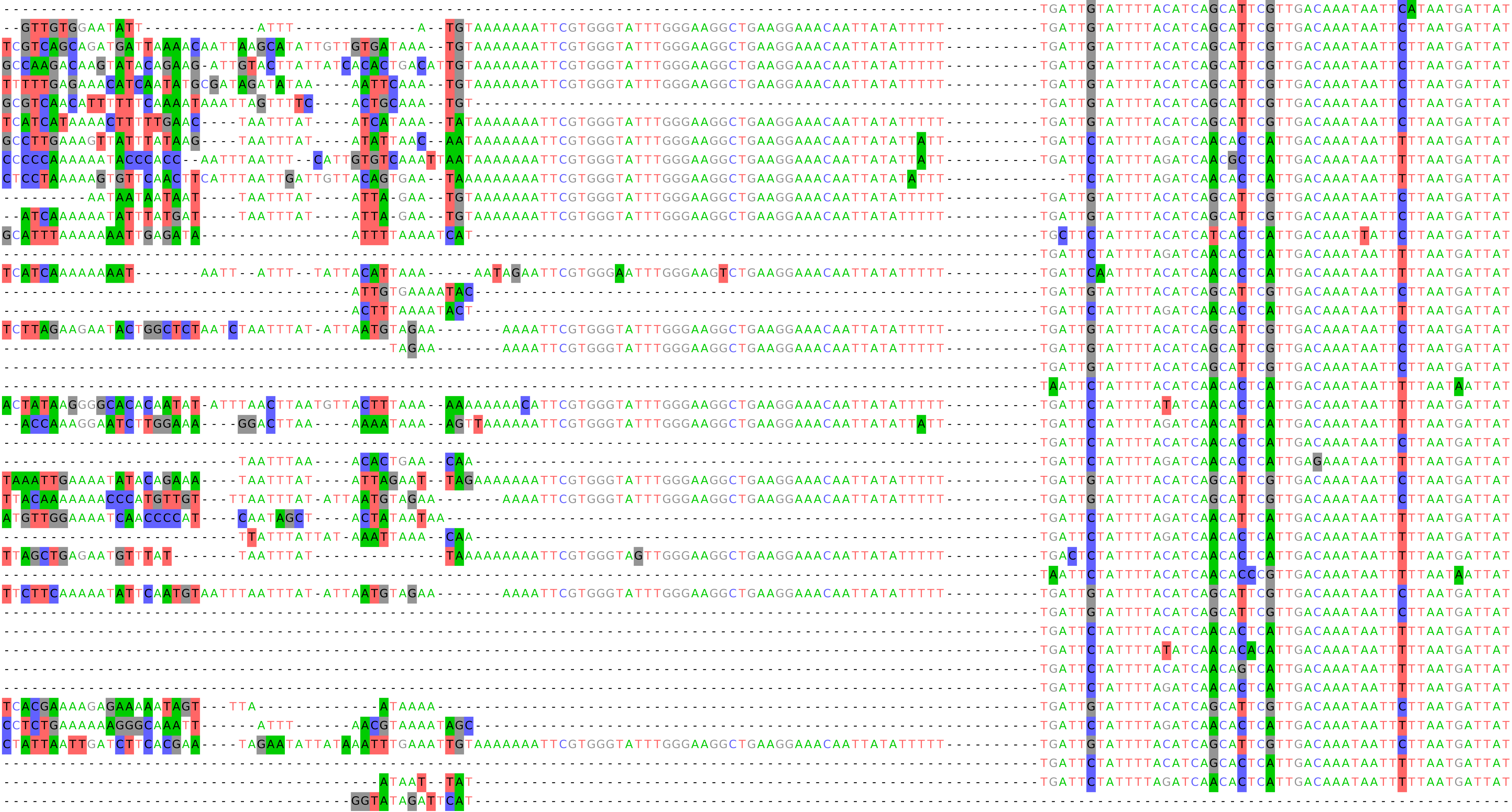


Start crop Point

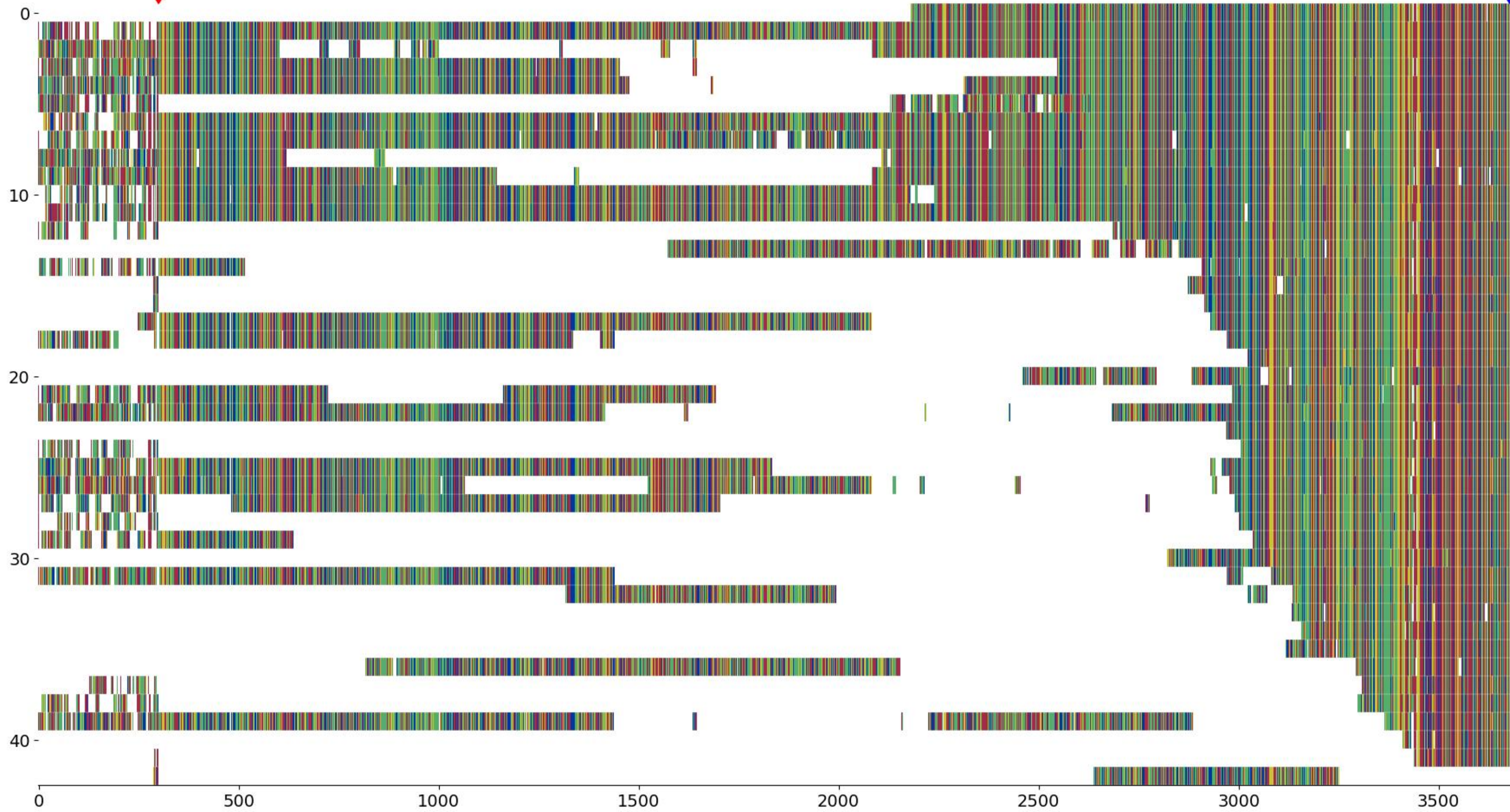
End crop Point

