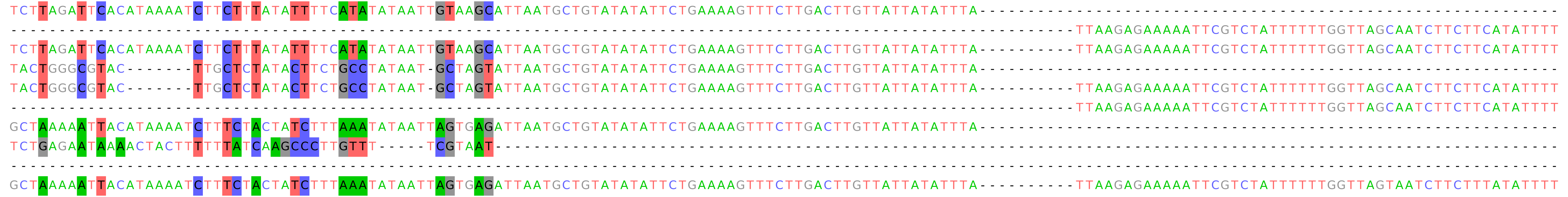
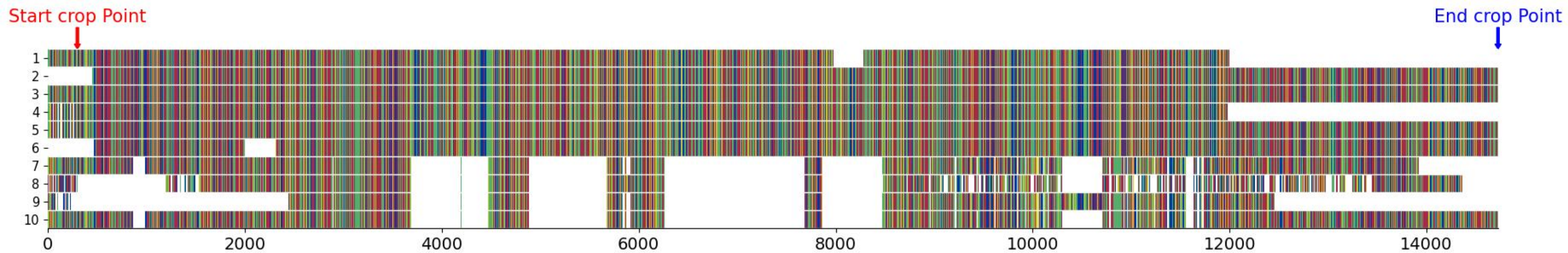


