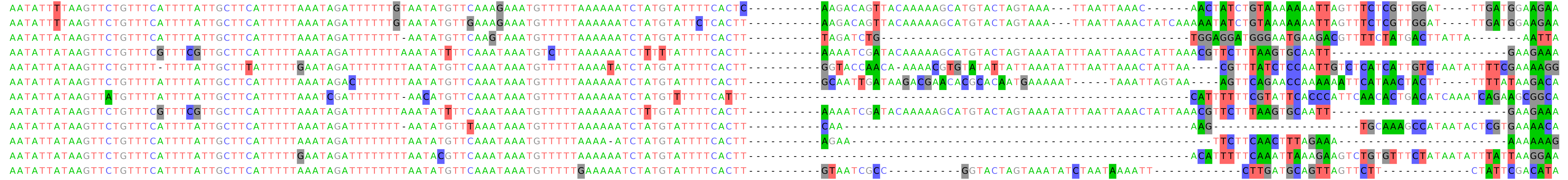


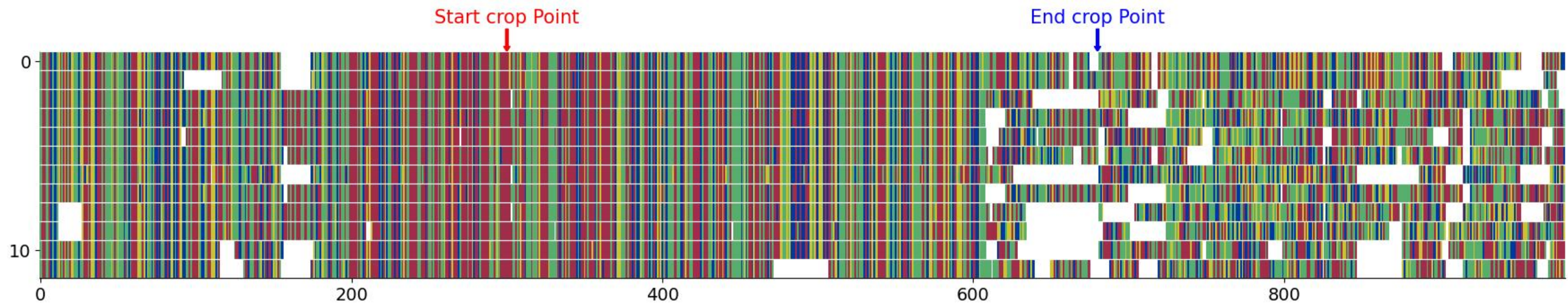
Start crop Point