MSA length = 3379



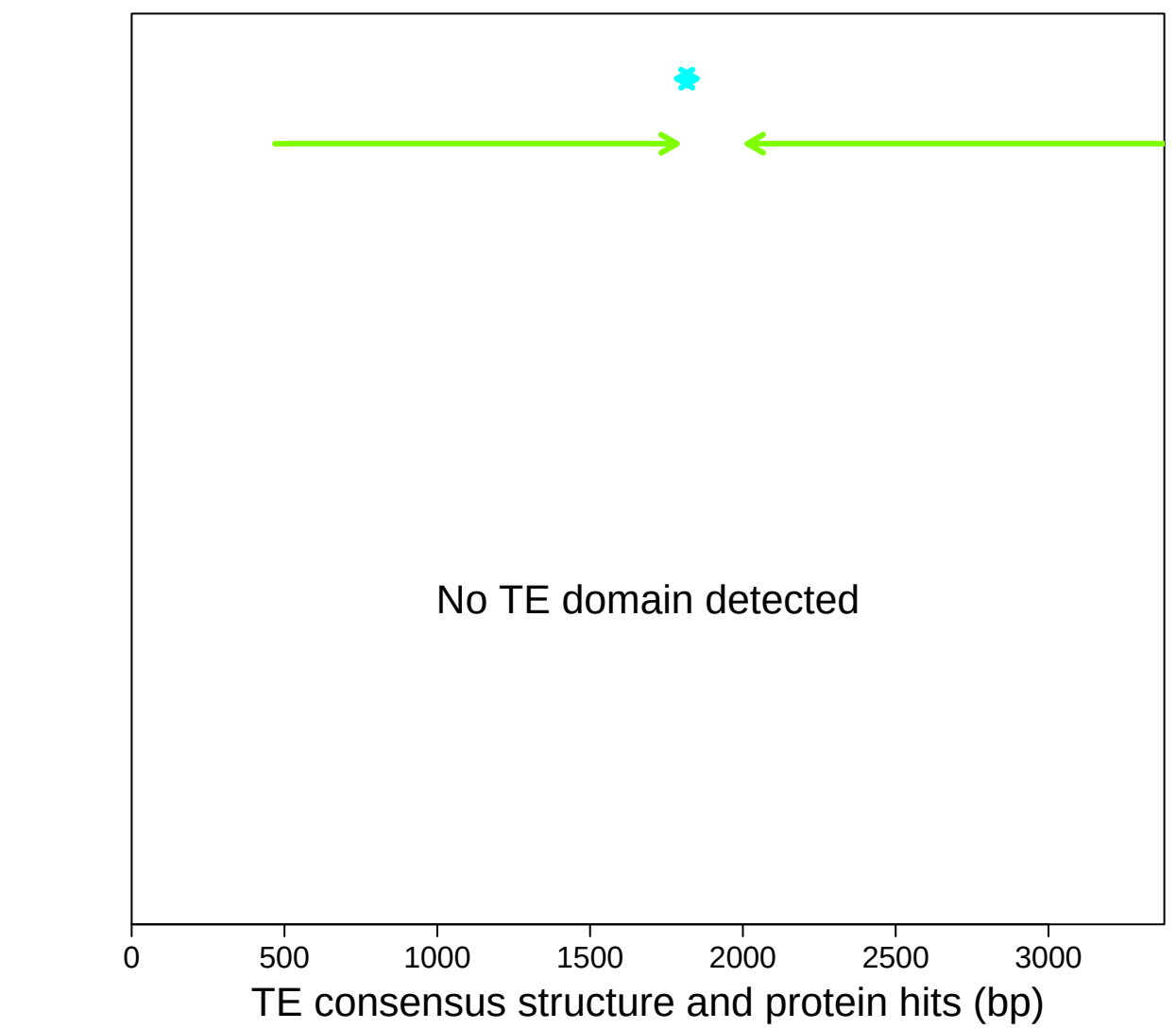
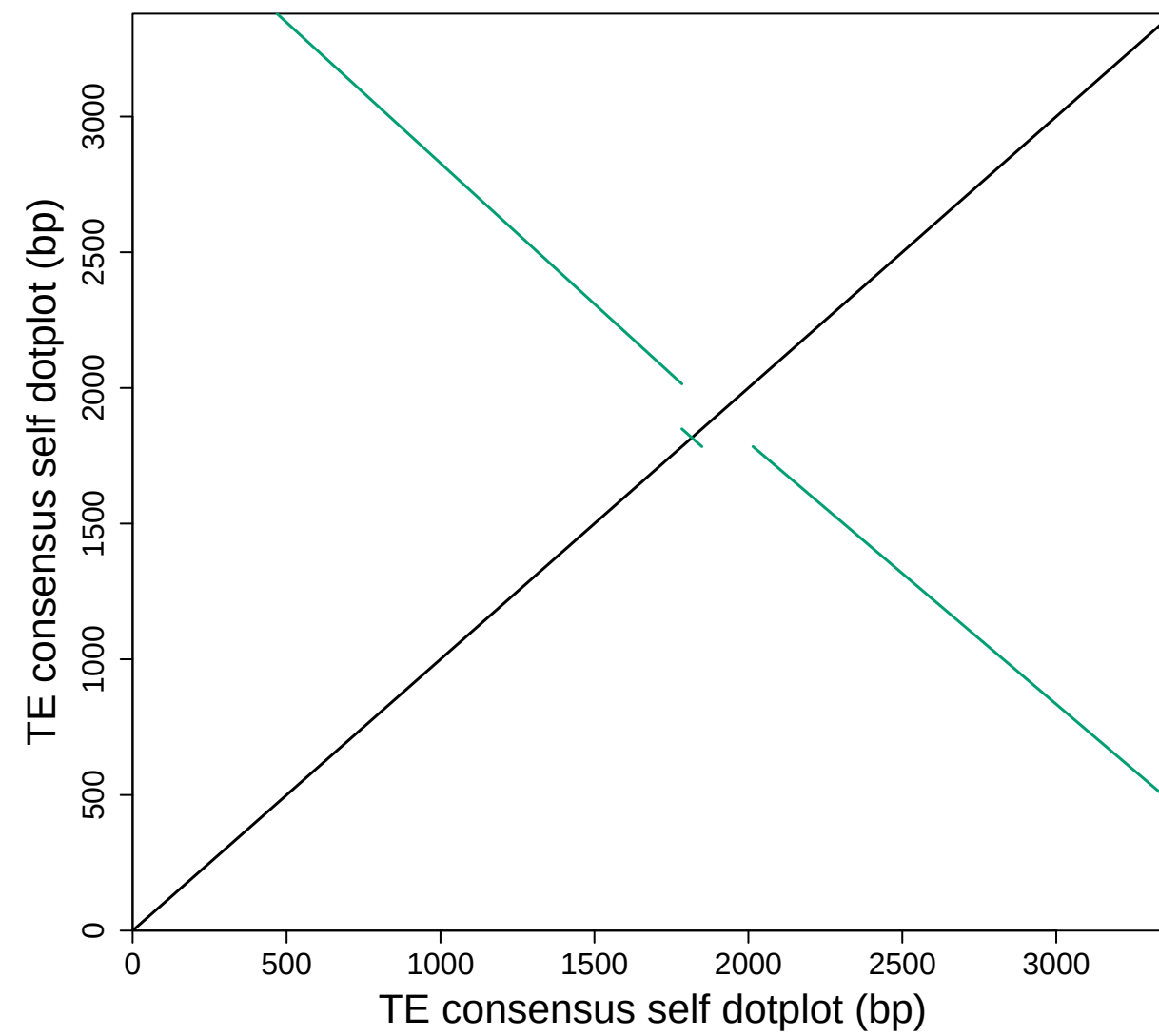