Start crop Point

End crop Point

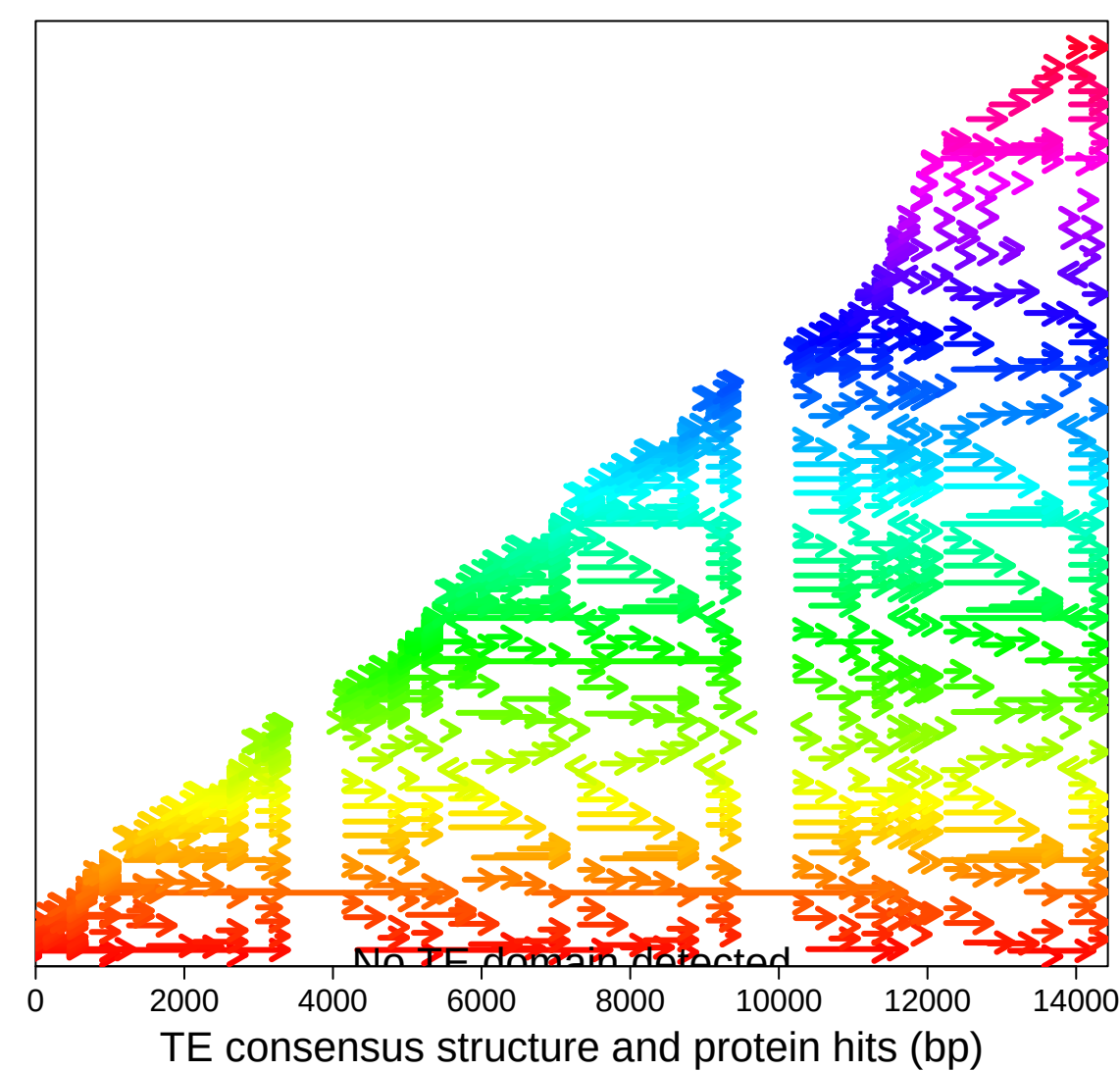
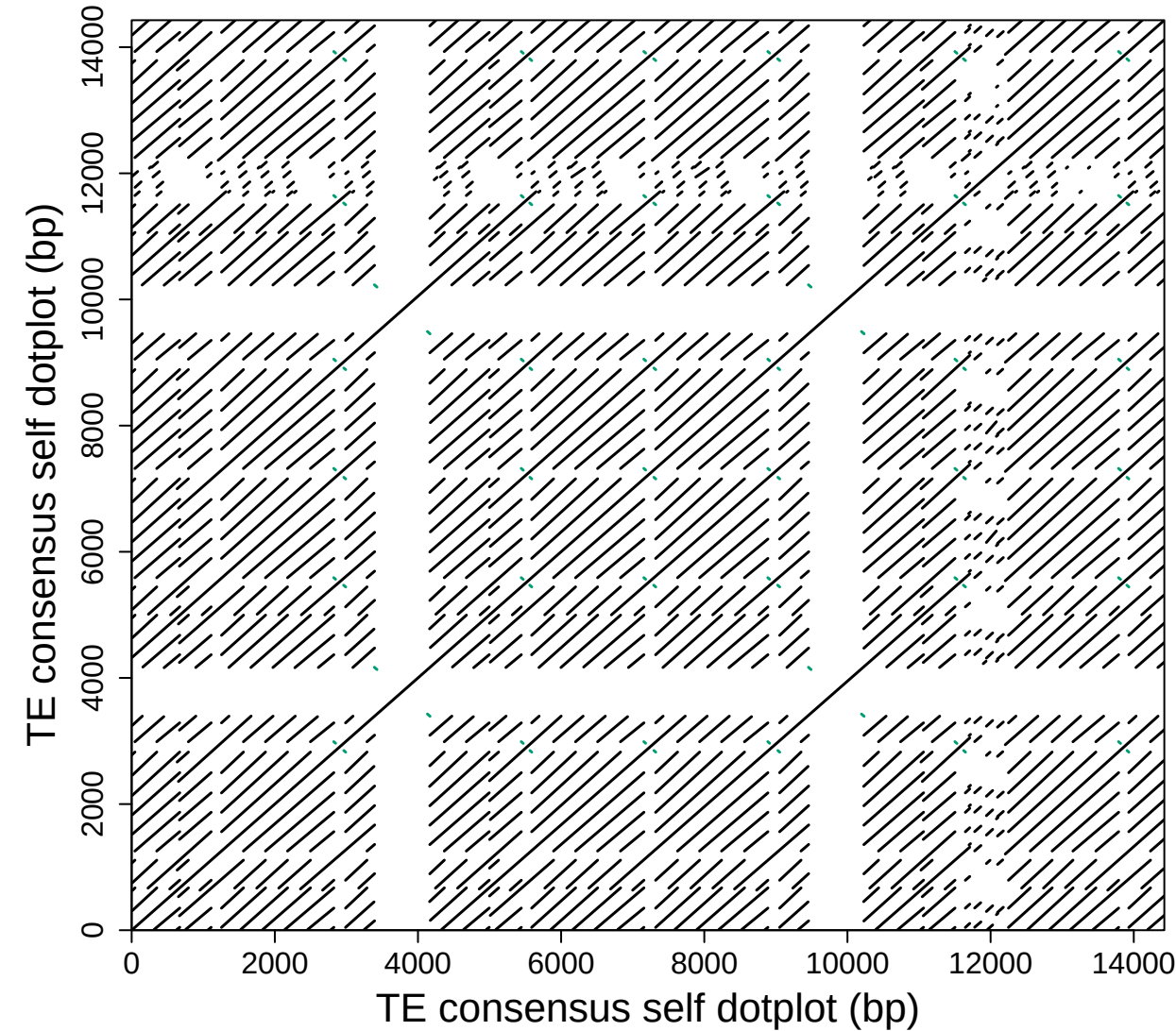