MSA length = 381

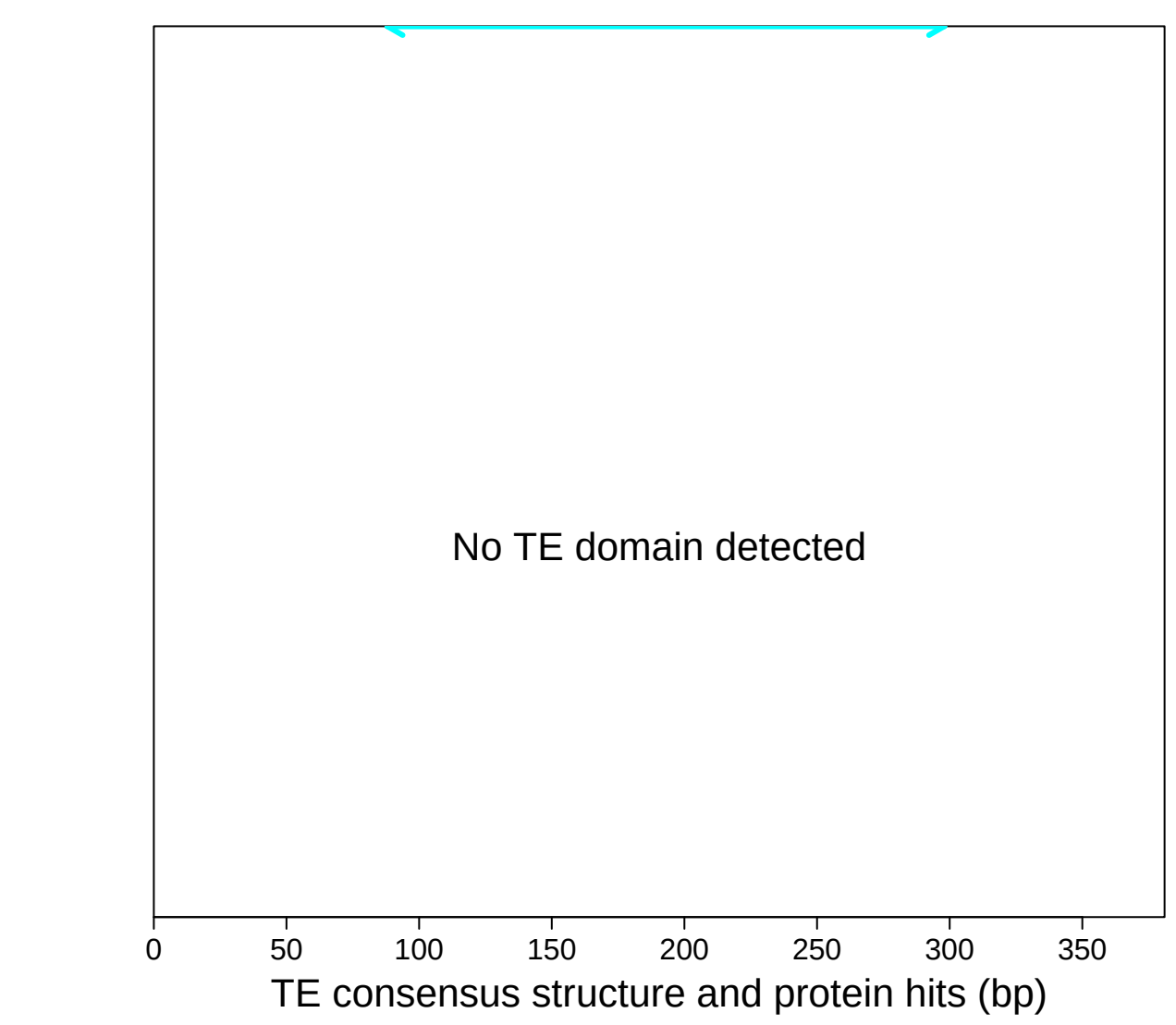
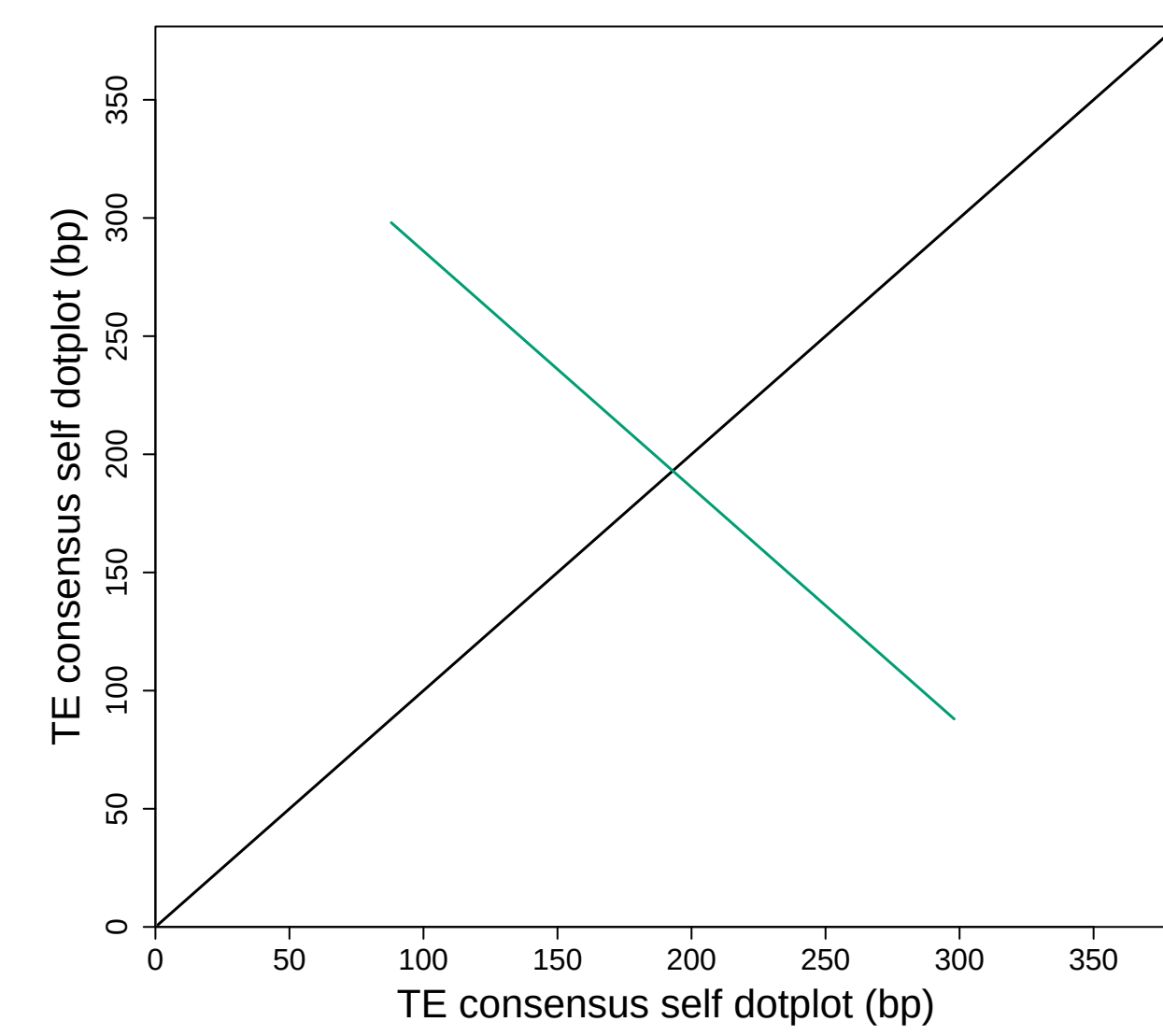
