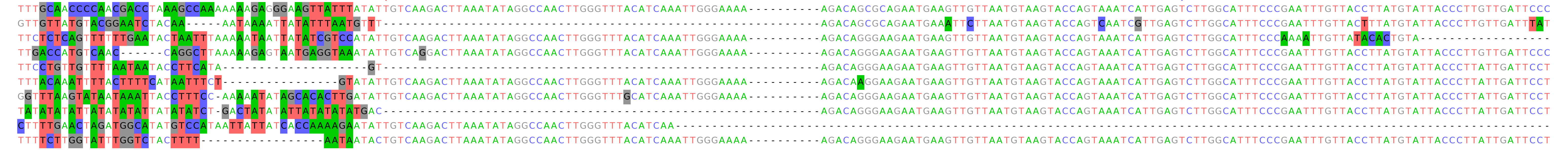


Start Crop Point

End Crop Point

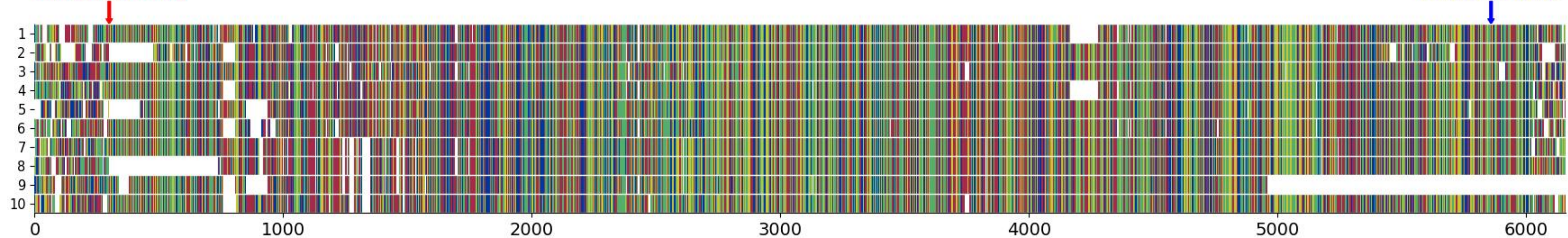