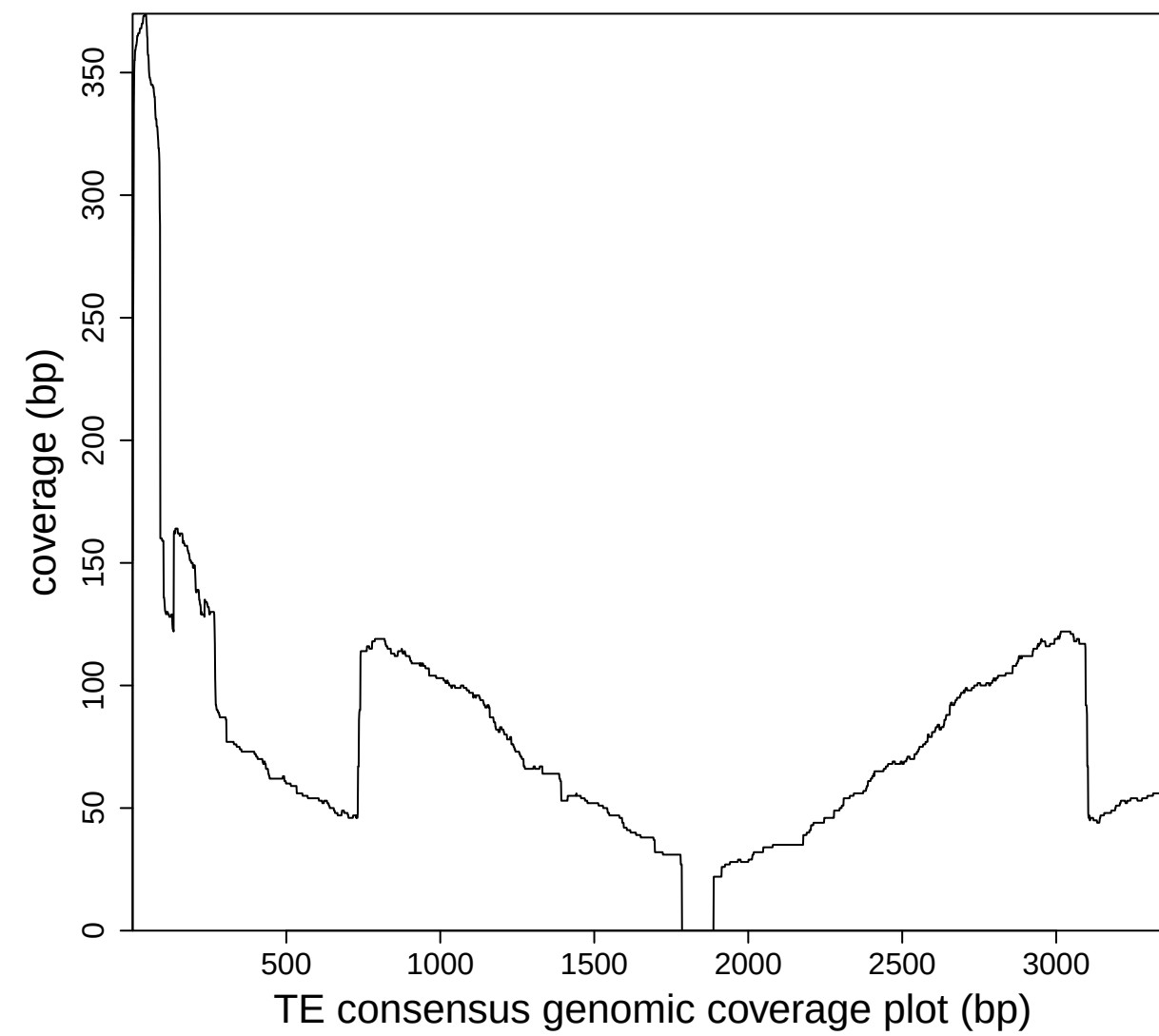
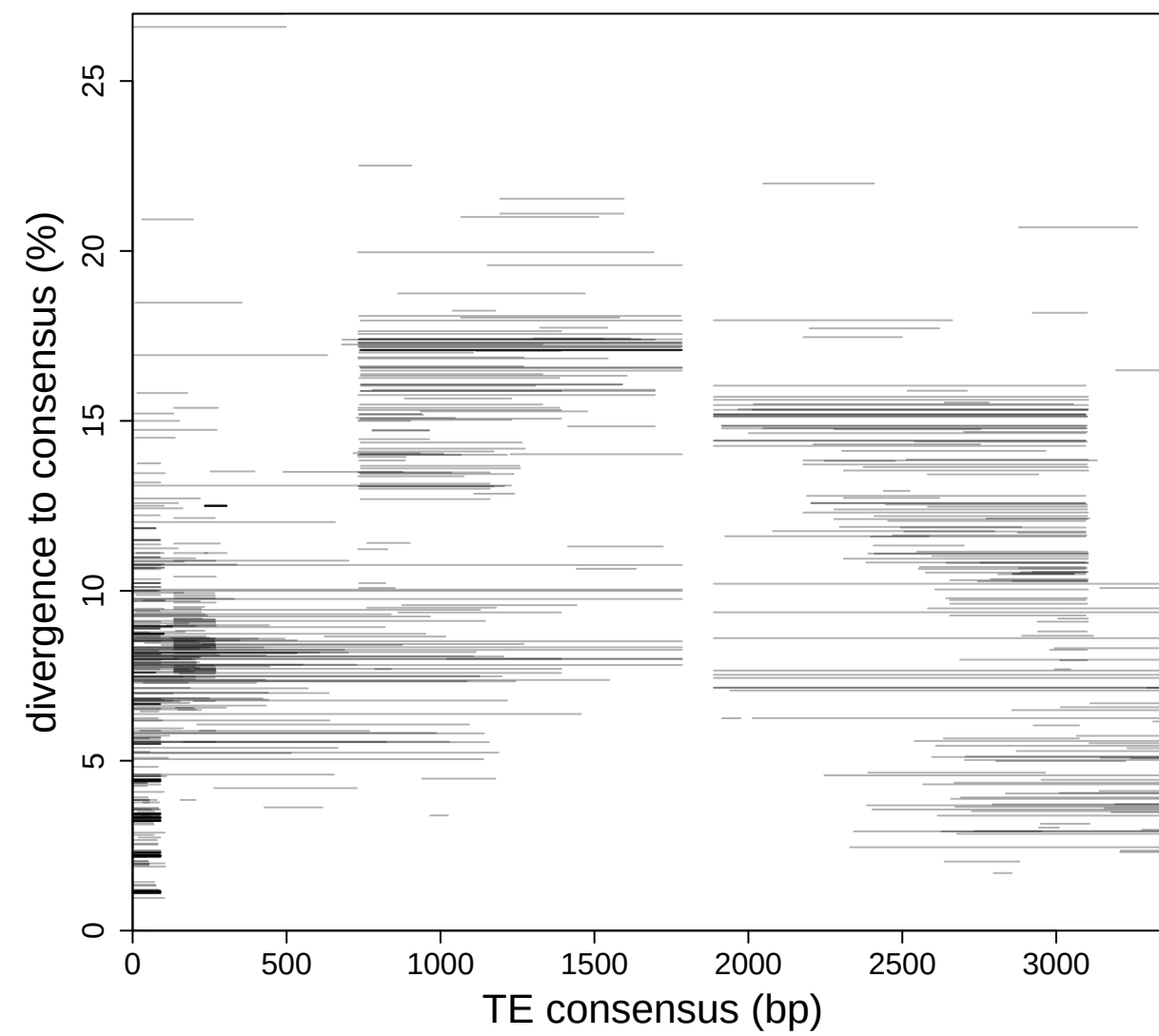
Start crop Point

End crop Point