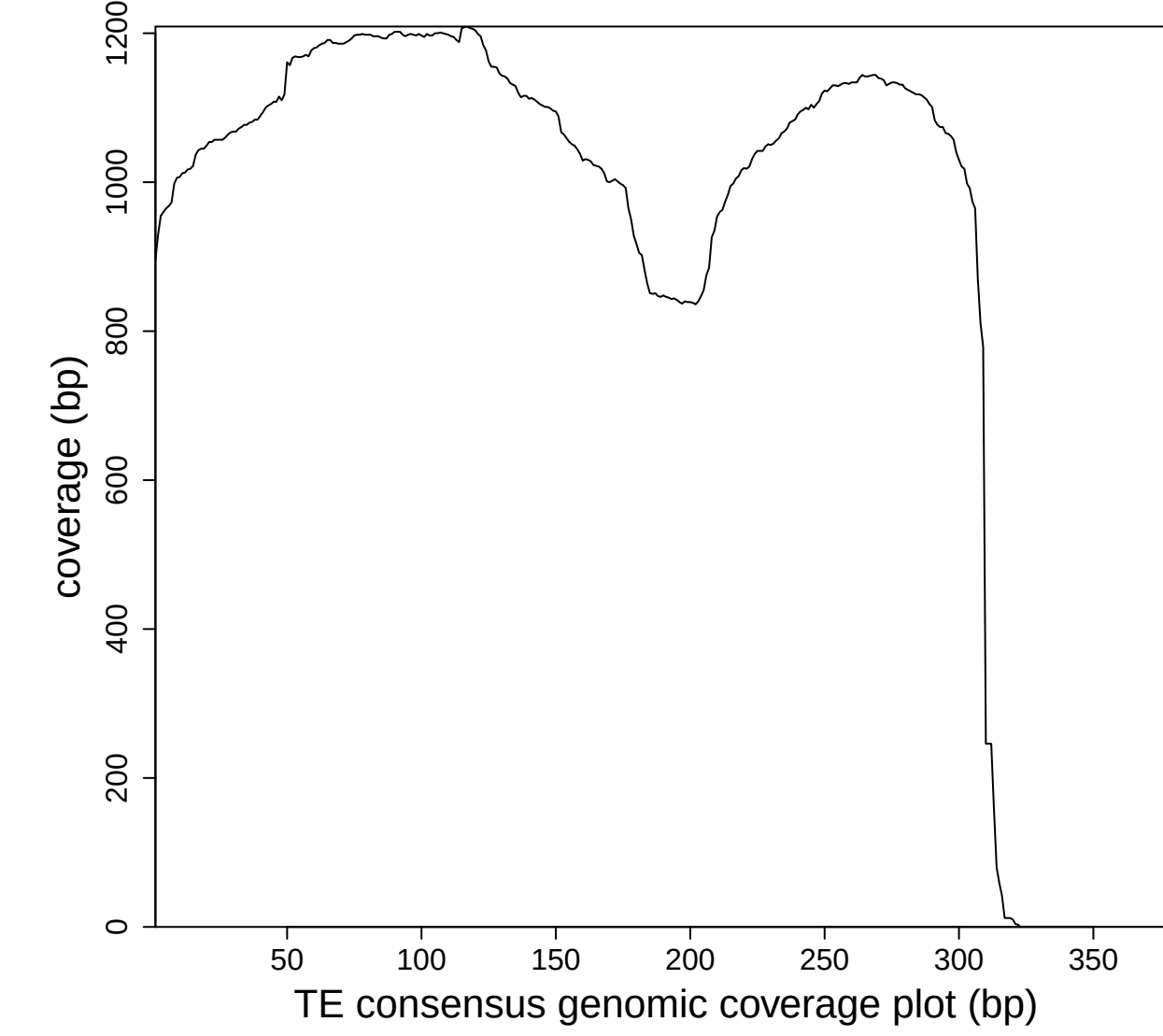
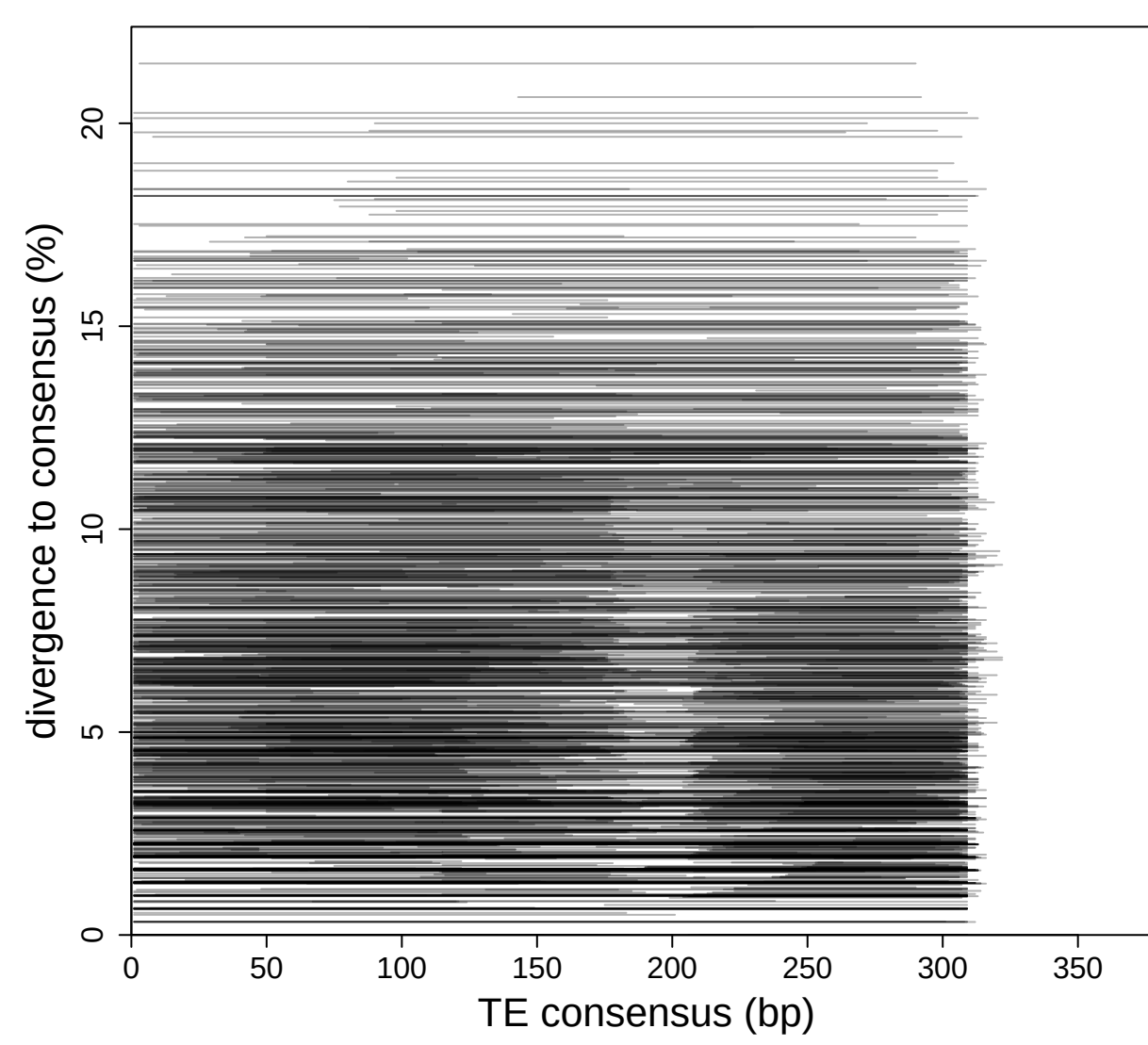
End crop Point