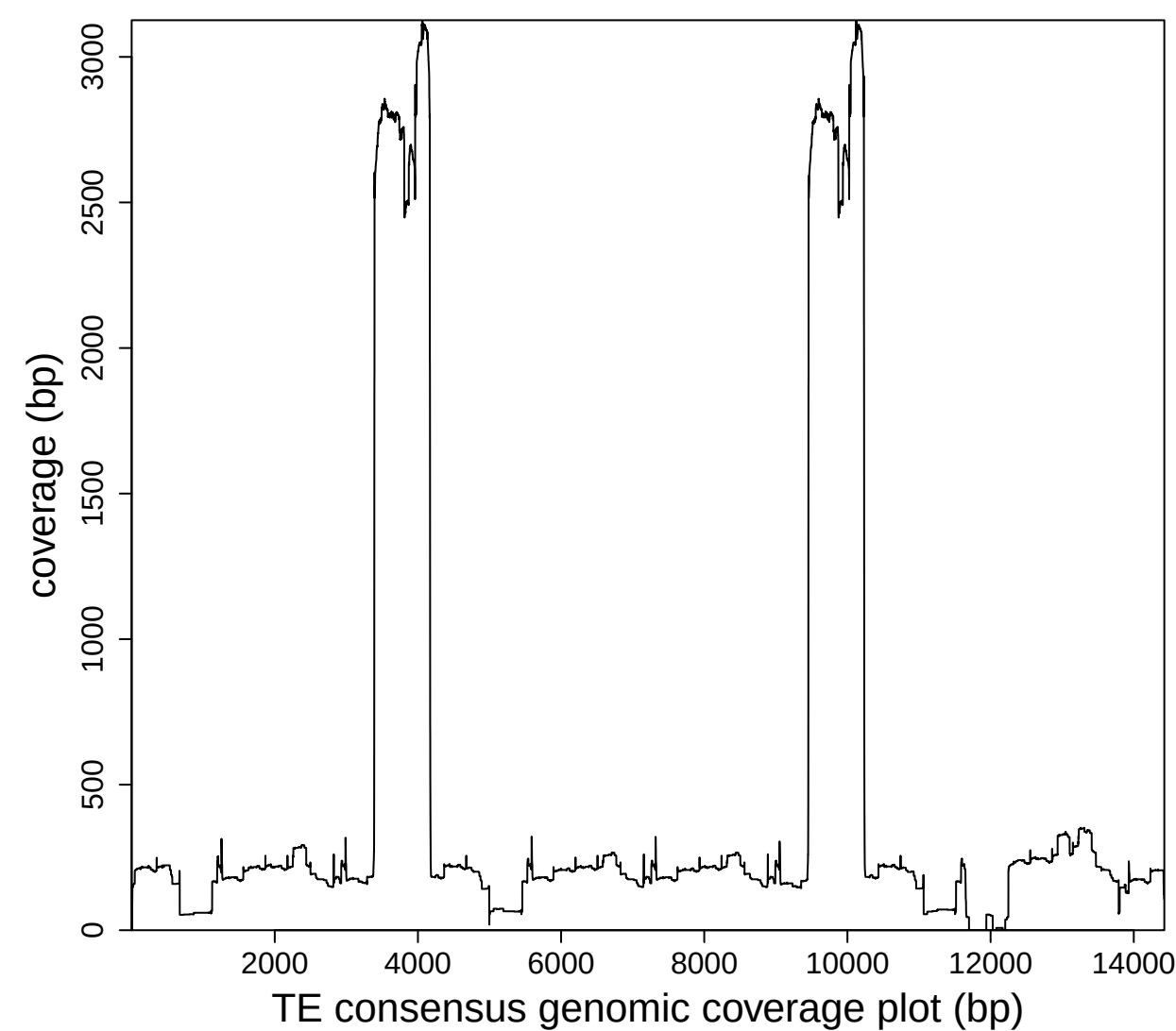
