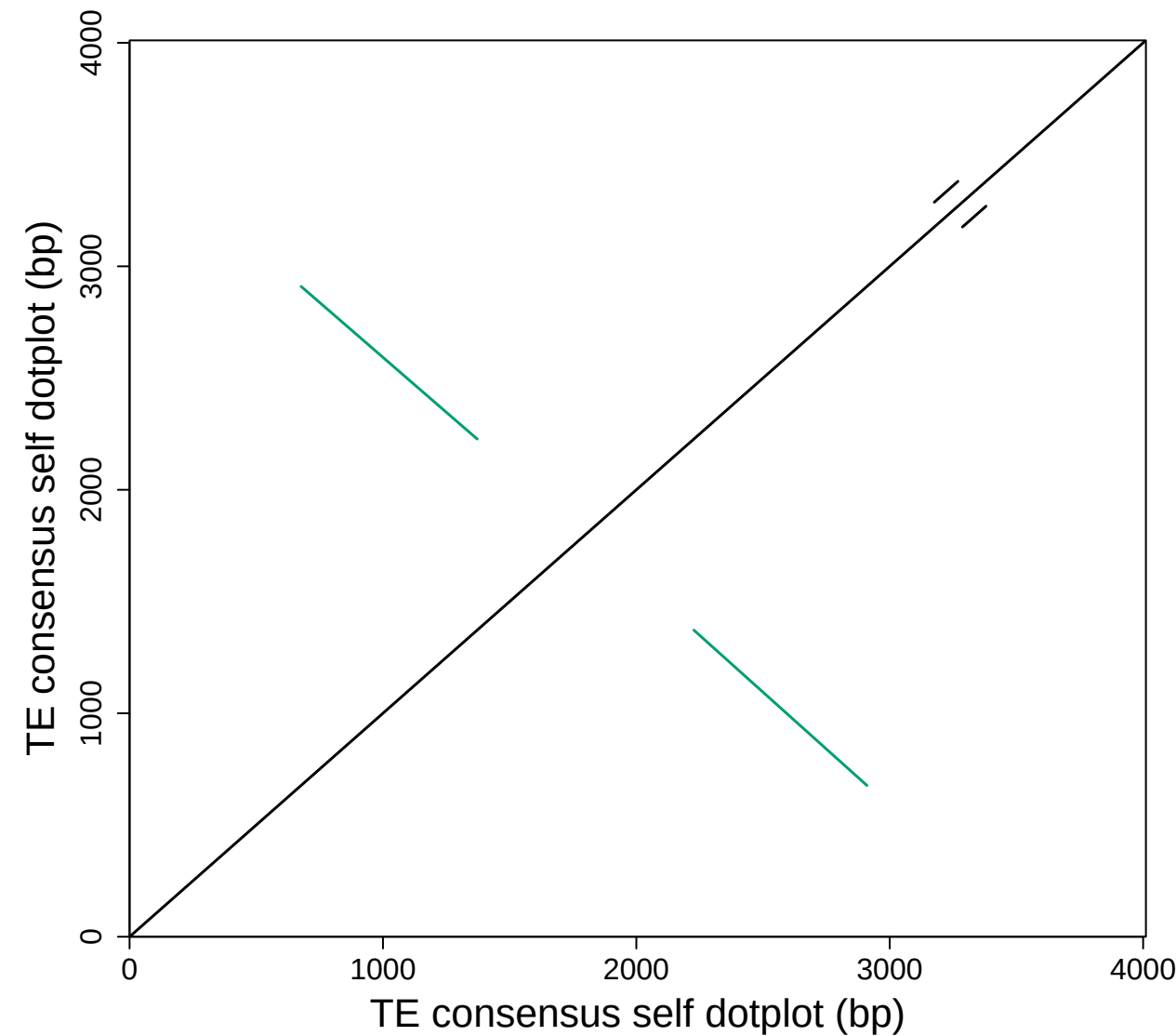
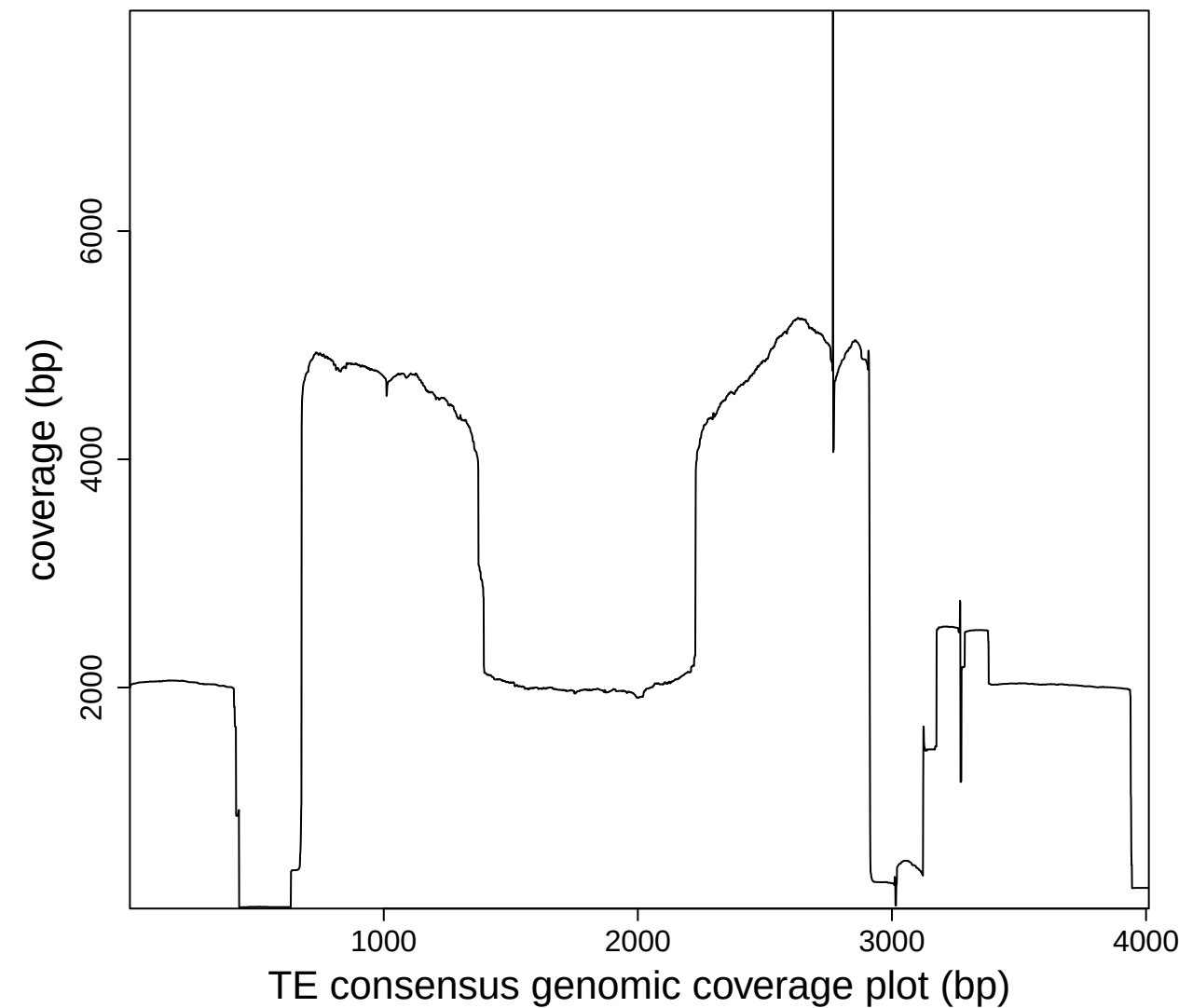
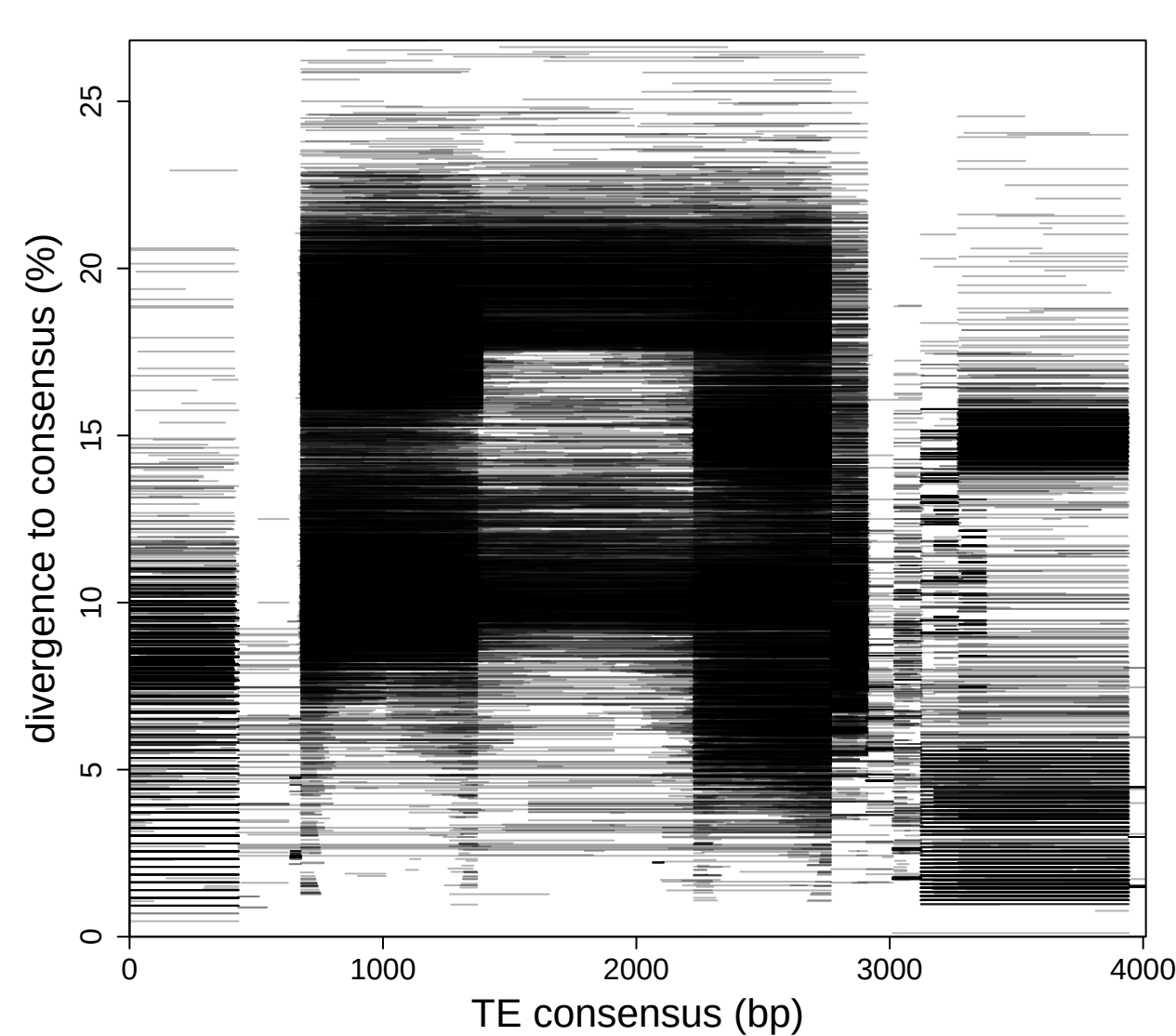
.fasta.b.bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa
size: 4393bp; fragments: 35053; full length: 0 (>=3953.7bp)

After TEtrimmer Extended plot Blue lines are boundaries

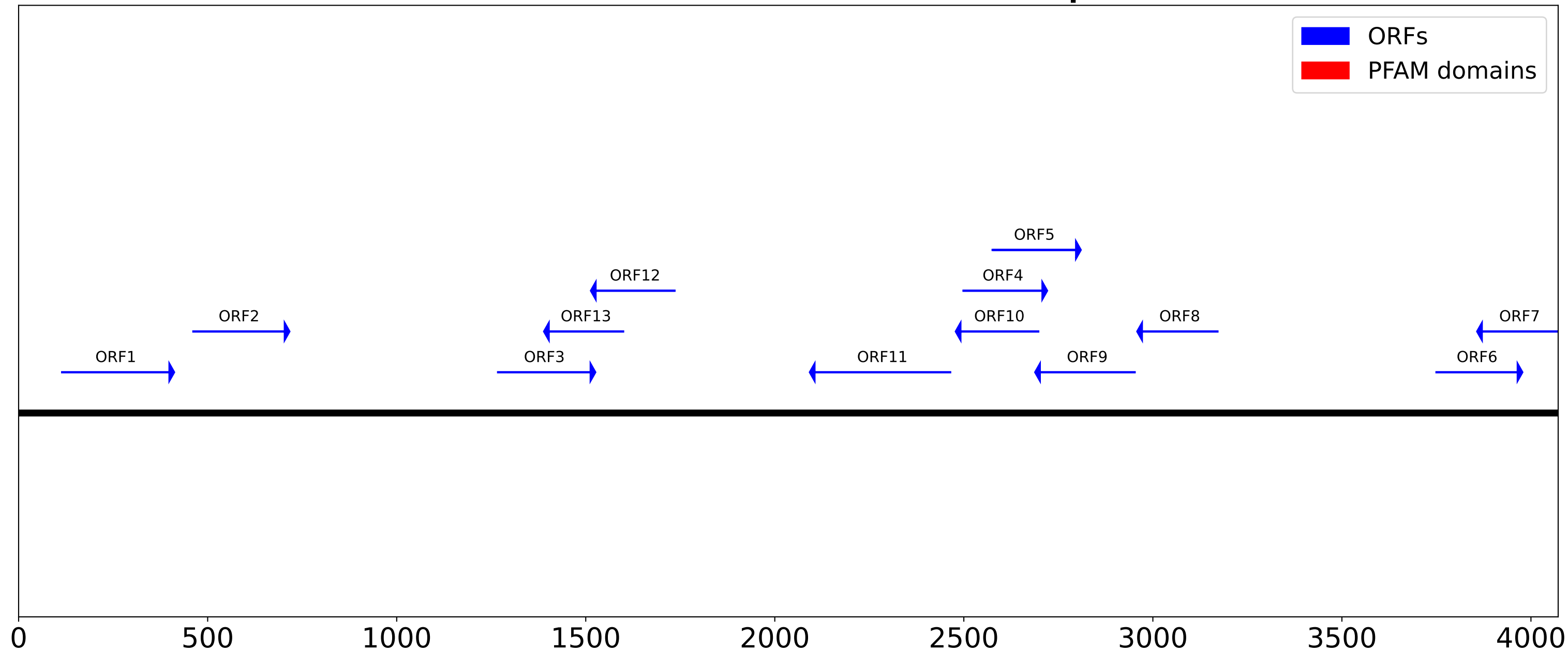


TE: ltr_1_family_557
size: 4011bp; fragments: 23502; full length: 0 (>=3609.9bp)