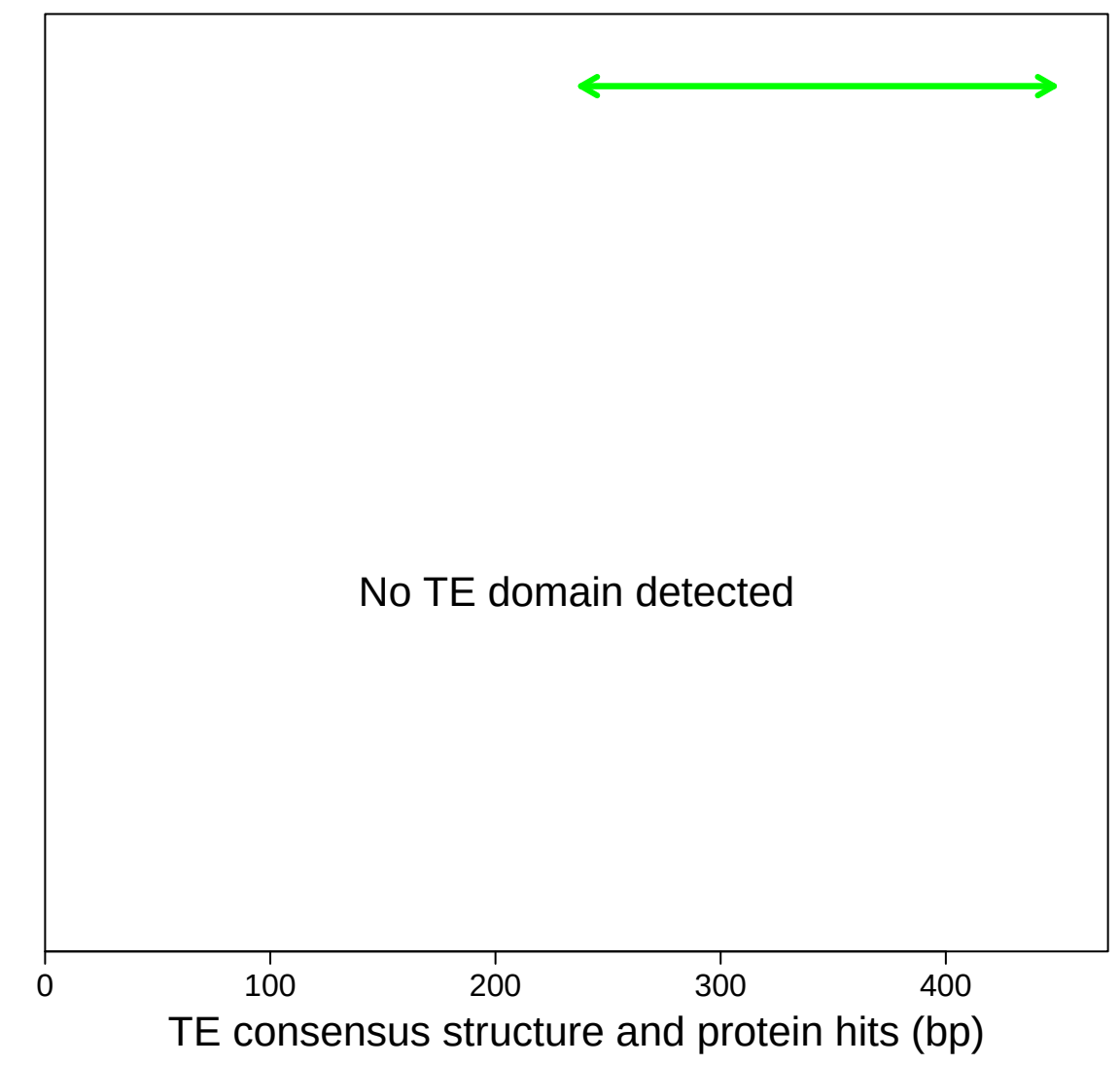
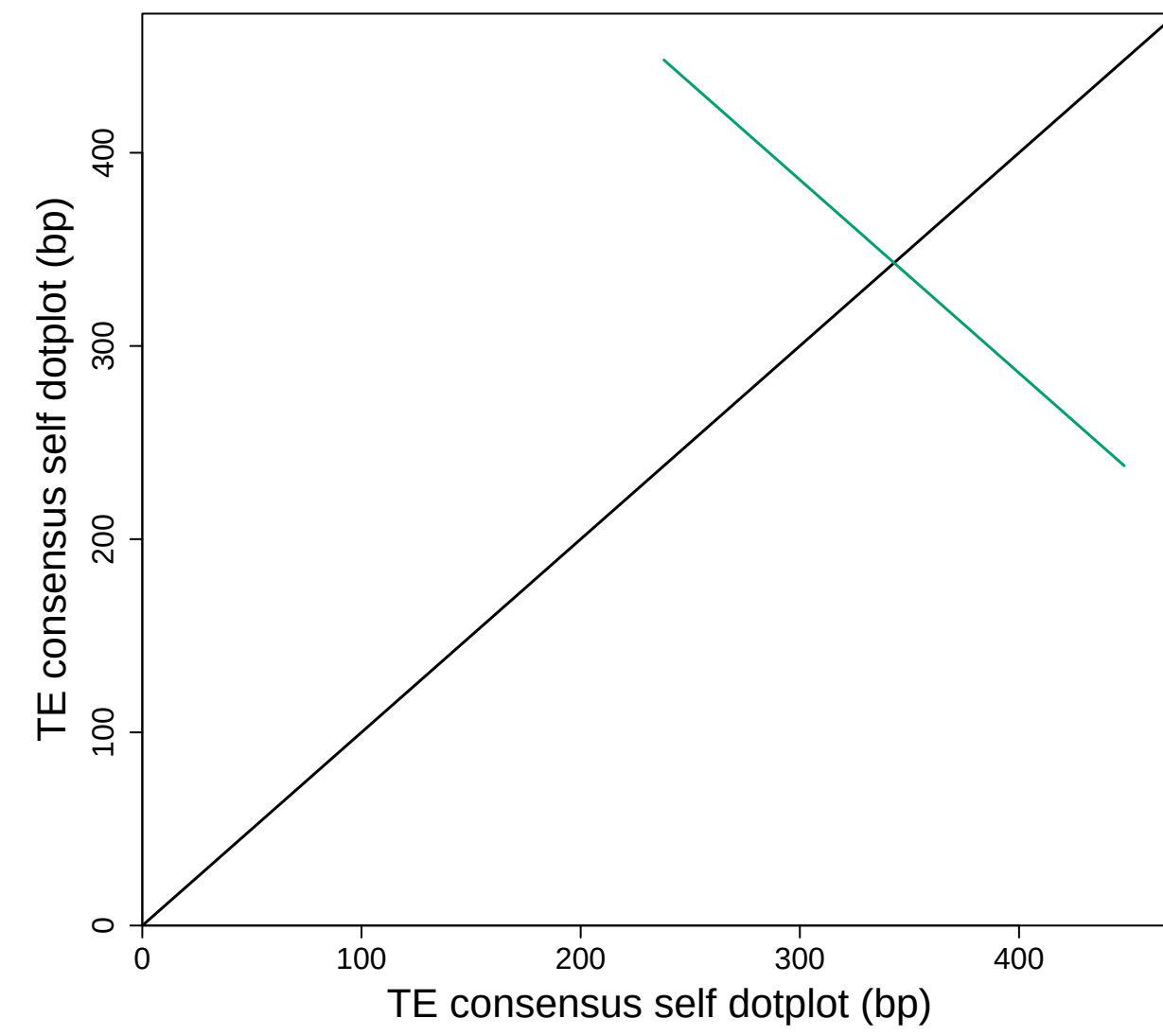
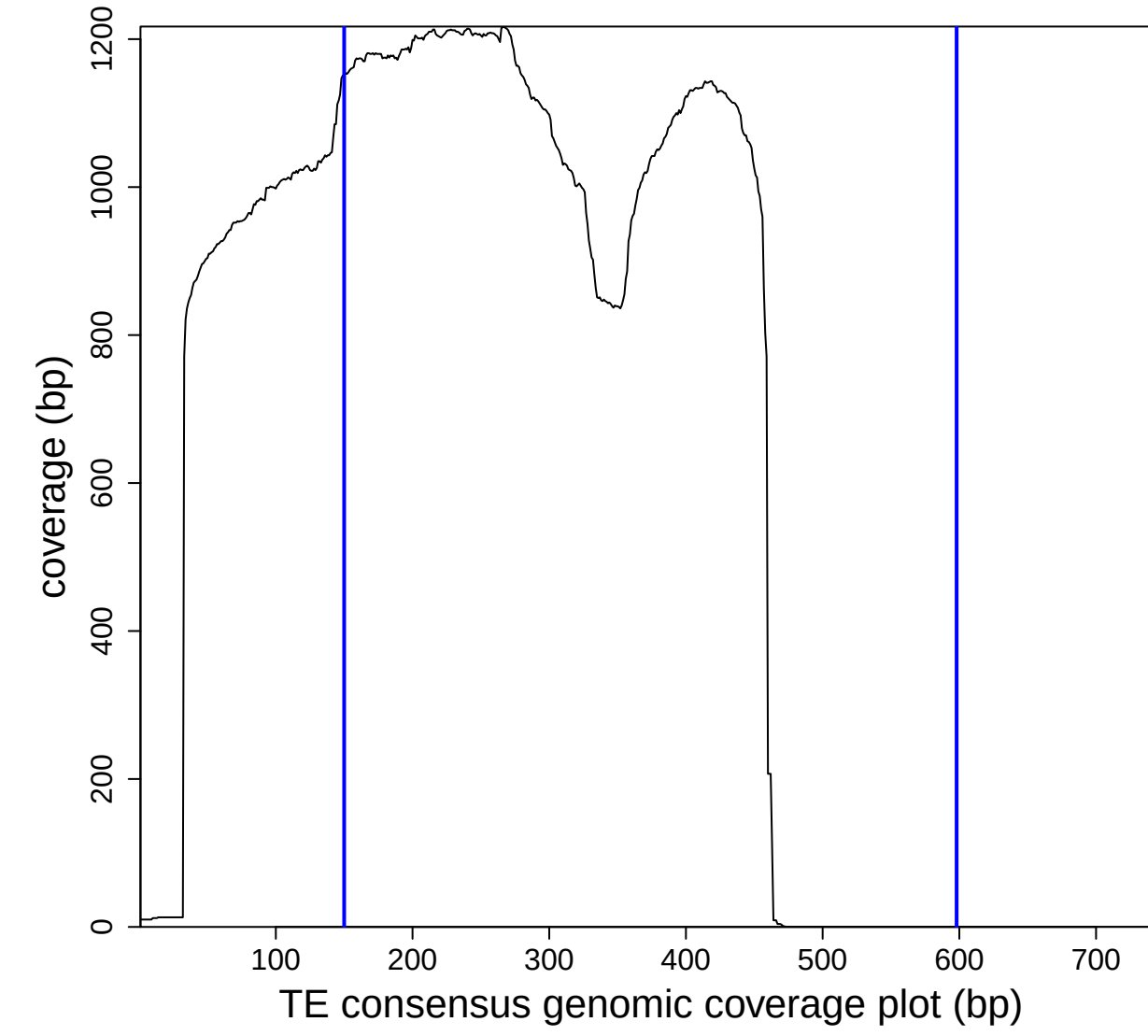