MSA length = 5560



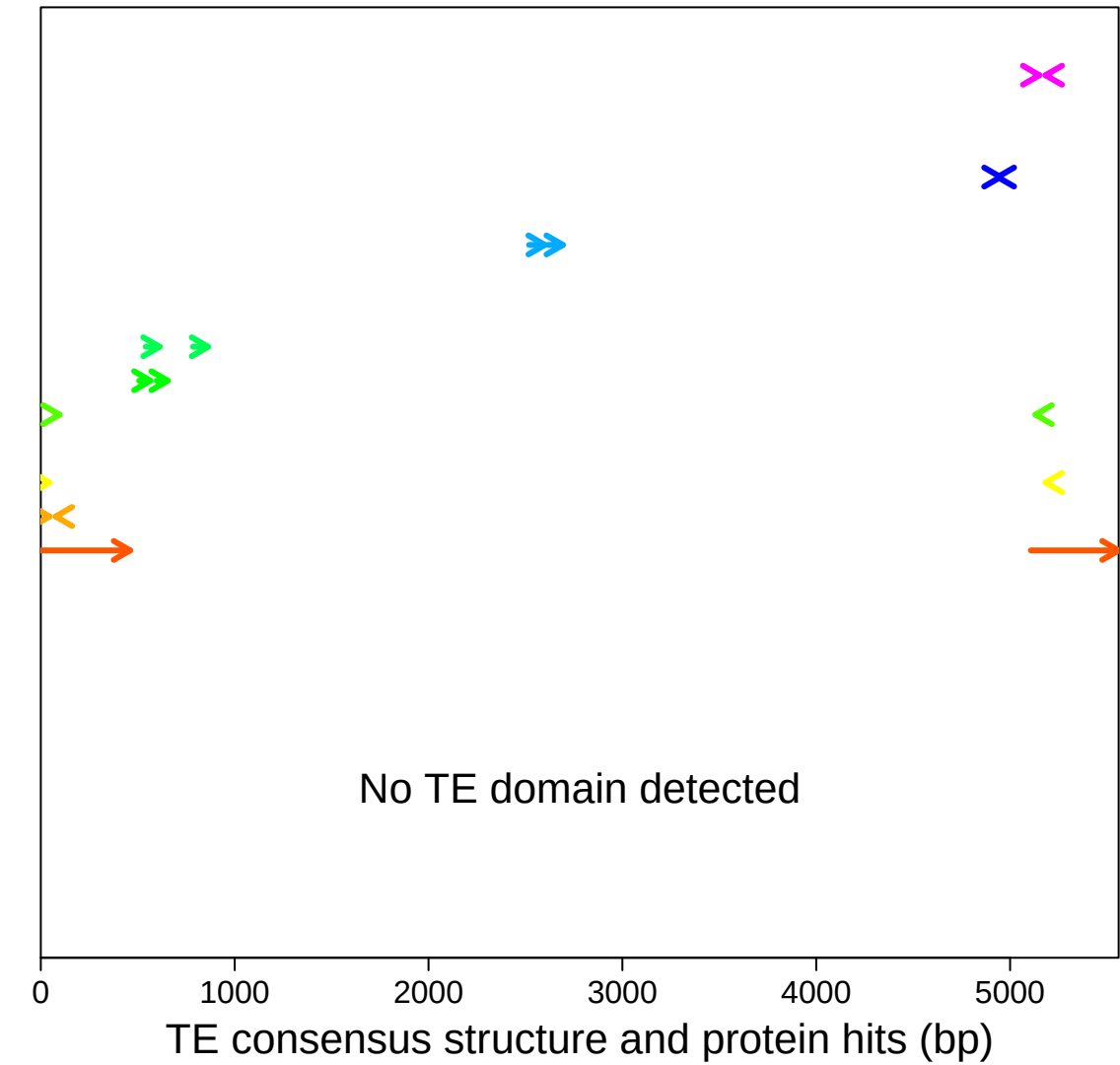
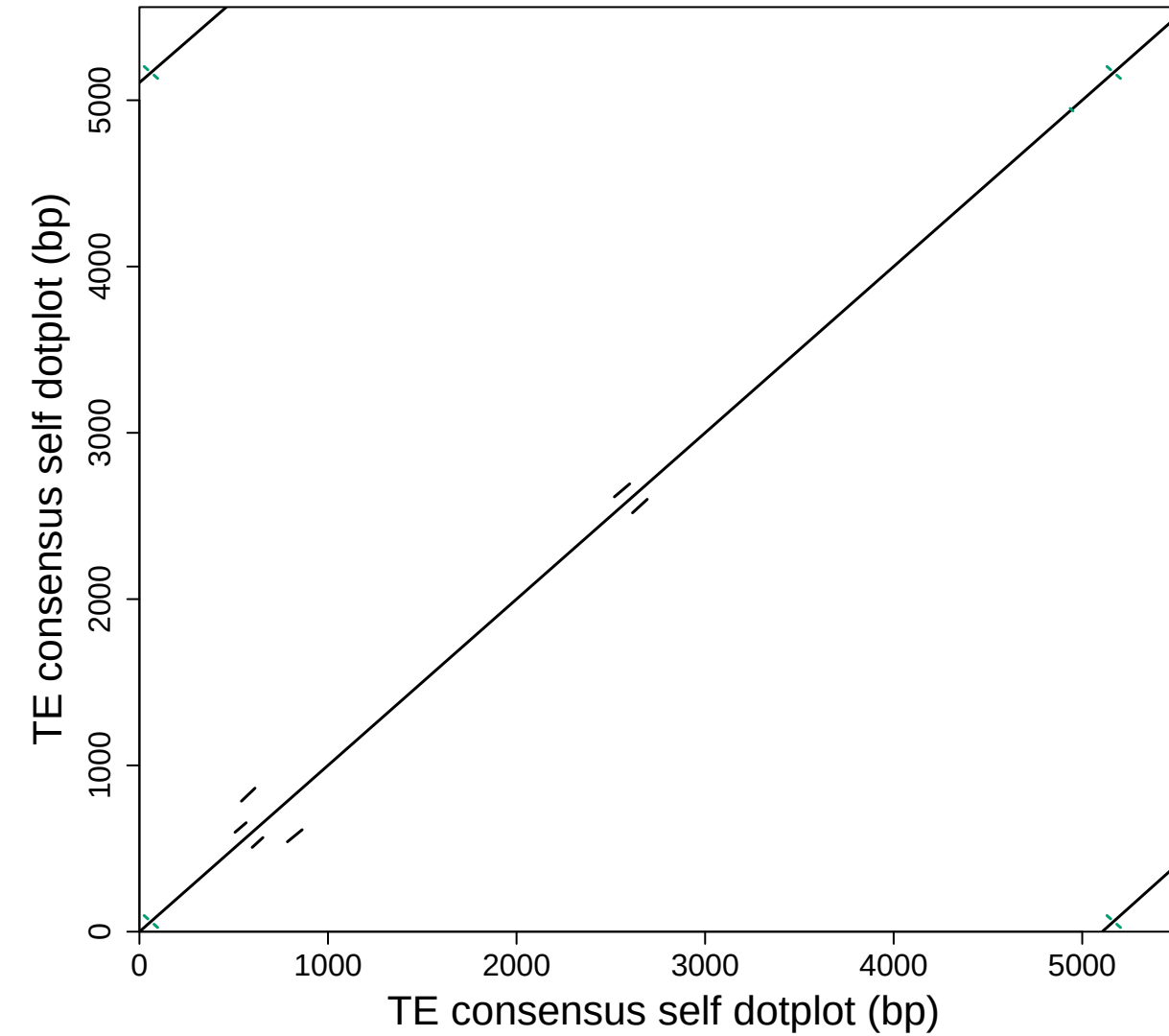