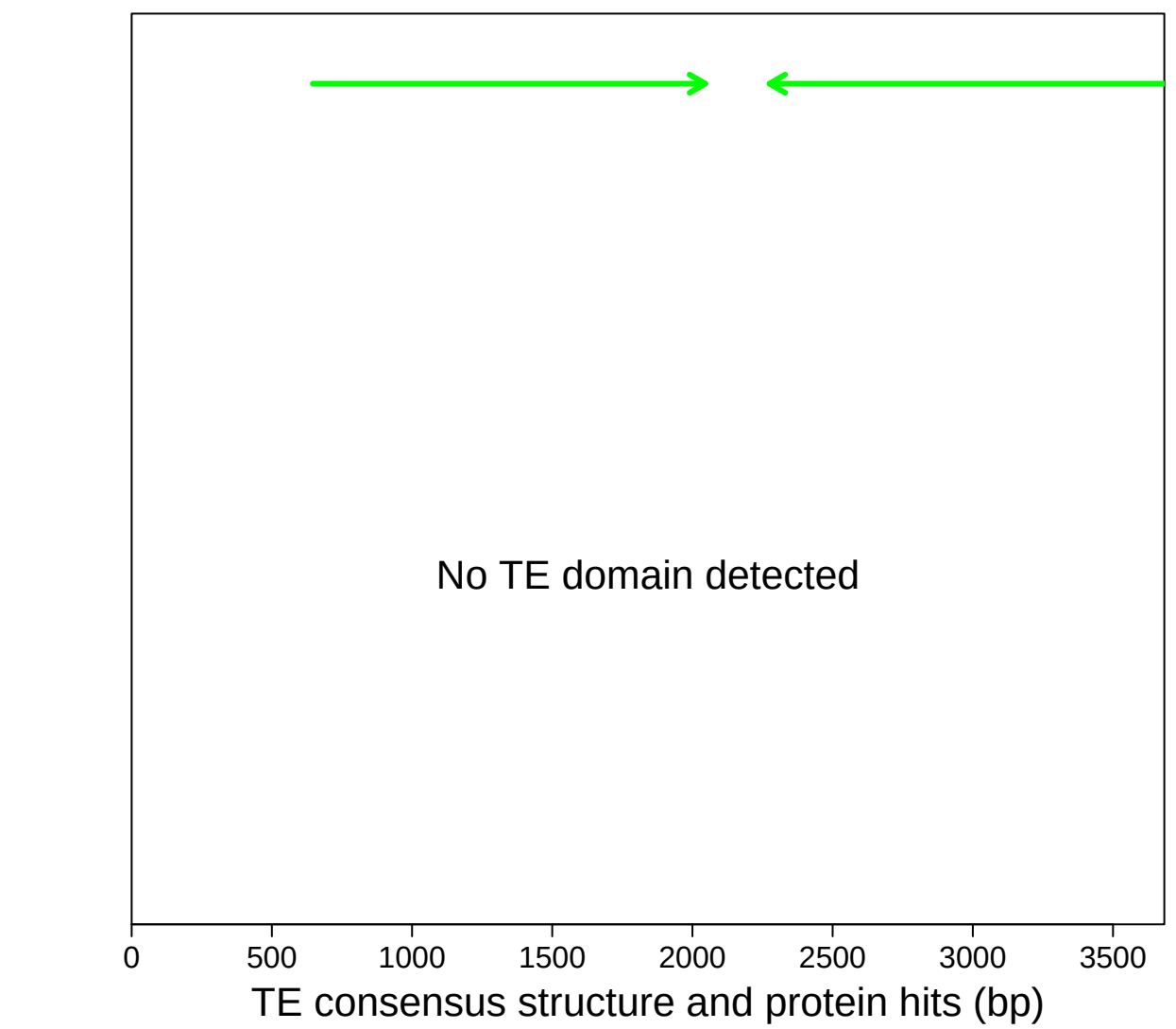
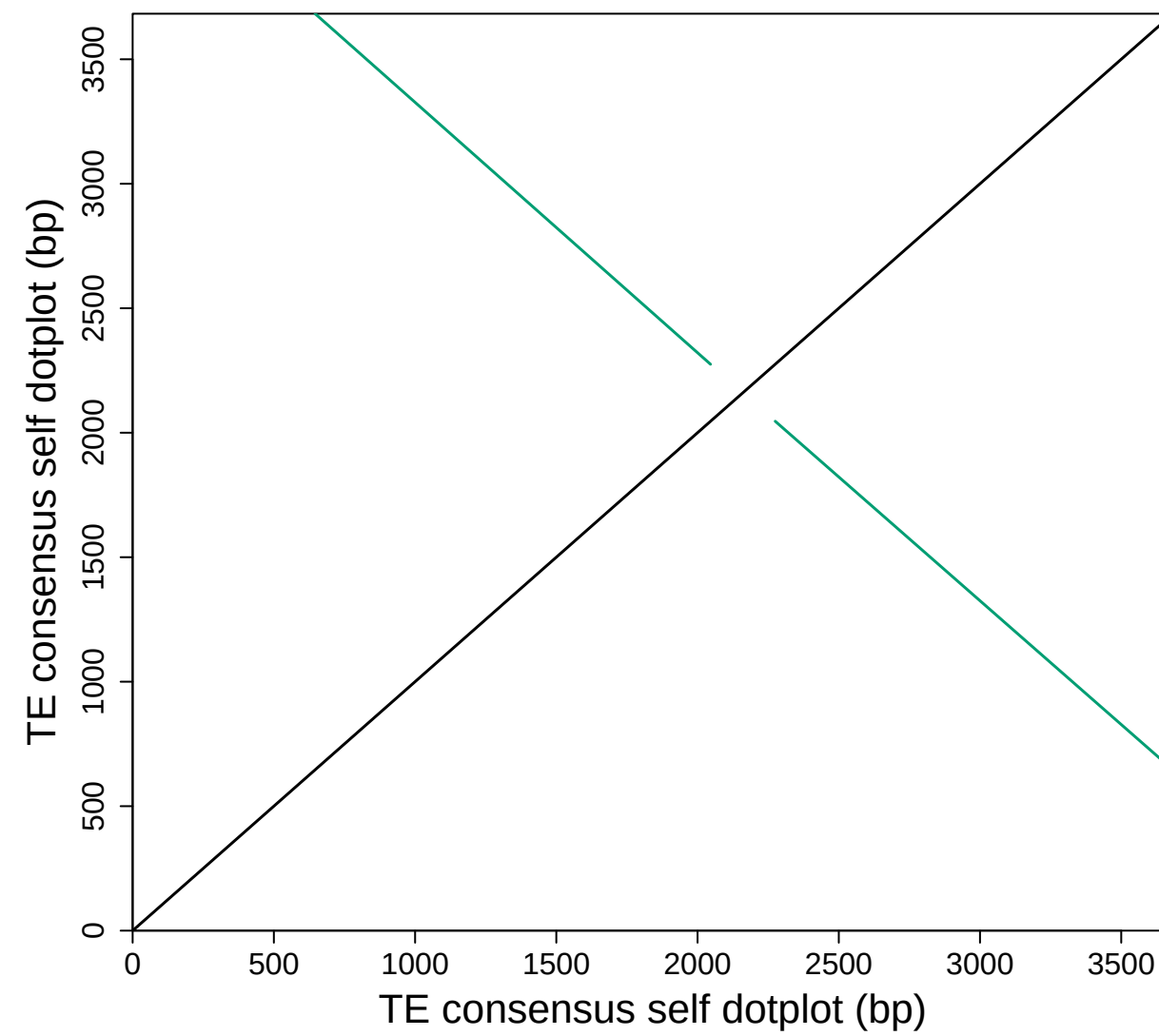
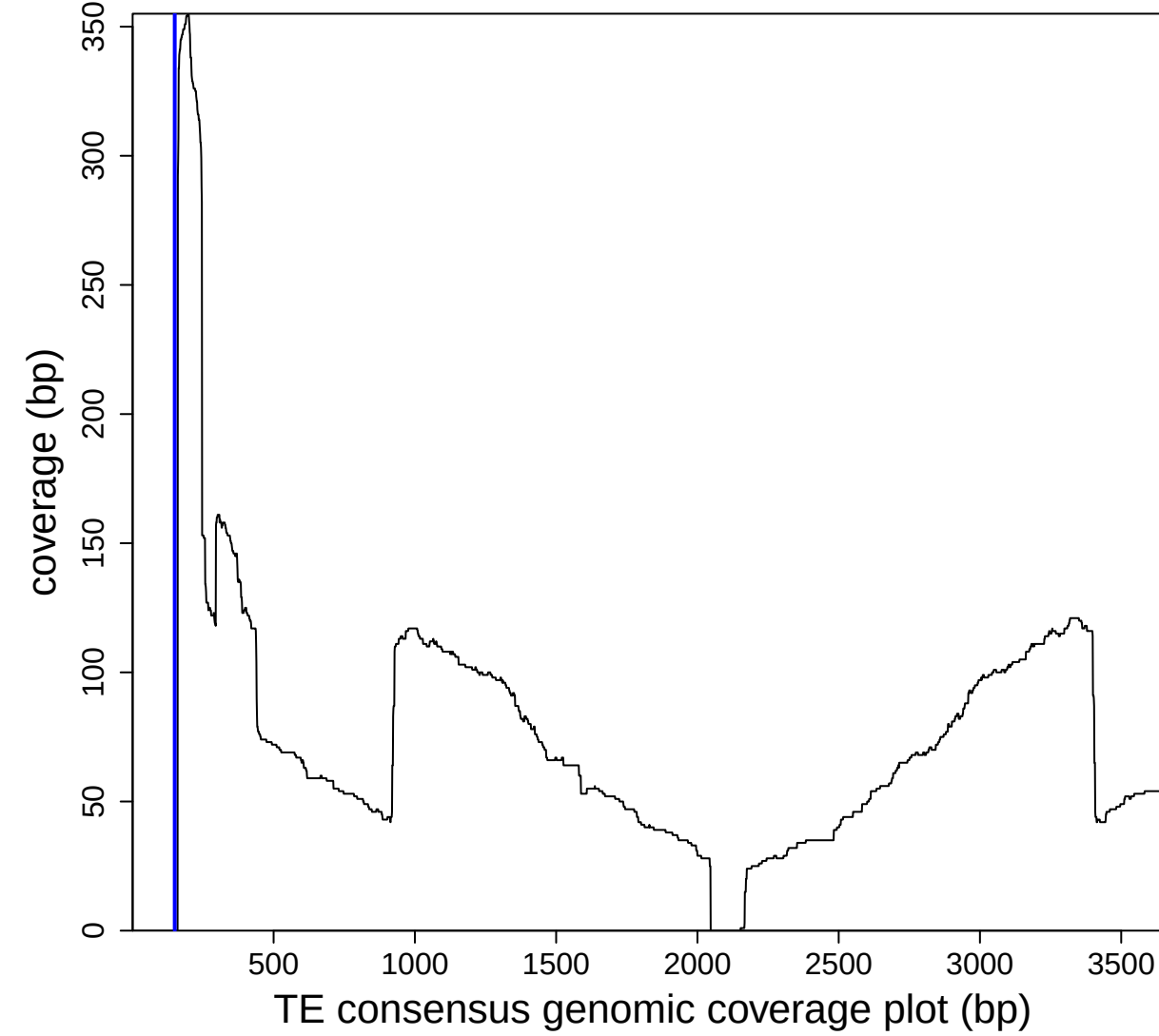