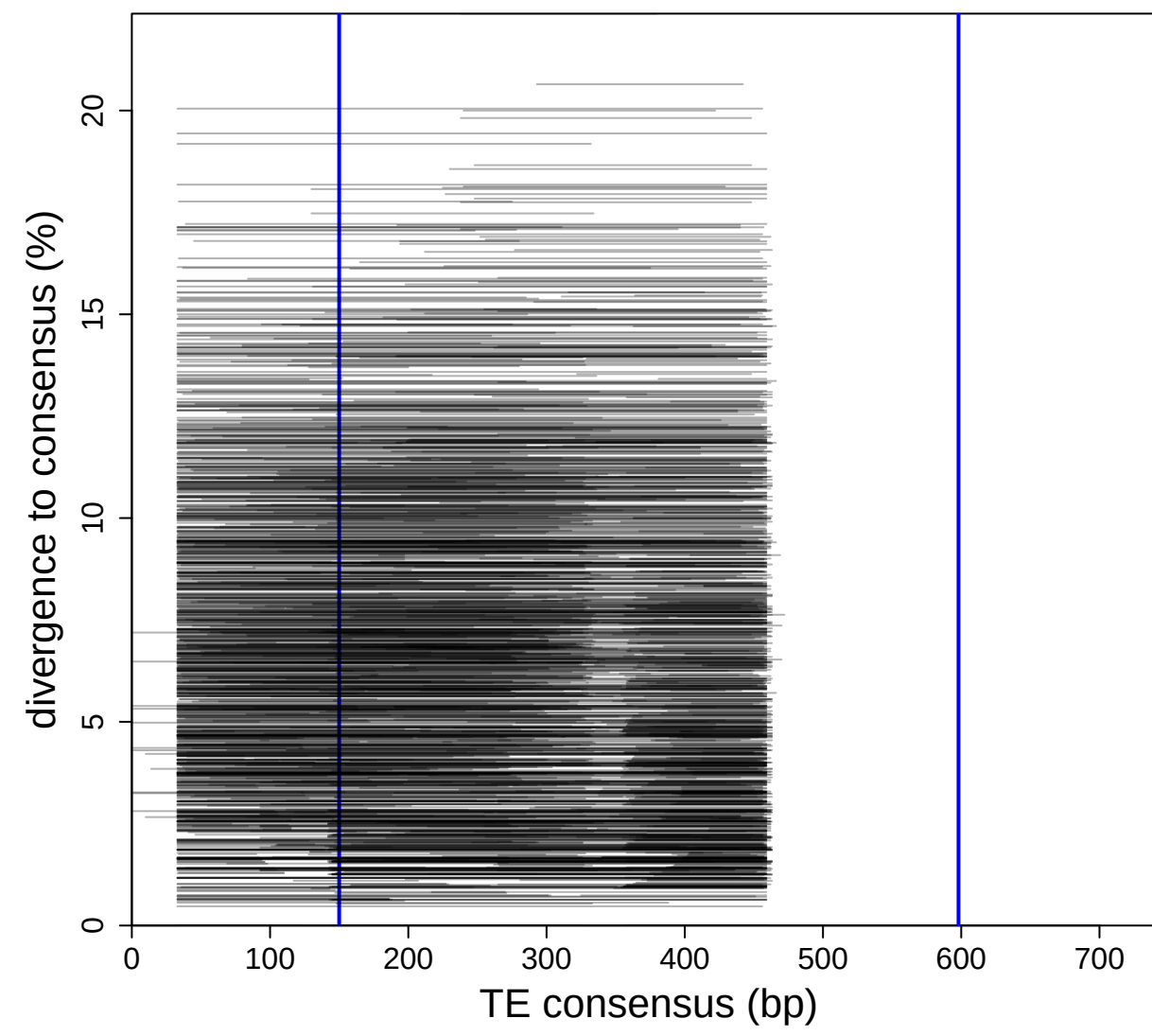
ed_uf.bed g_3.bed fm_1.bed 0 0 bcln.fa aln.fa cl.fa gs.fa ce.