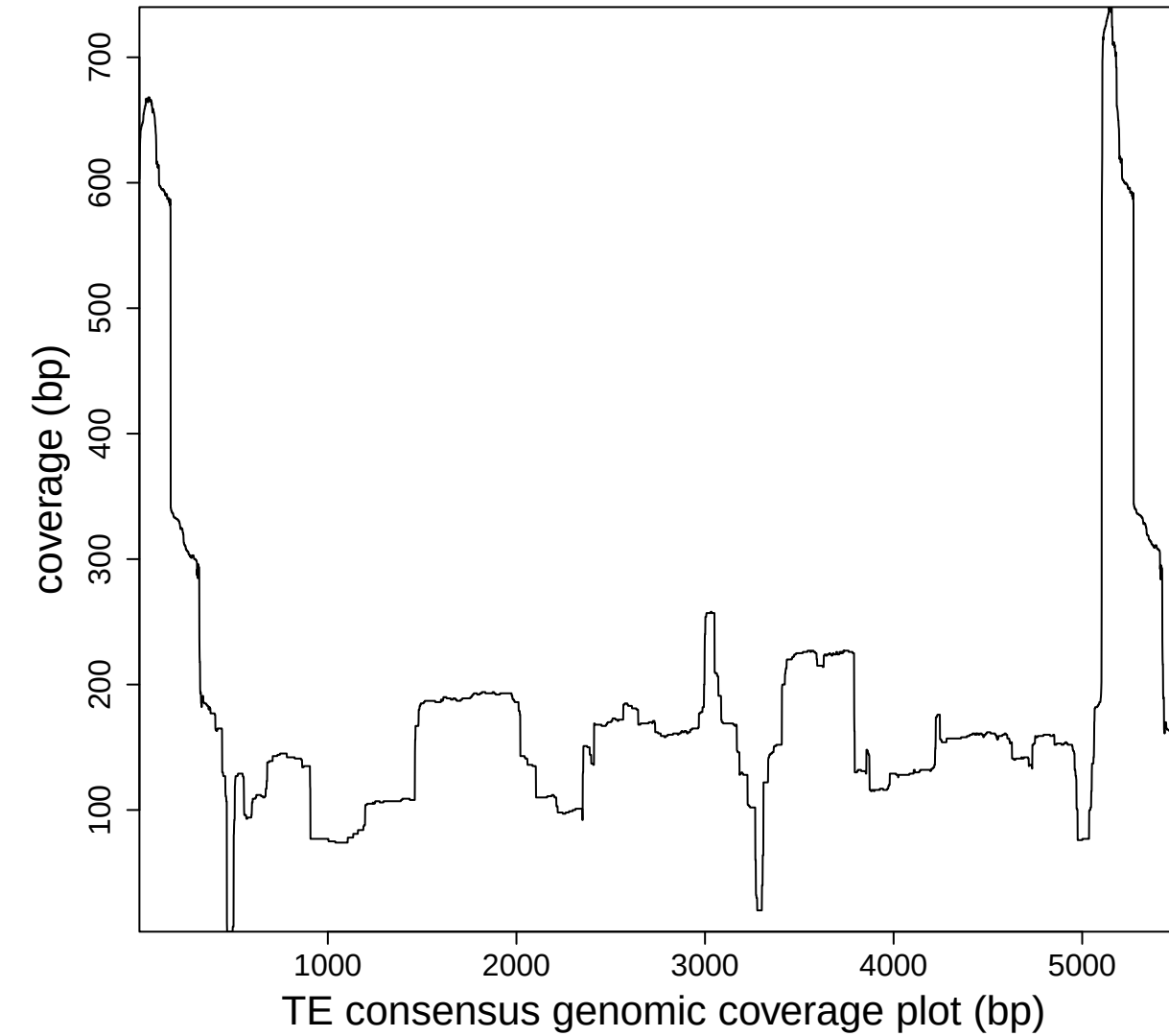
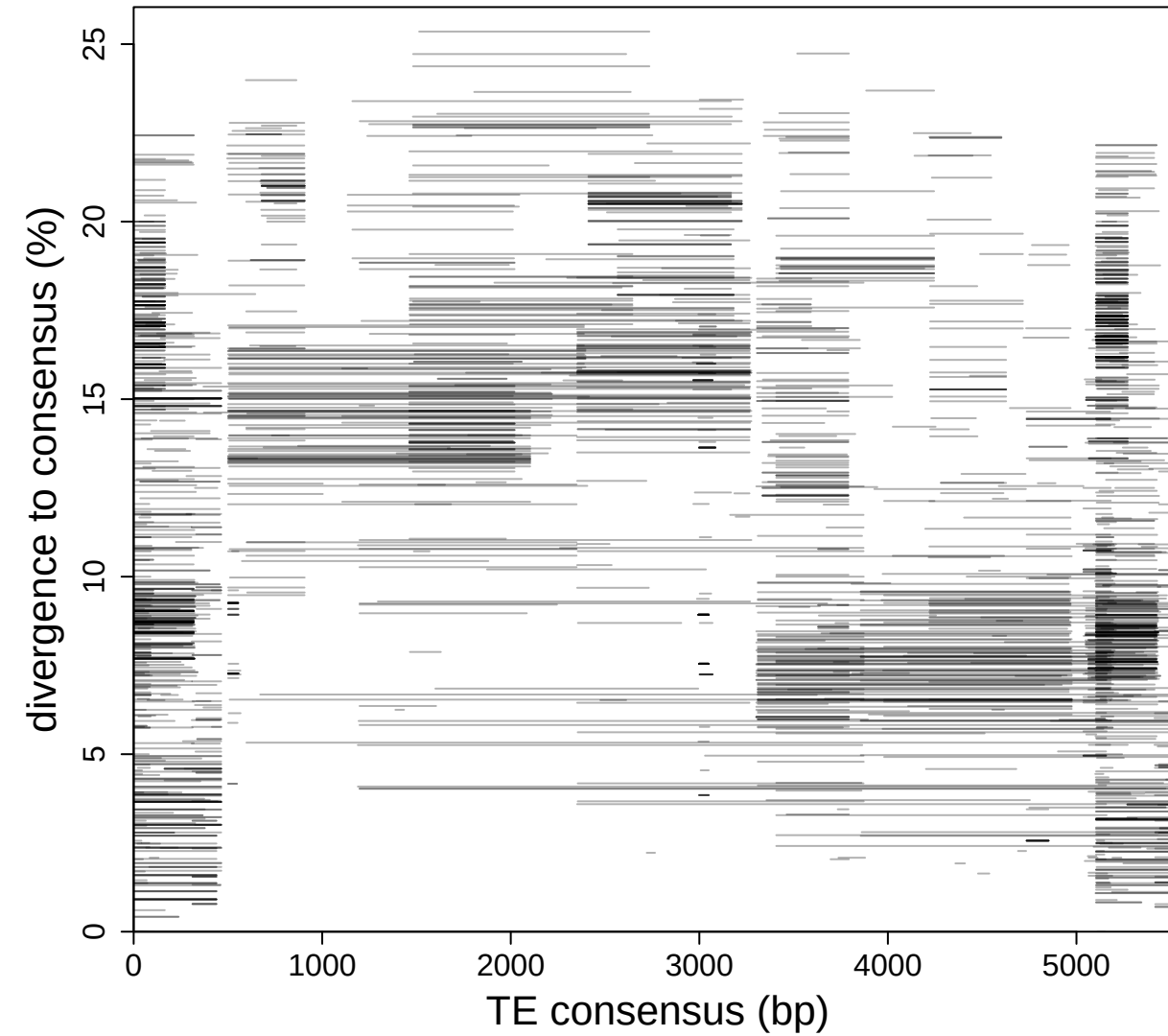
Start crop Point