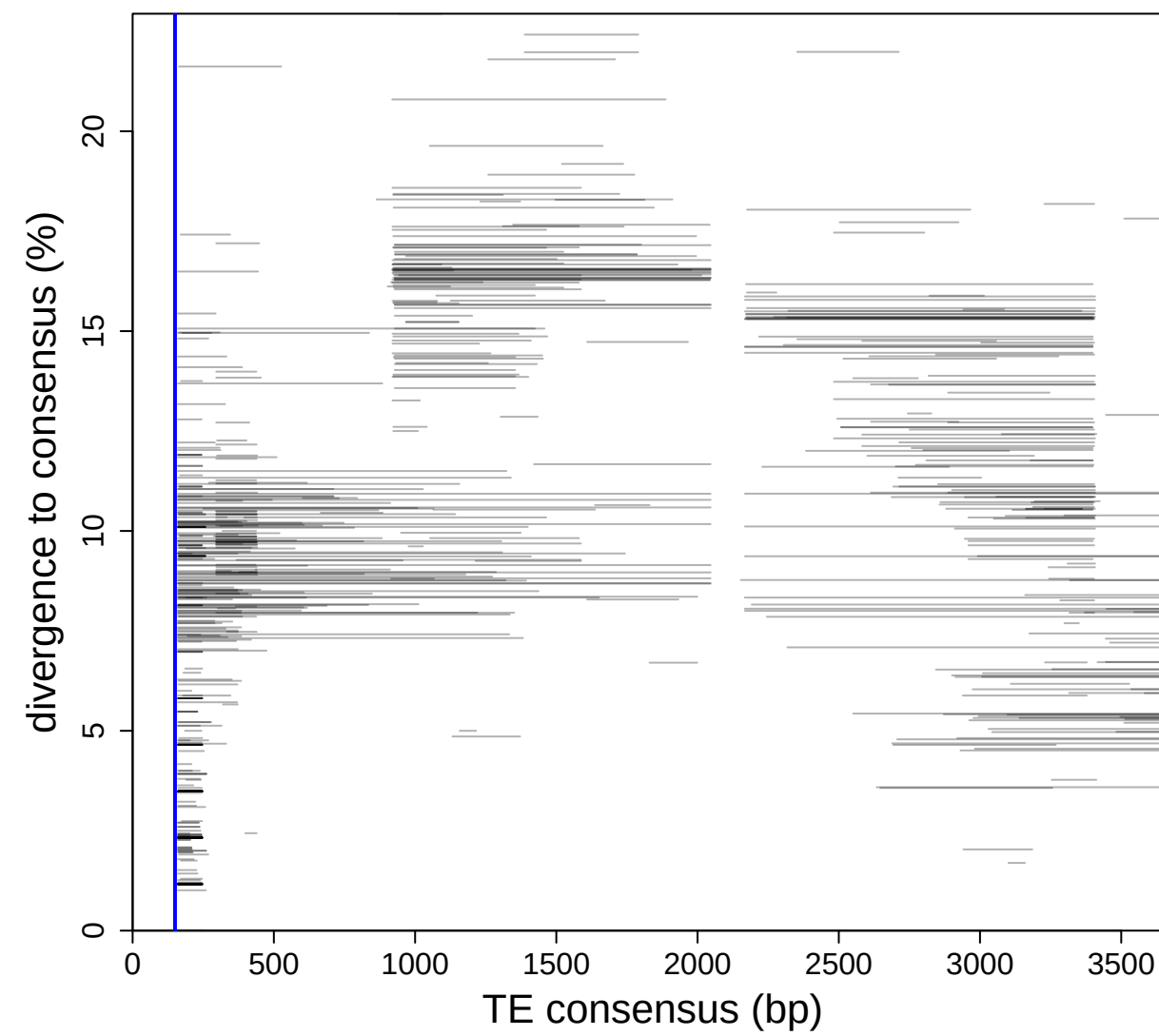
1.bed_fm 1.bed_0_0_n.bed_g 1.bed_fm 2.bed_0_0_bcln.fa aln.