size: 381bp; fragments: 1782; full length: 0 (>=342.9bp)

After TEtrimmer 381 bp

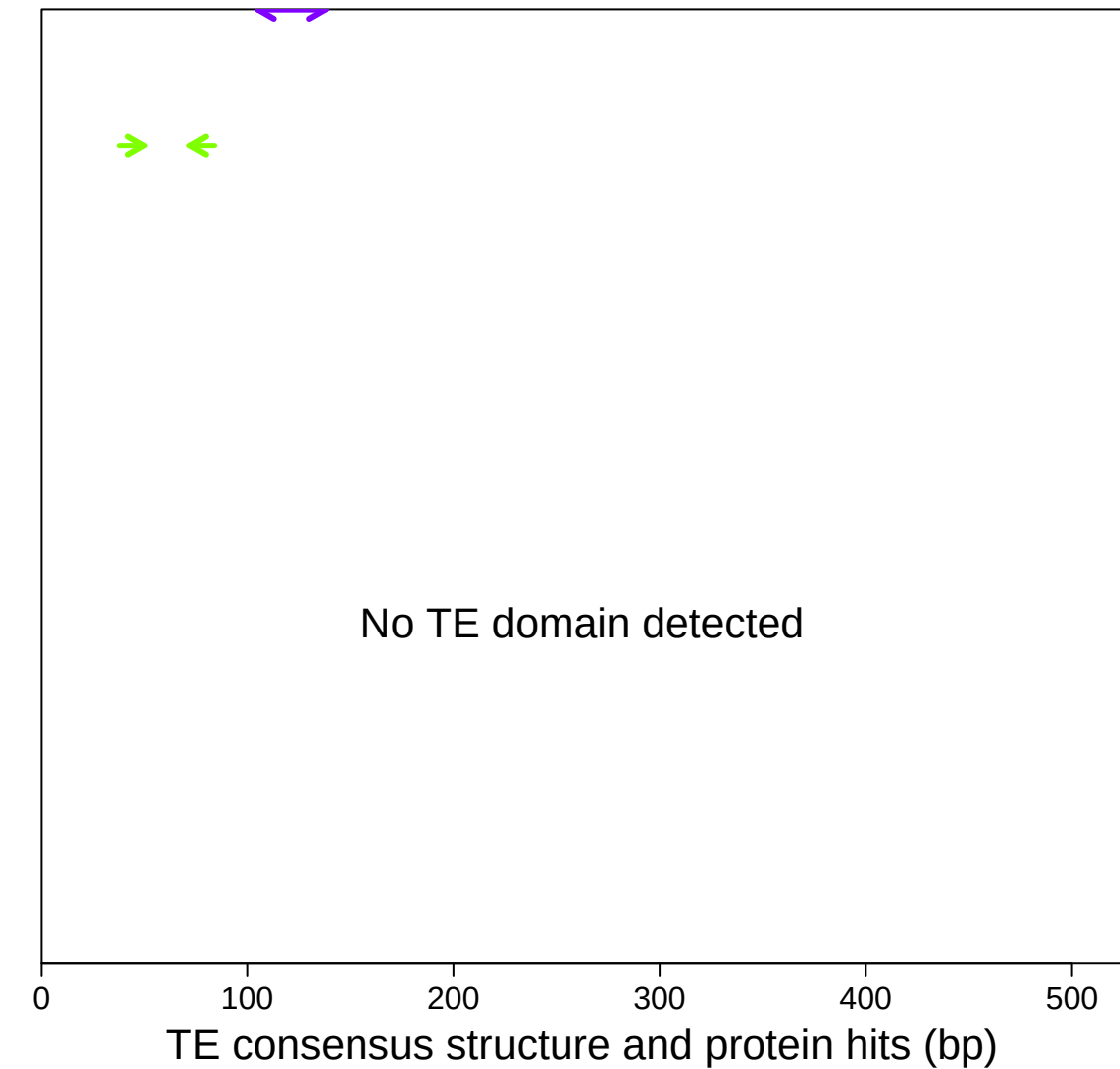
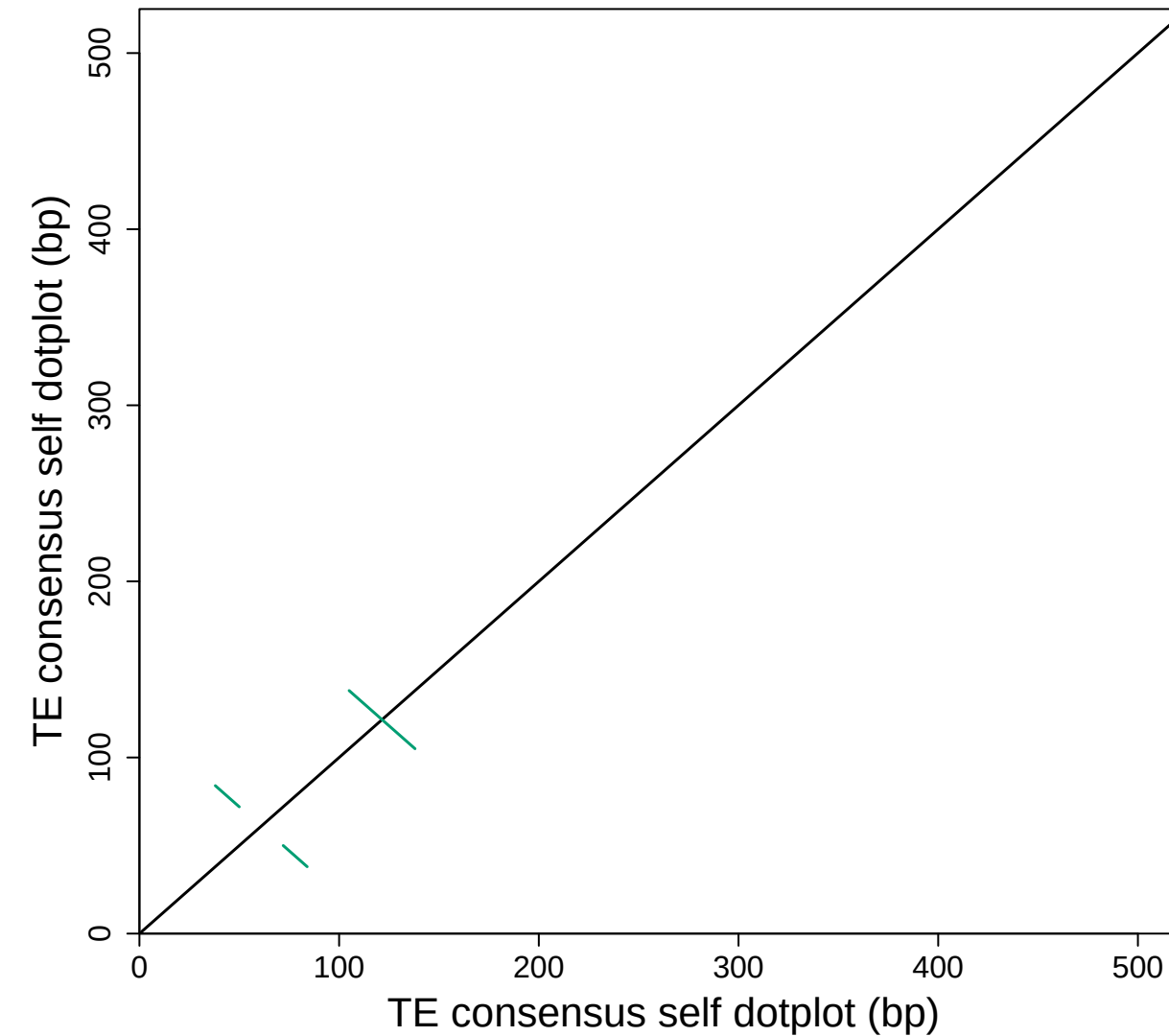
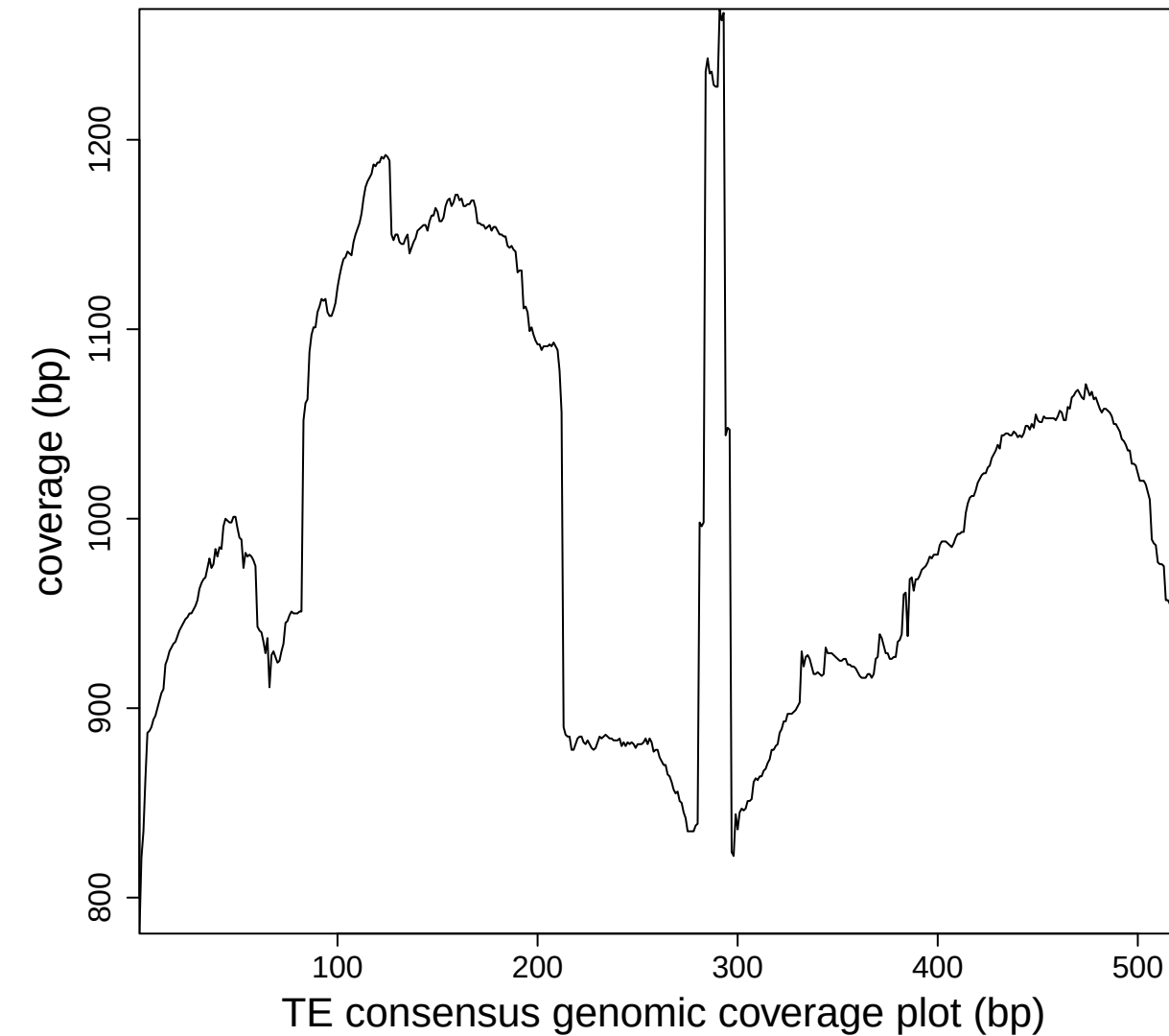
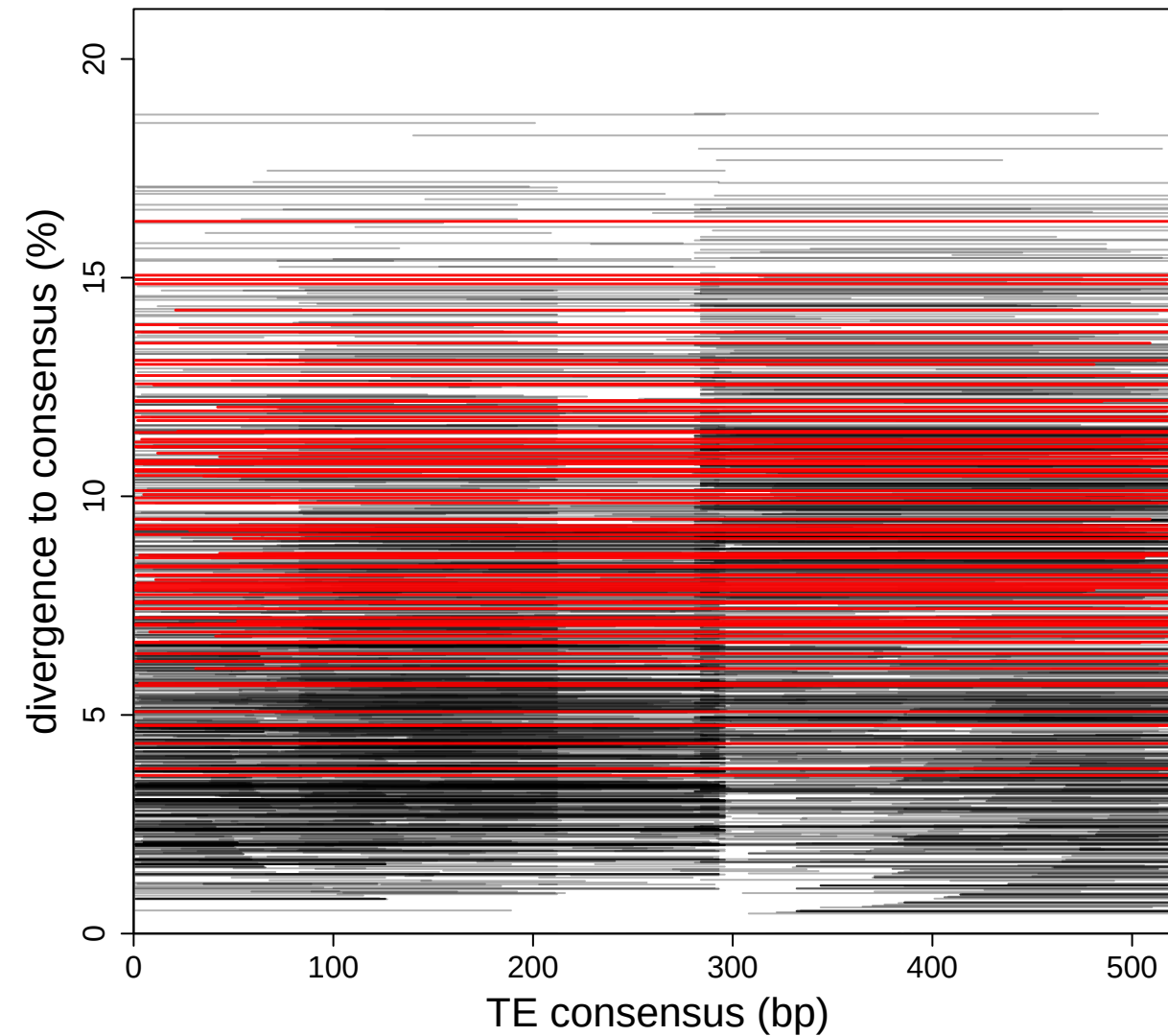