End crop Point



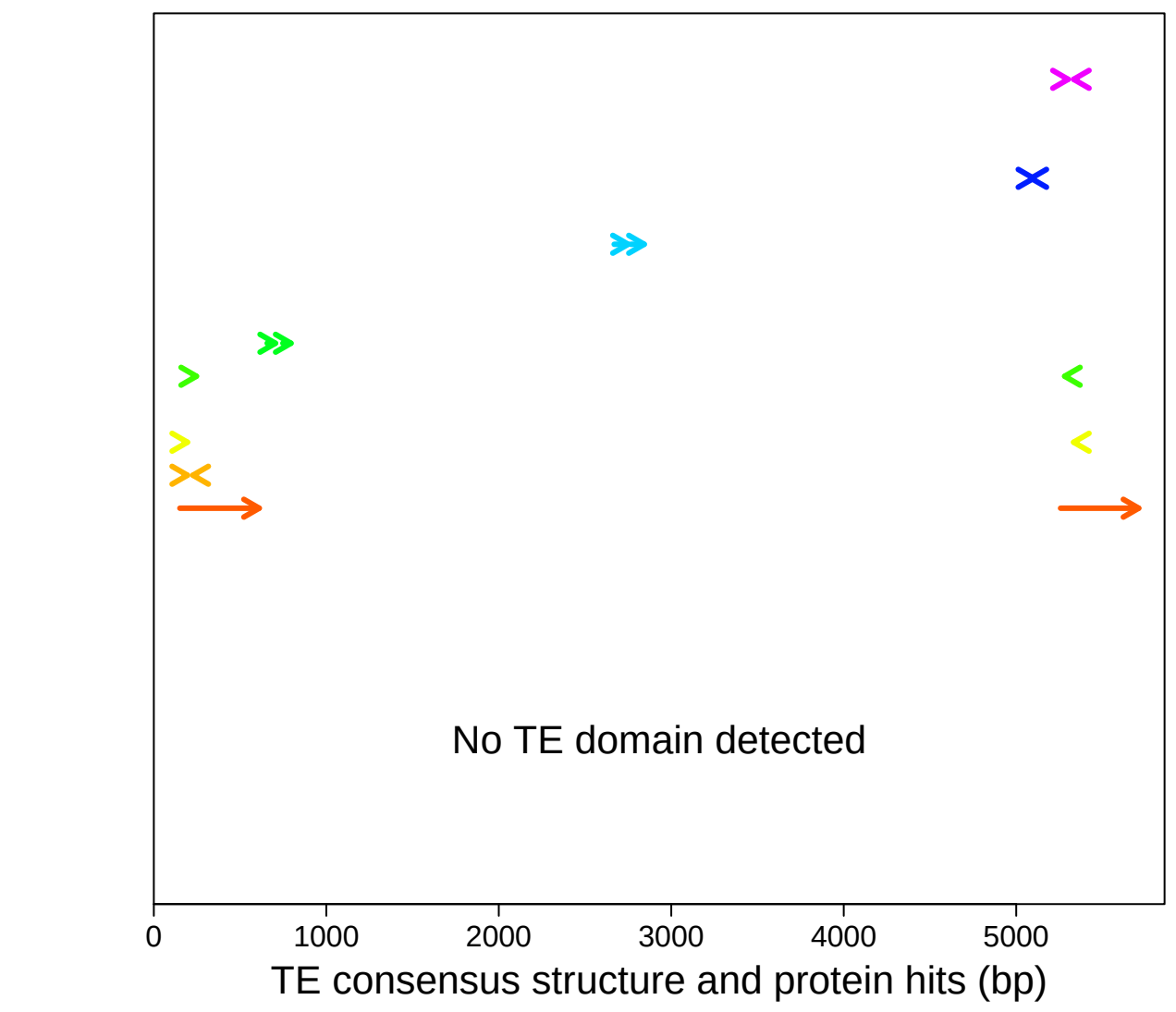
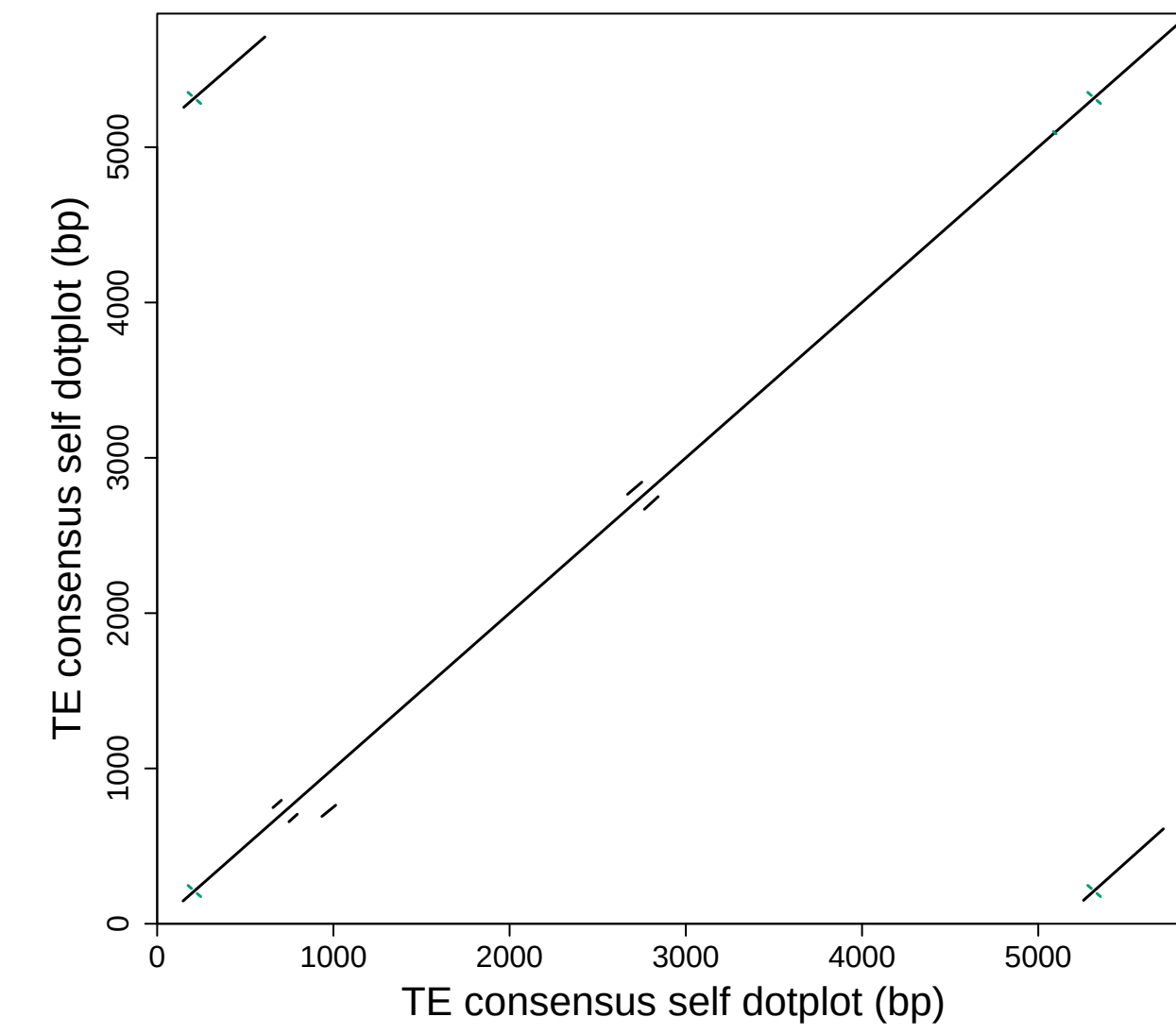
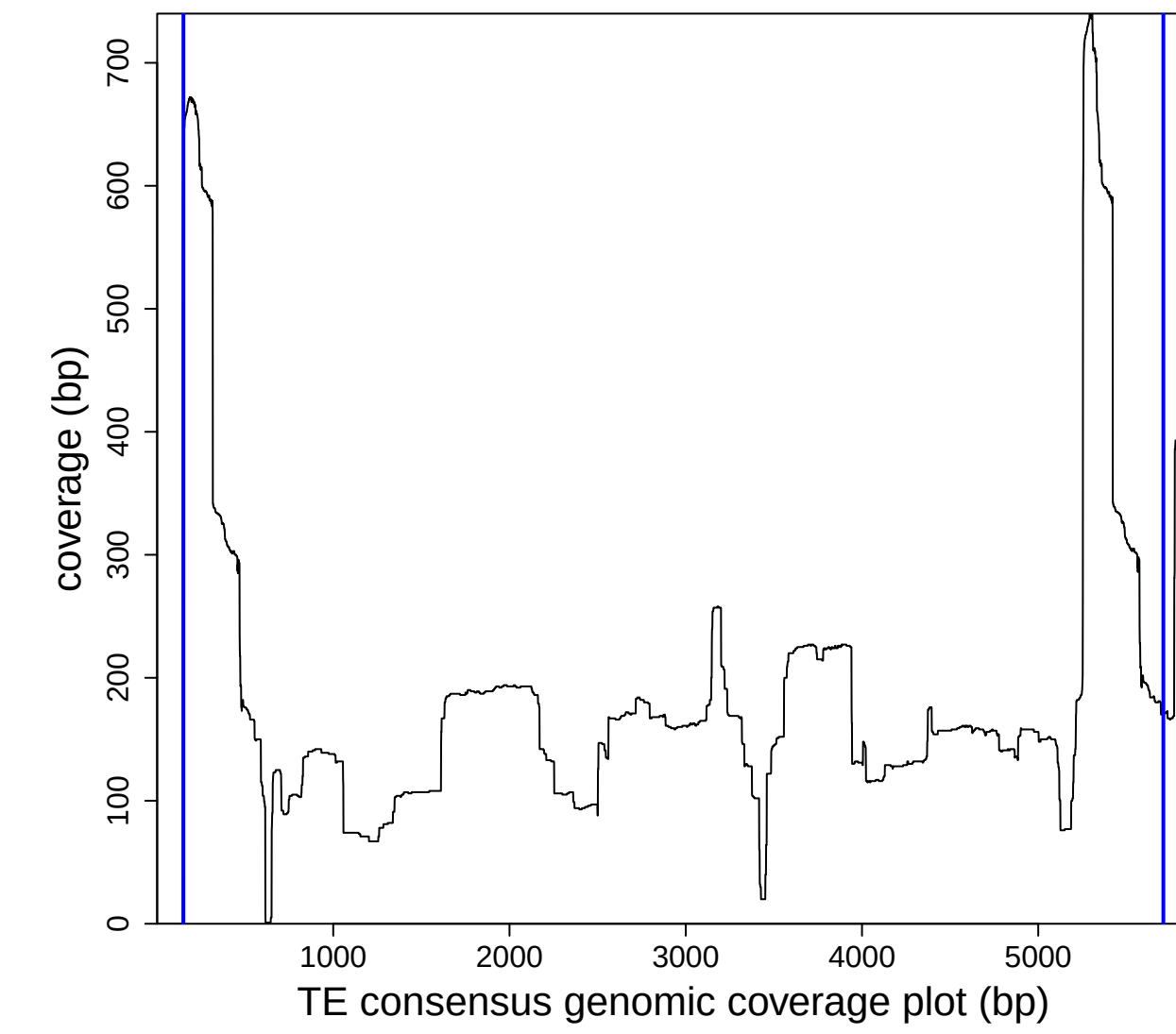