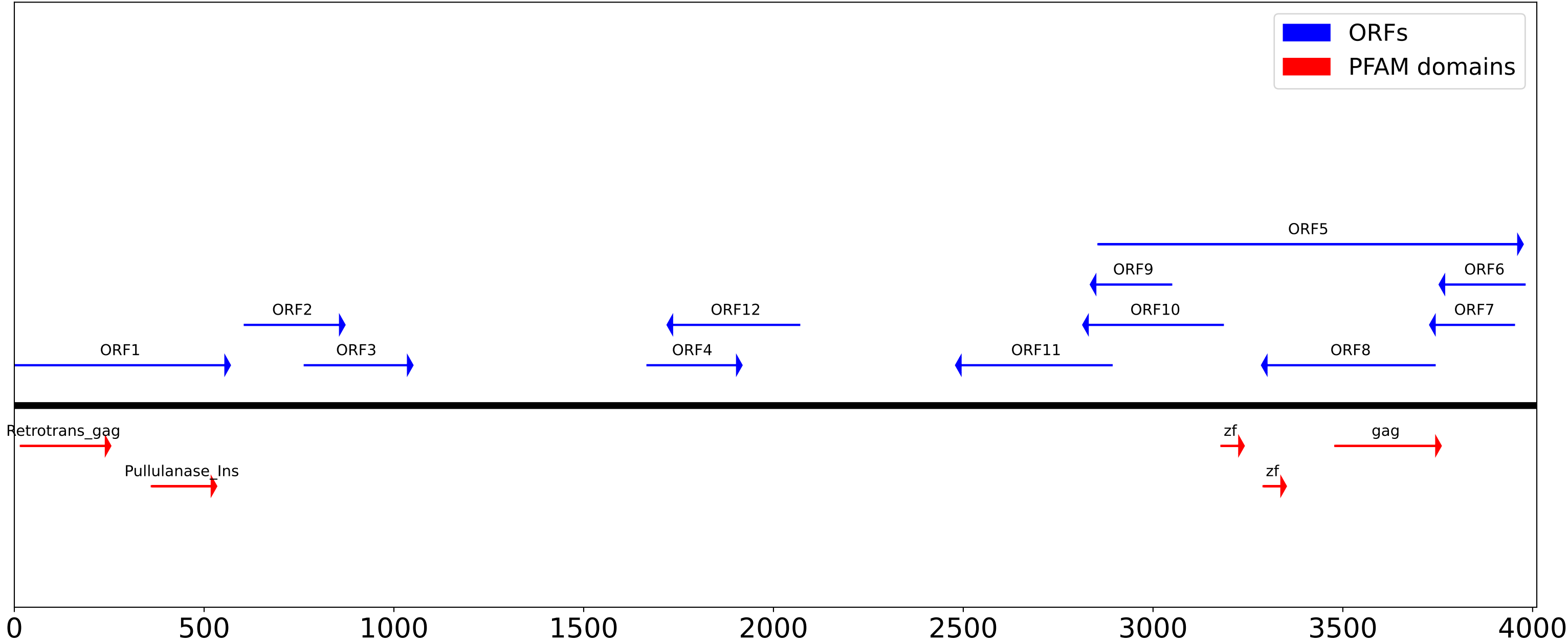
Before TEtrimmer 4011 bp



After TEtrimmer ORF and PFAM domain plot



Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