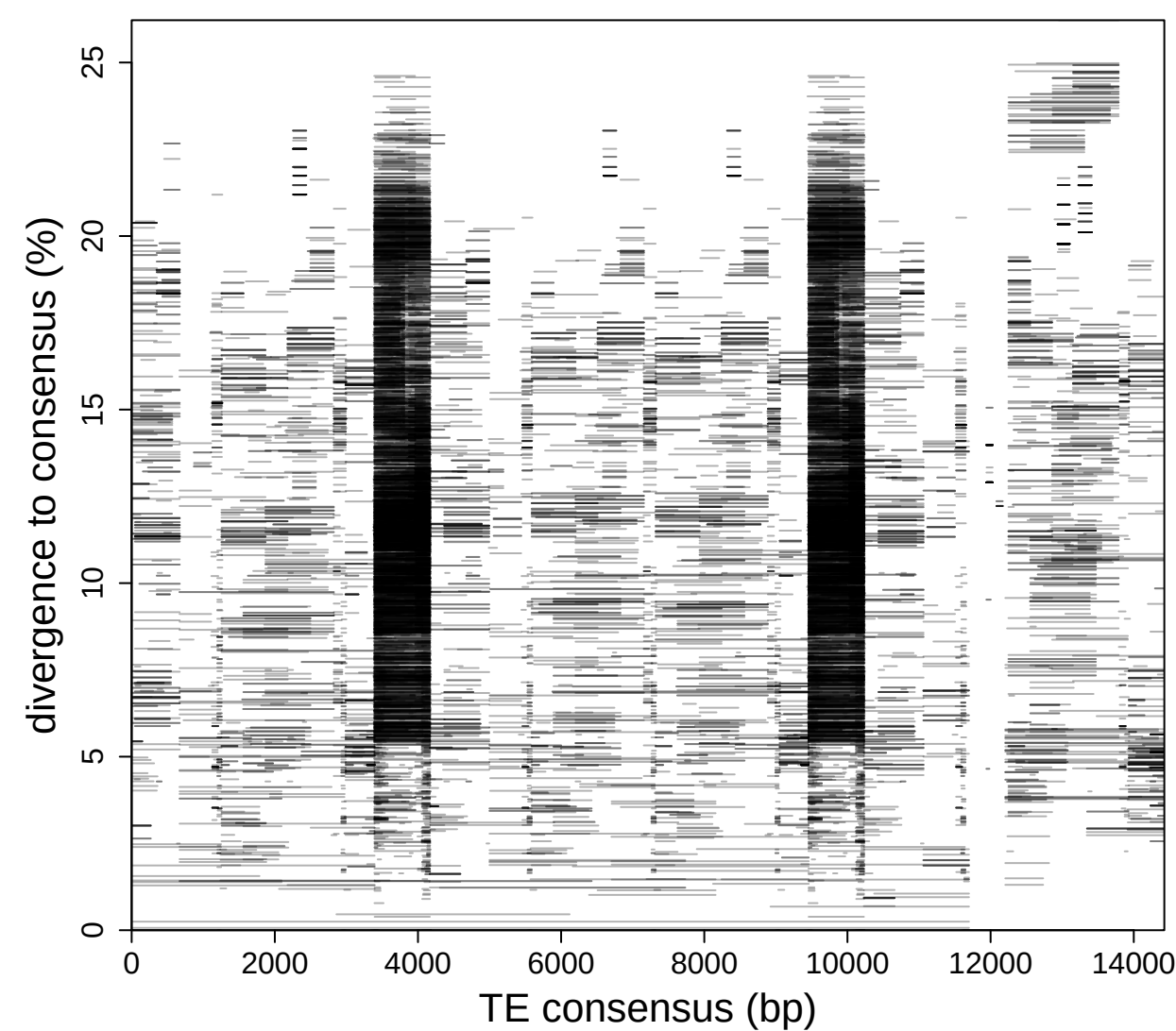
MSA length = 14427