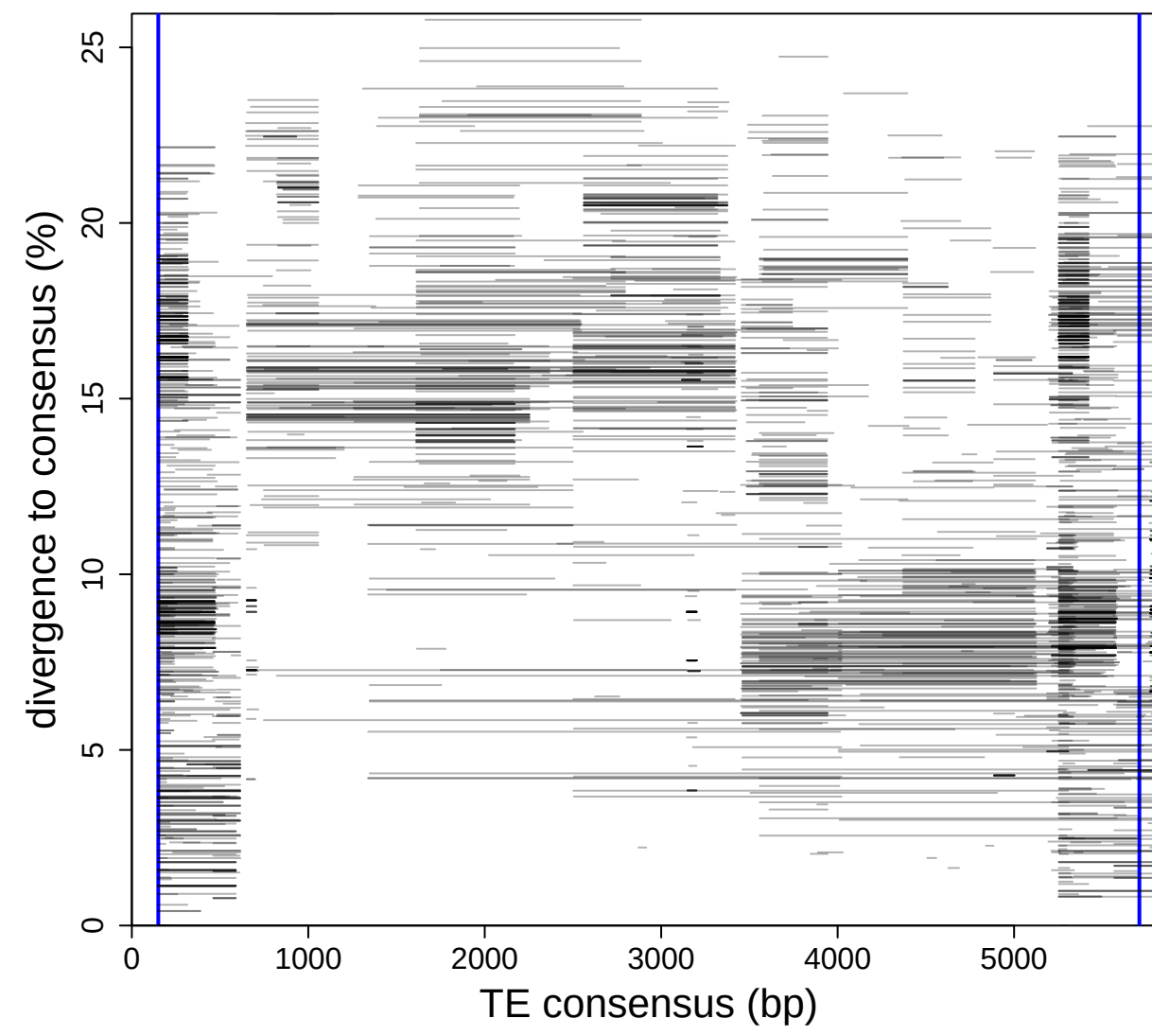
bed uf.bed g 4.bed fm 1.bed 0 0 bcln.fa aln.fa cl.fa gs.fa ce
size: 5560bp; fragments: 2449; full length: 0 (>=5004bp)