size: 3379bp; fragments: 705; full length: 0 (>=3041.1bp)

After TEtrimmer 3379 bp



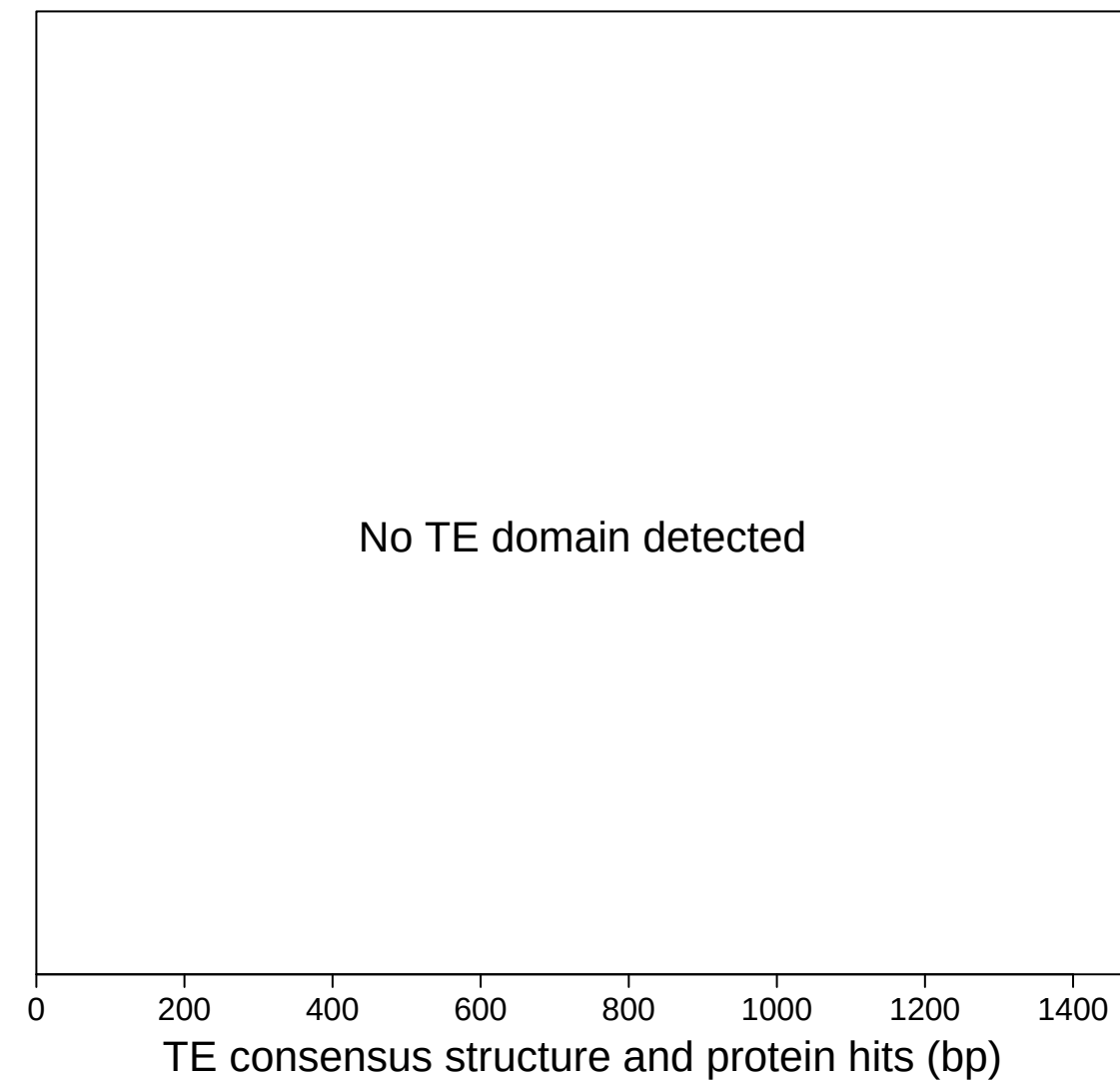
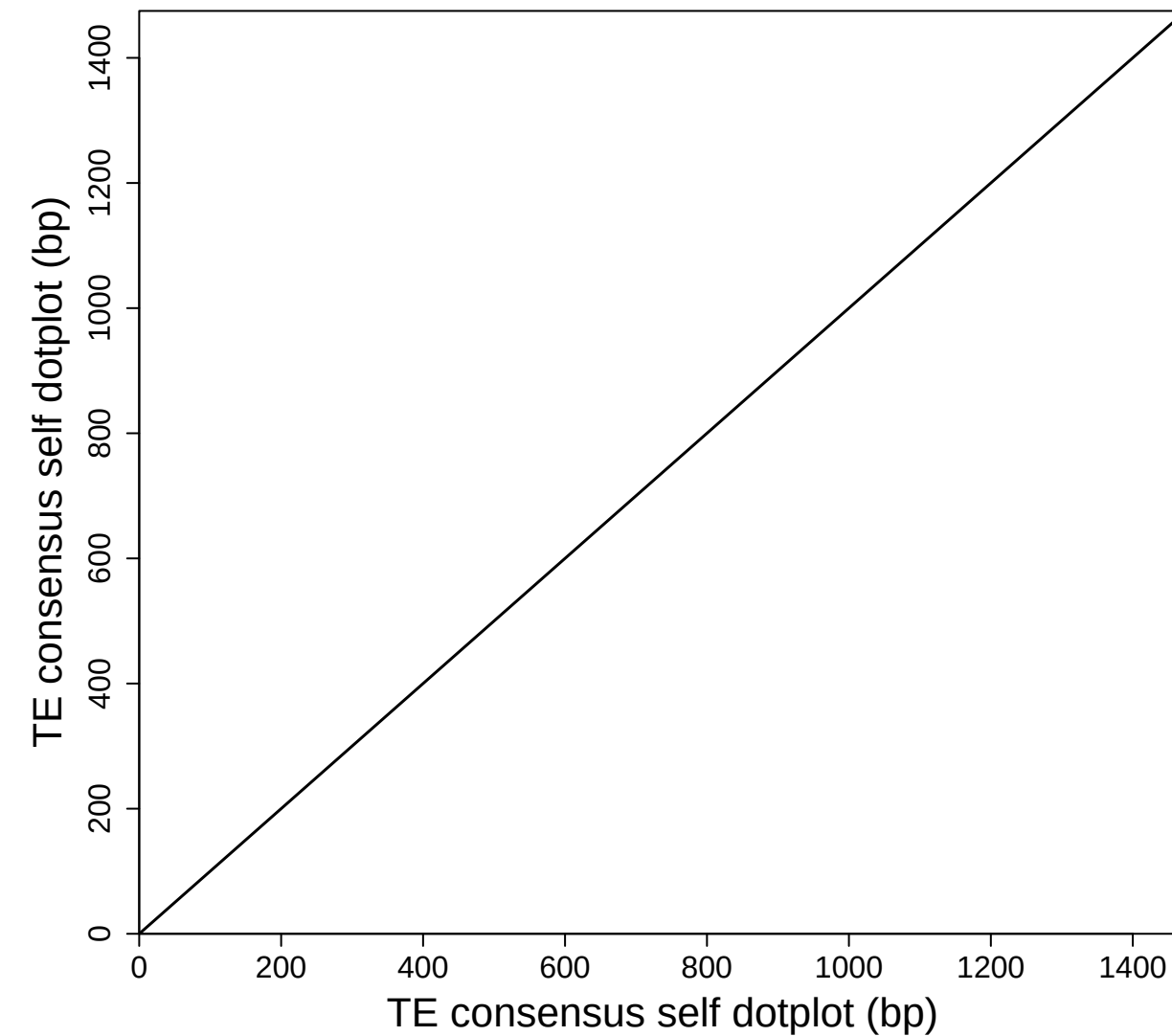
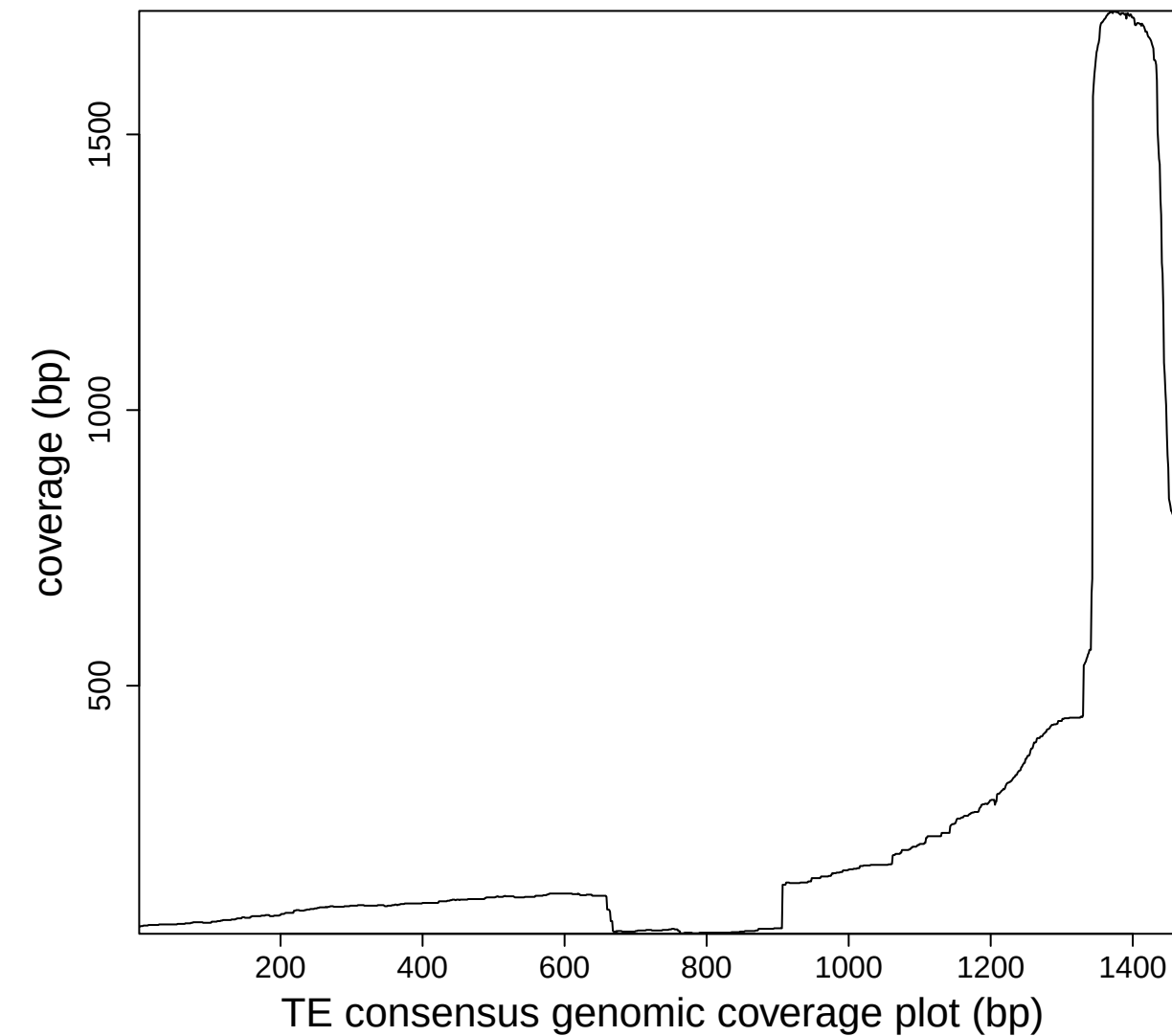
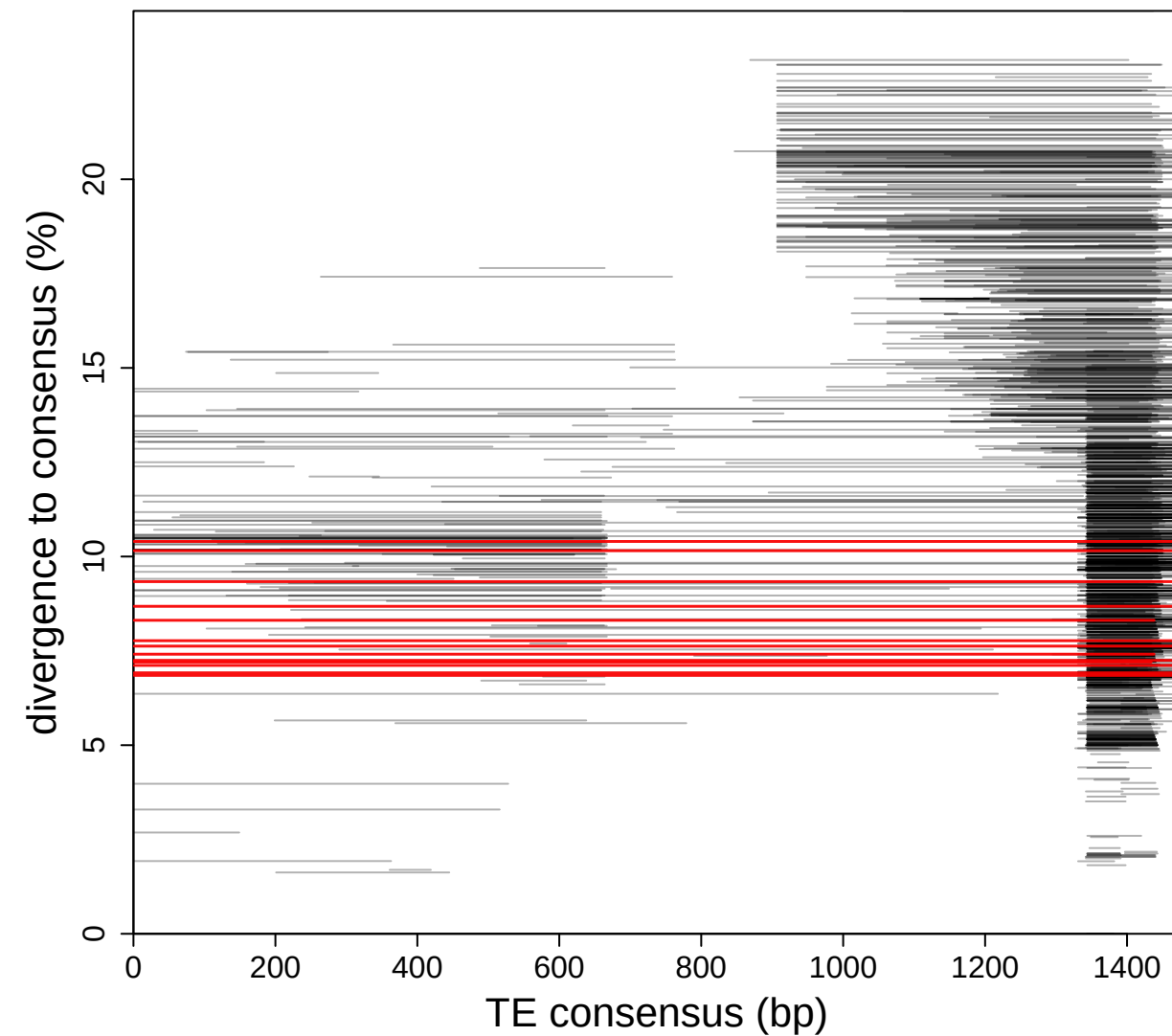