hily_48.fasta.b.bed_uf.bed_g_3.bed_fm_1.bed_0_0_bcln.fa_aln.fa
size: 748bp; fragments: 1925; full length: 0 (>=673.2bp)

After TEtrimmer Extended plot Blue lines are boundaries

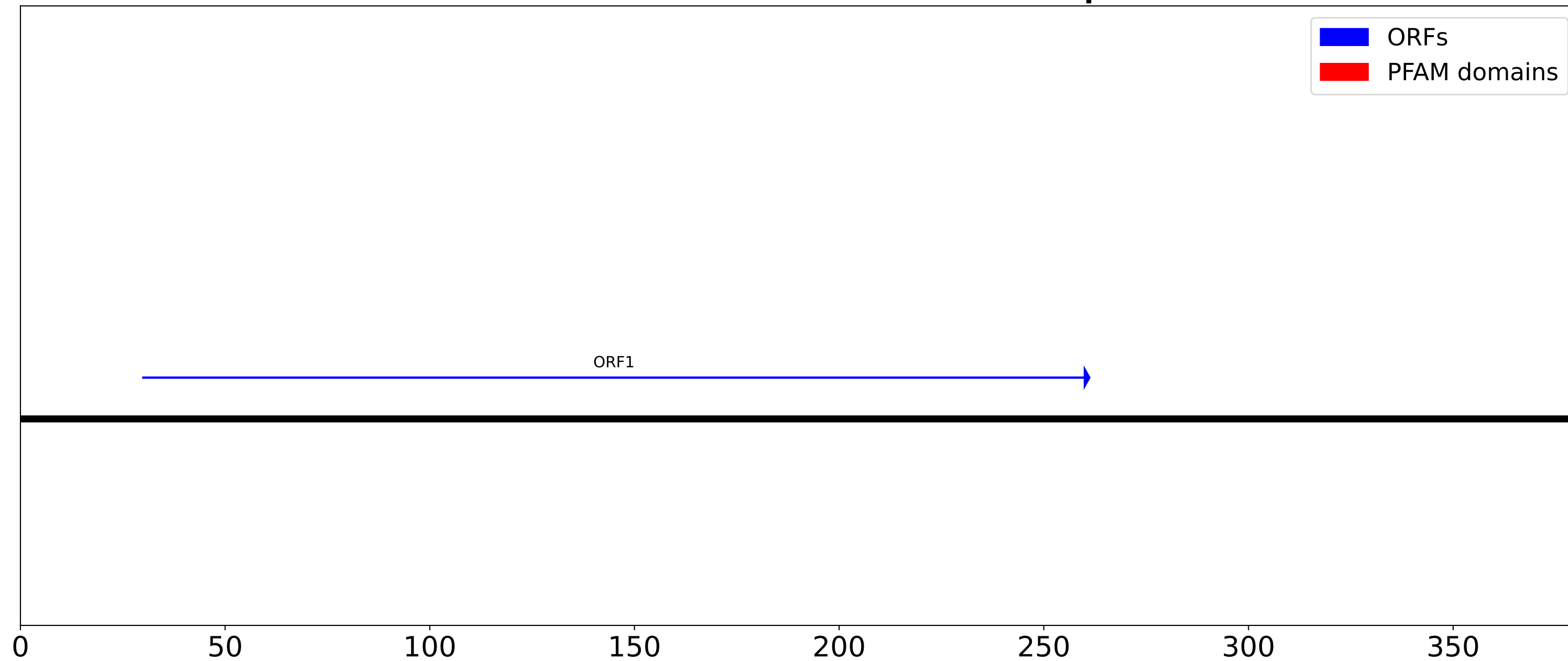