After TEtrimmer 5560 bp



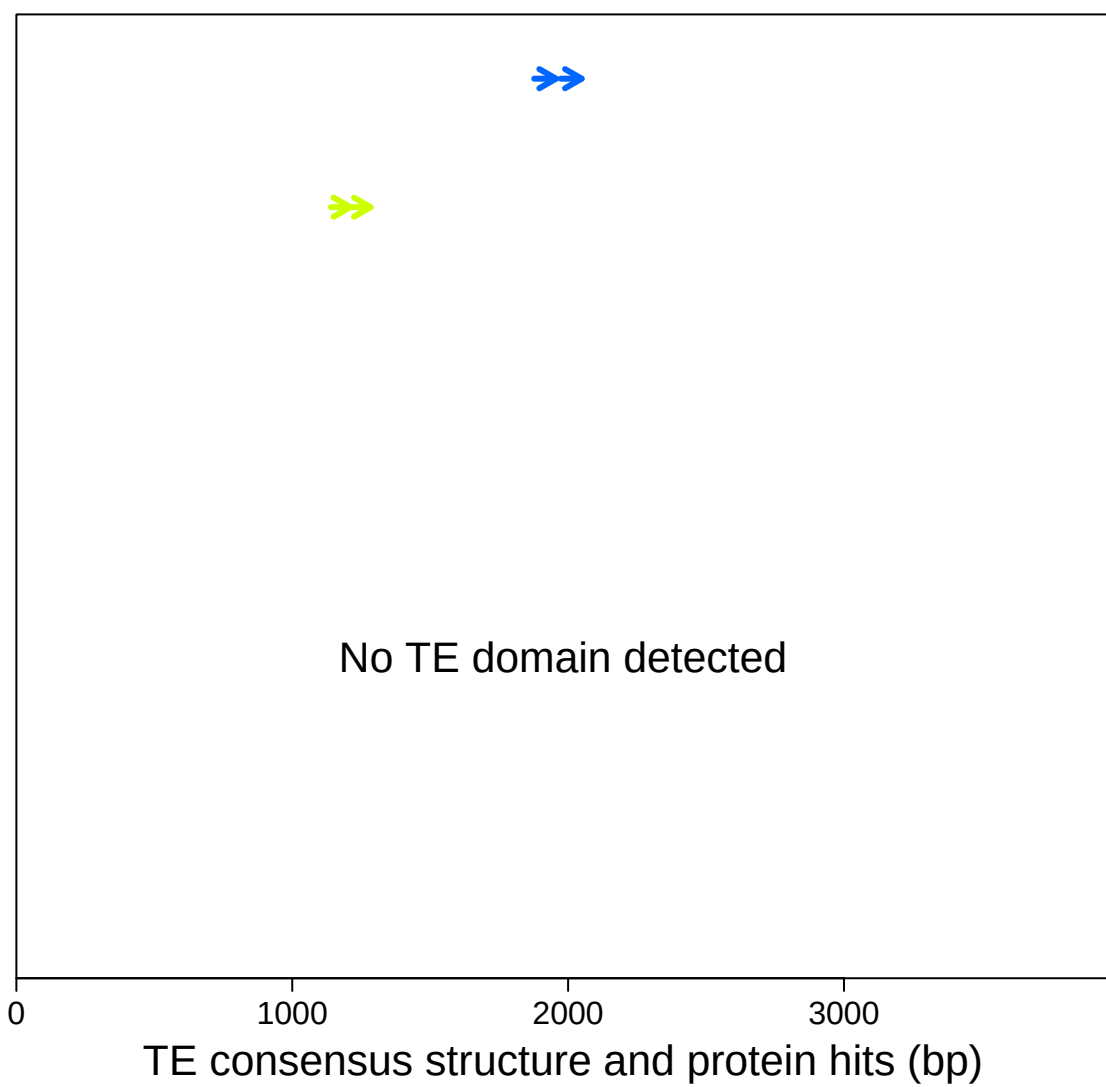
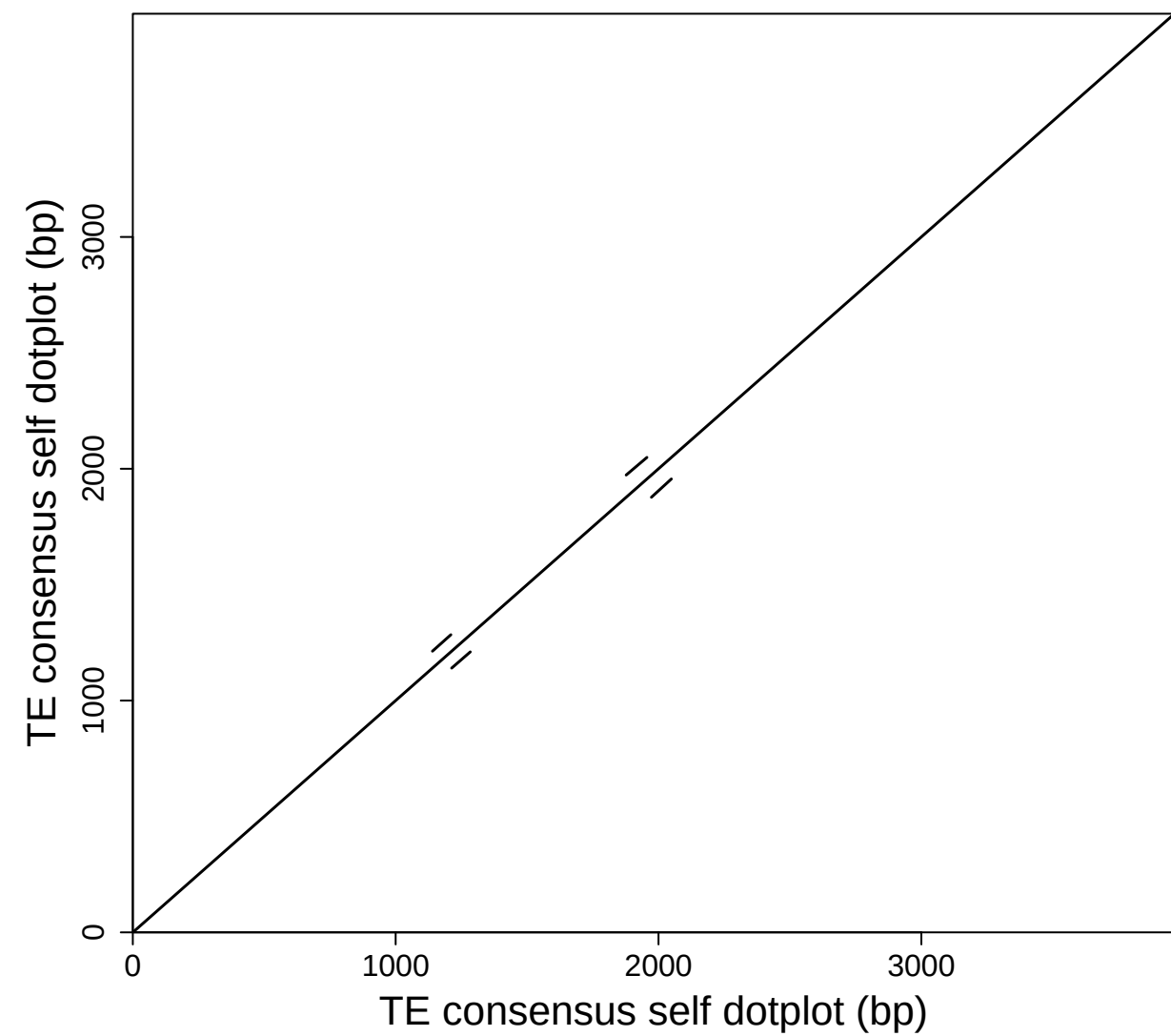
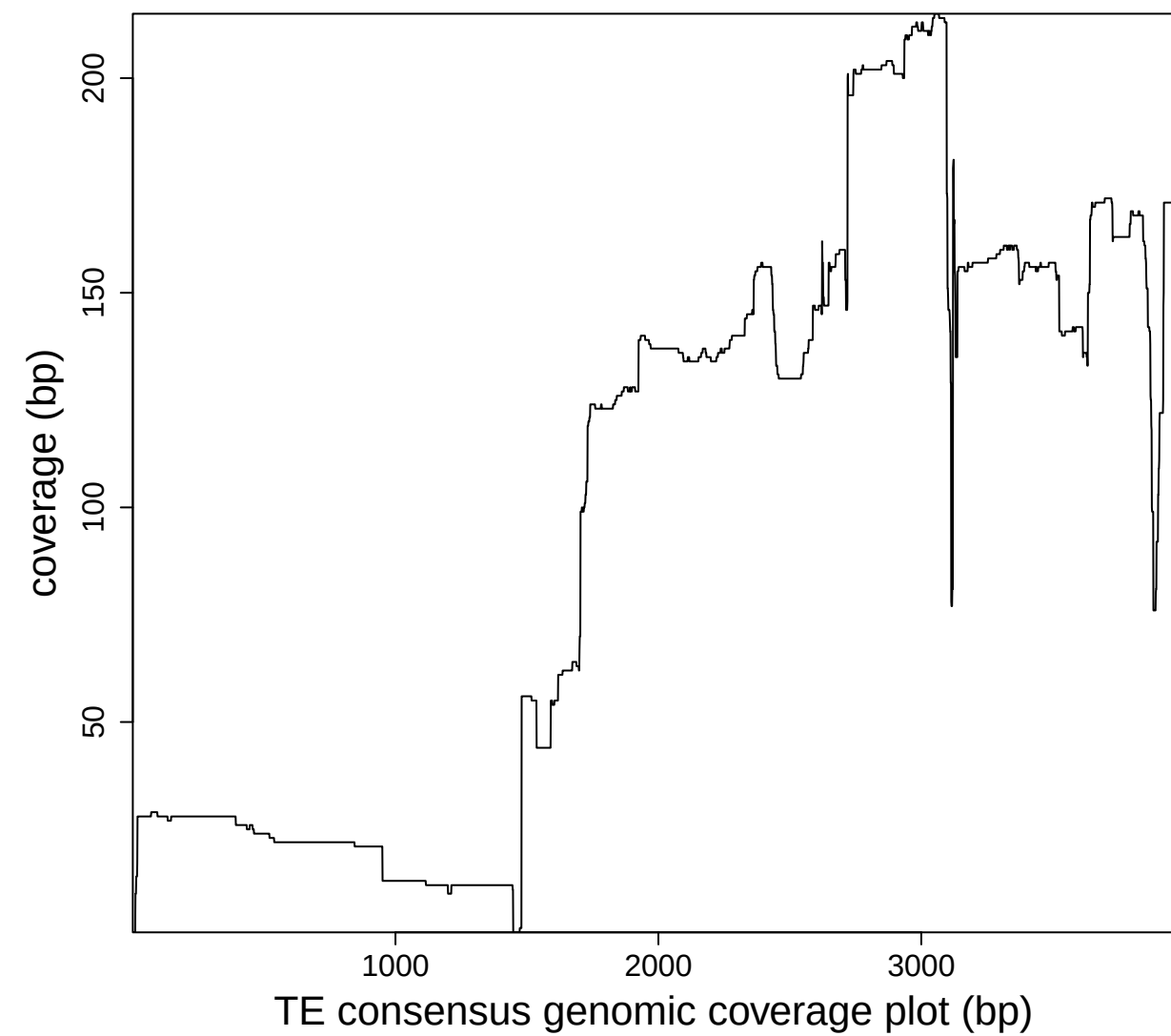