uf.bed g 1.bed fm 1.bed 0 0 n.bed g 1.bed fm 2.bed 0 0 bc
size: 3683bp; fragments: 672; full length: 0 (>=3314.7bp)

After TEtrimmer Extended plot Blue lines are boundaries

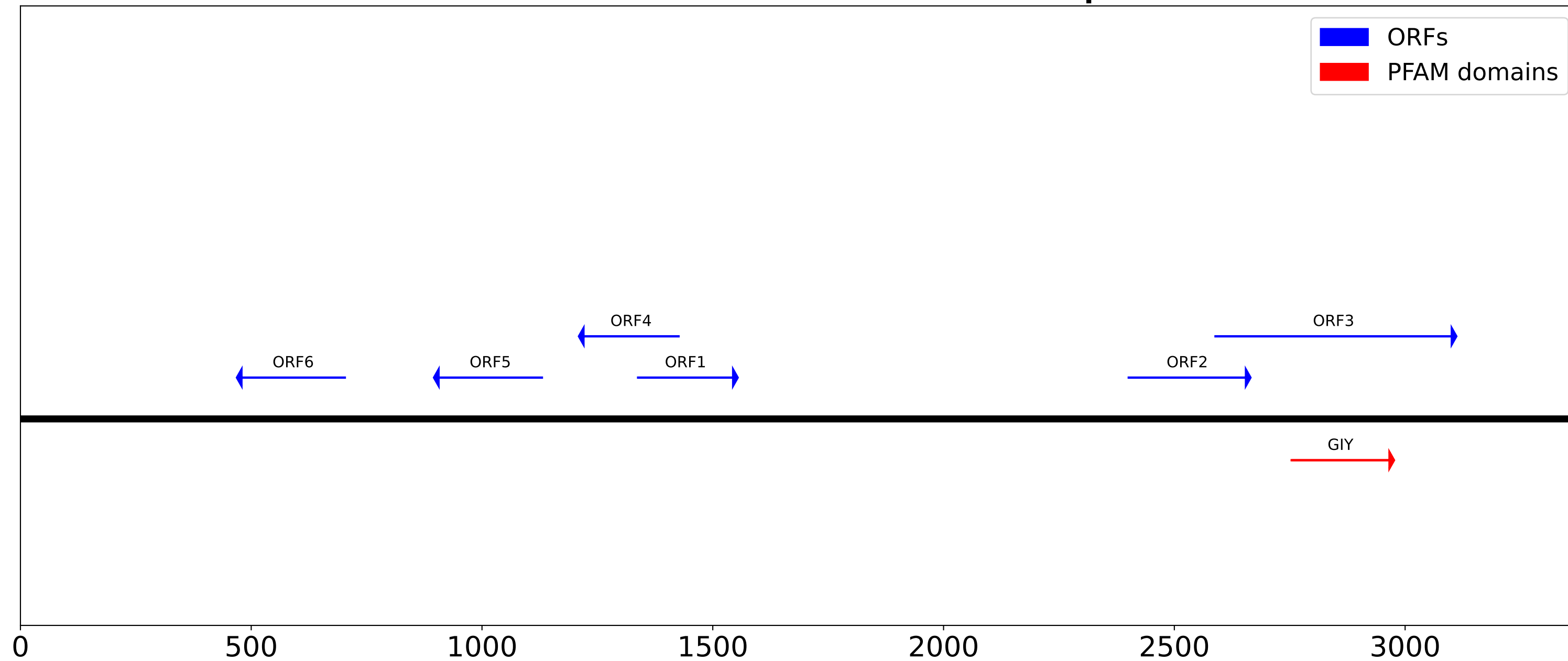


TE: rnd_5_family_10213