TE: rnd_4_family_48
size: 525bp; fragments: 2732; full length: 85 (≥ 472.5 bp)

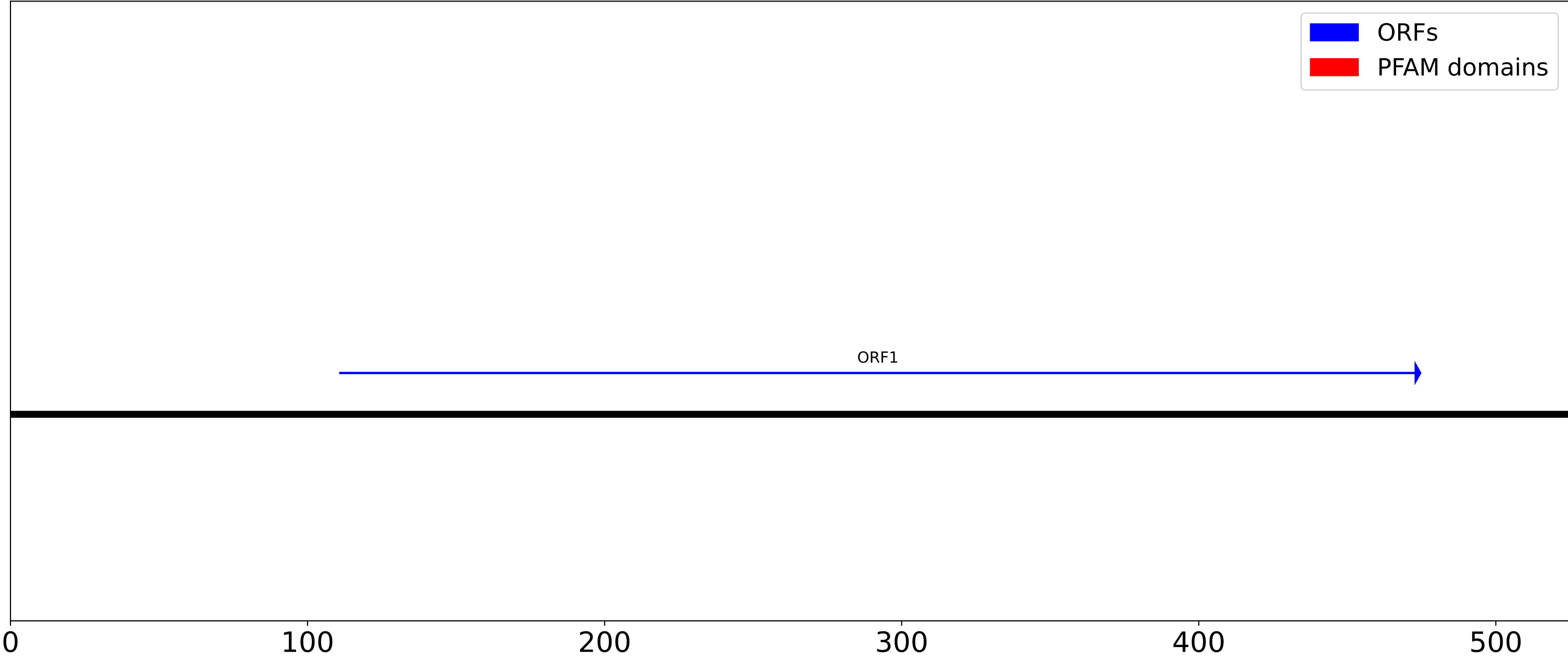


Before TEtrimmer 525 bp

After TEtrimmer ORF and PFAM domain plot



Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