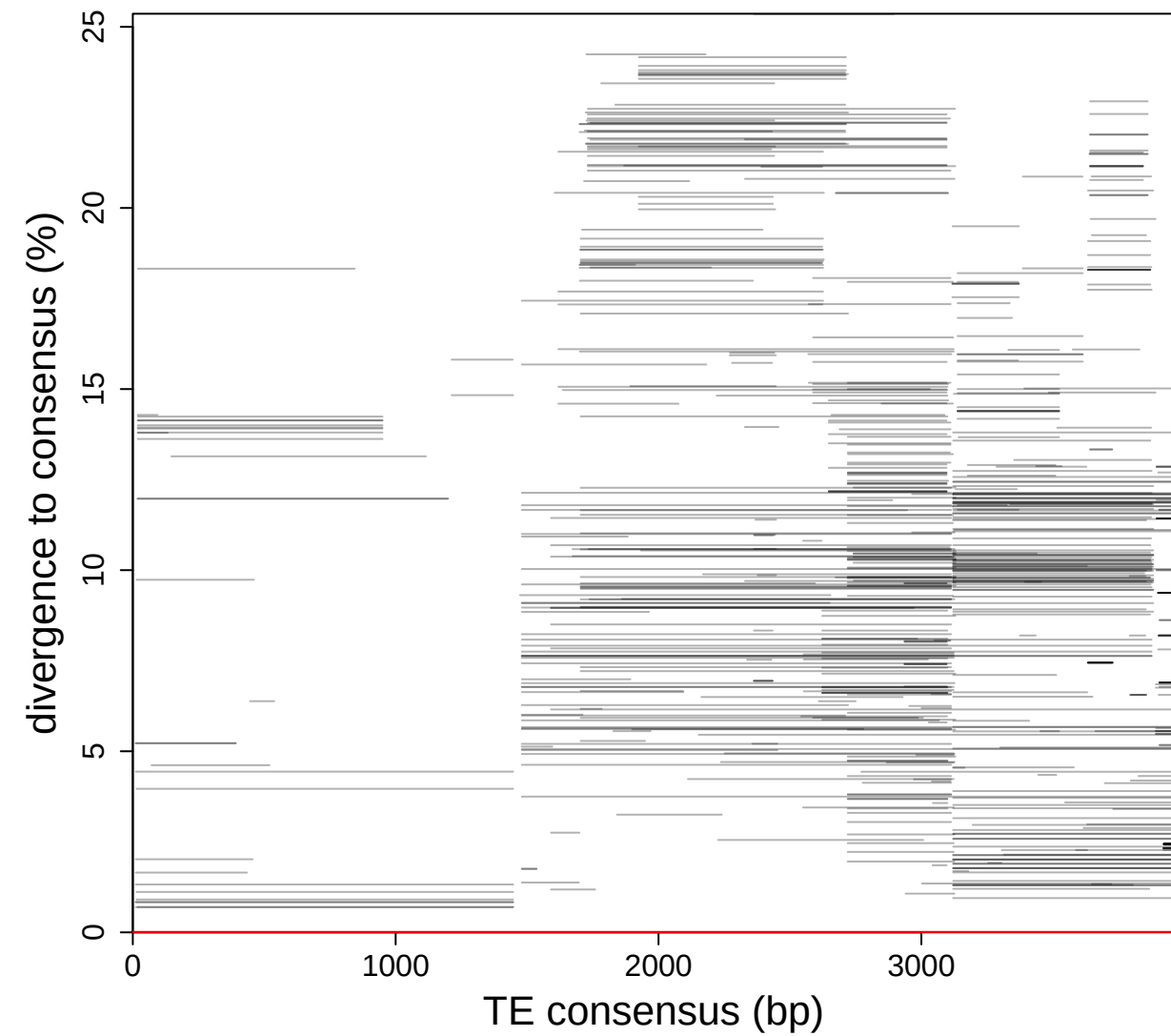
y_1036.fasta.b.bed_uf.bed_g_4.bed_fm_1.bed_0_0_bcln.fa_aln.f
size: 5860bp; fragments: 2698; full length: 0 (>=5274bp)