size: 1475bp; fragments: 1888; full length: 13 (≥ 1327.5 bp)

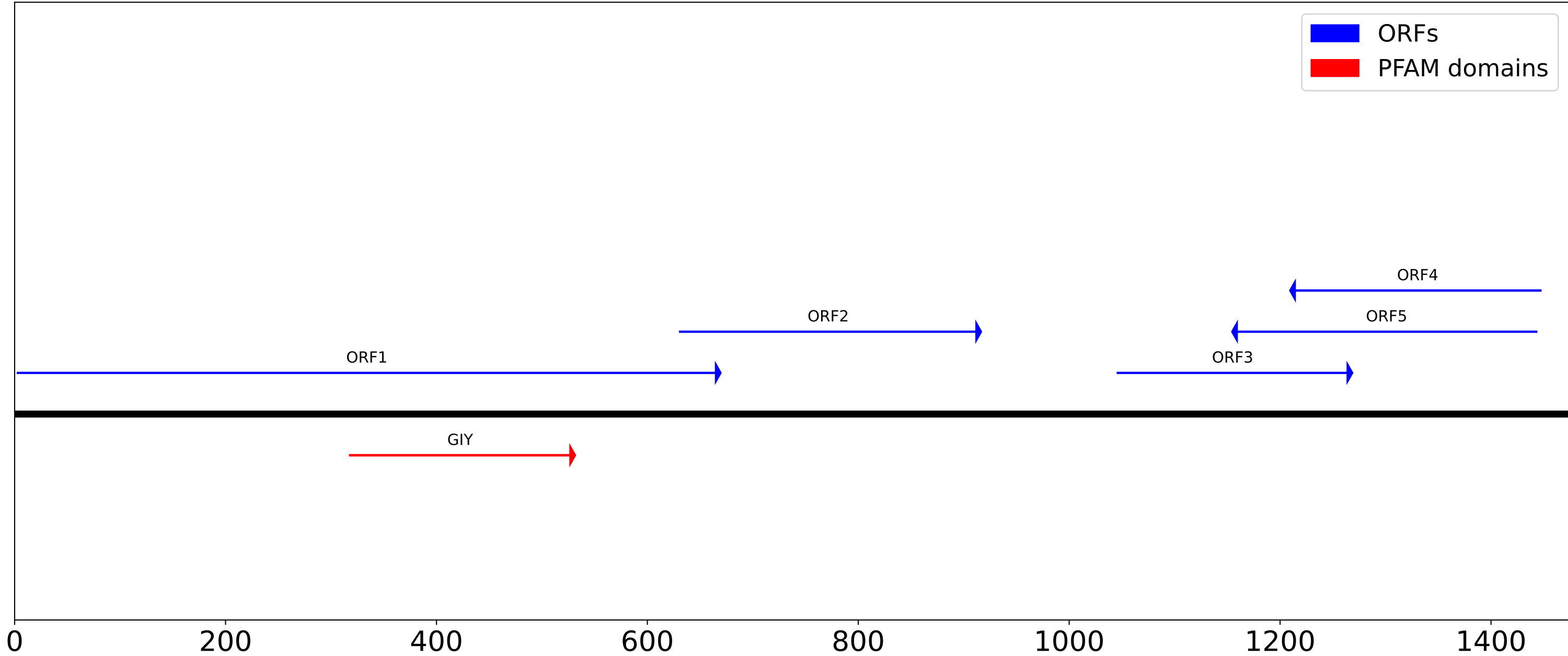
Before TEtrimmer 1475 bp



After TEtrimmer ORF and PFAM domain plot



Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)