After TEtrimmer Extended plot Blue lines are boundaries

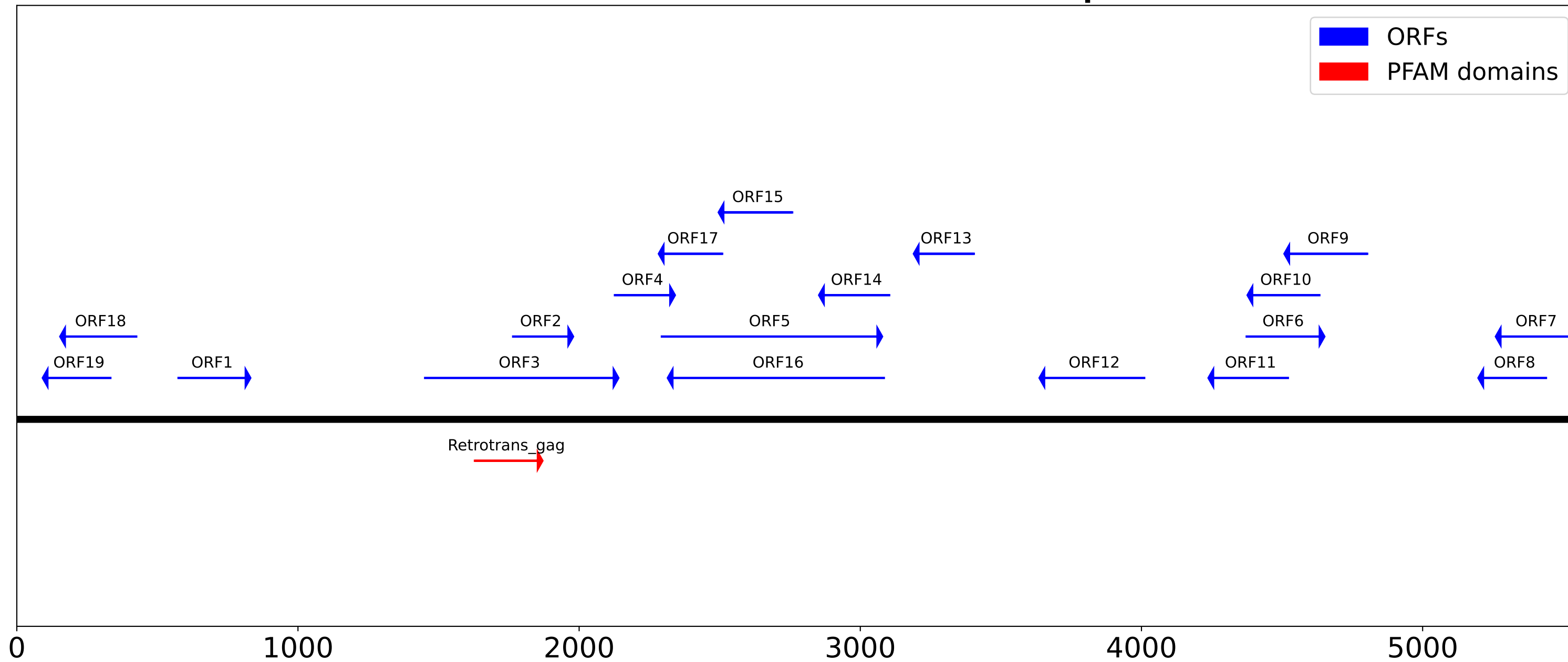


TE: ltr_1_family_1036
size: 3963bp; fragments: 673; full length: 1 (>=3566.7bp)

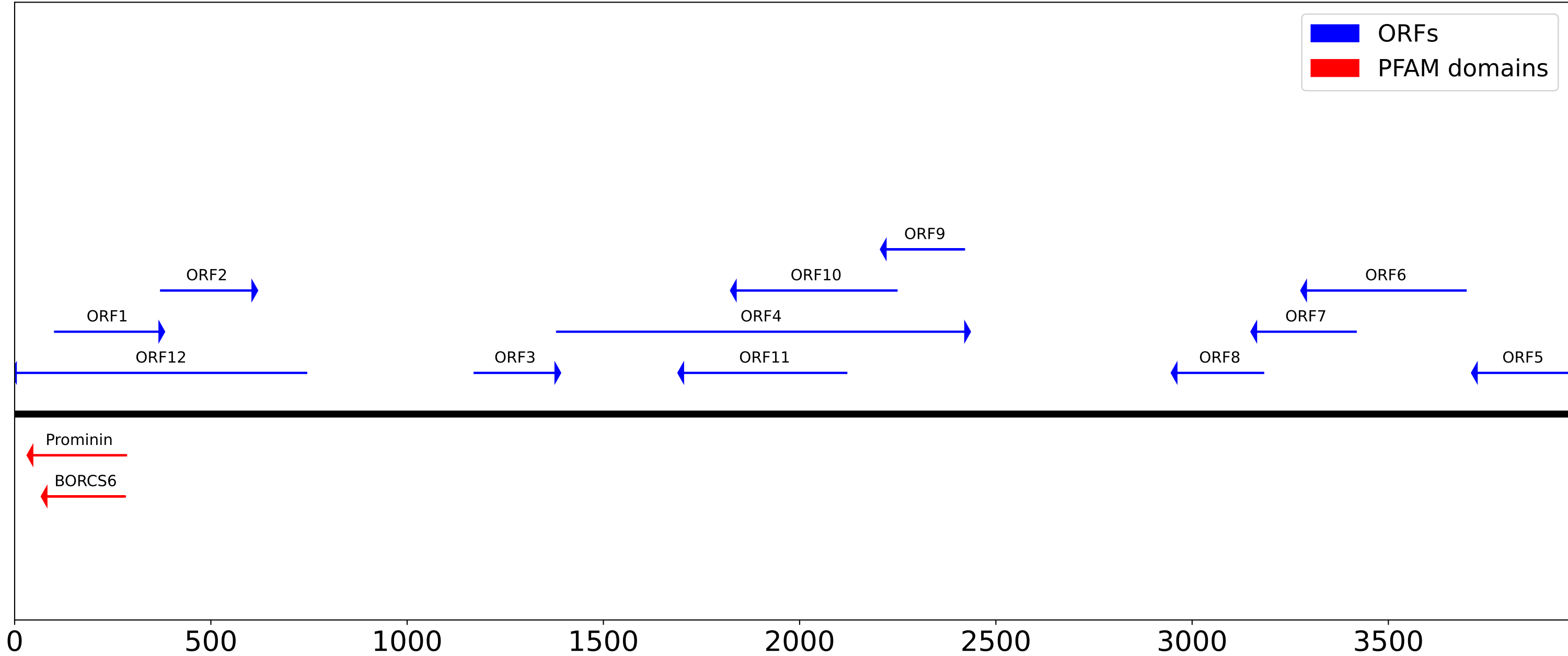
Before TEtrimmer 3963 bp



After TEtrimmer ORF and PFAM domain plot



Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