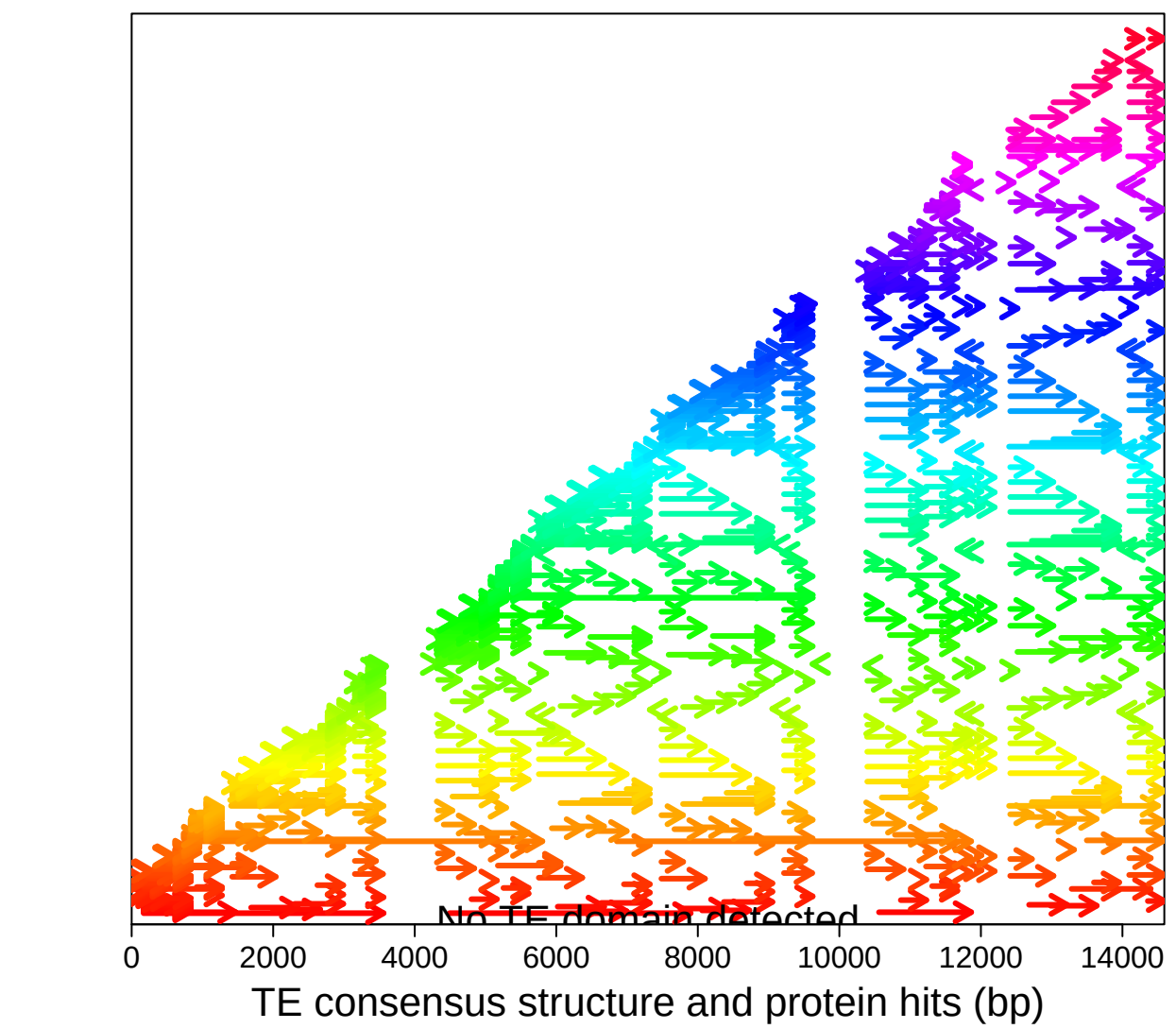
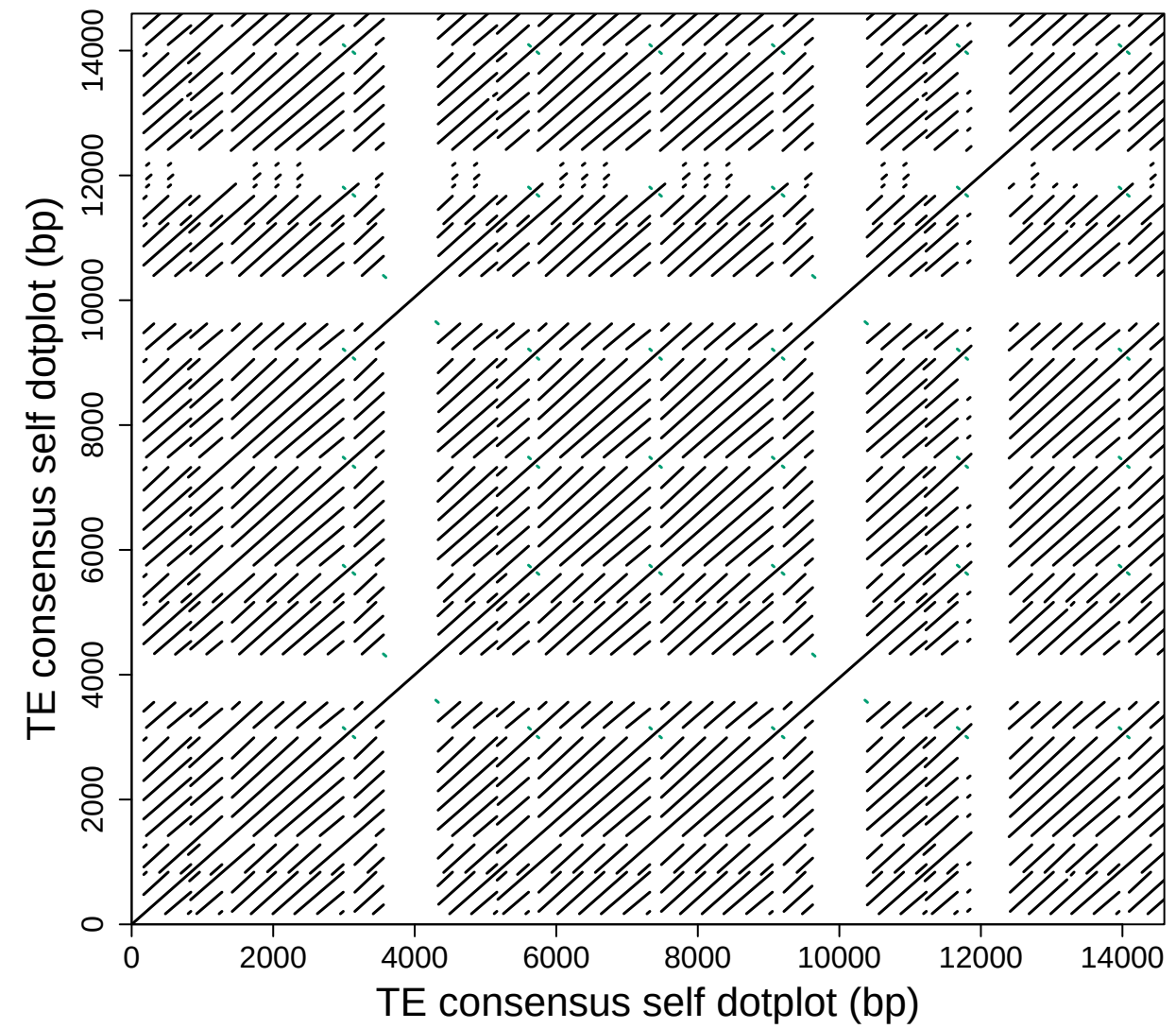
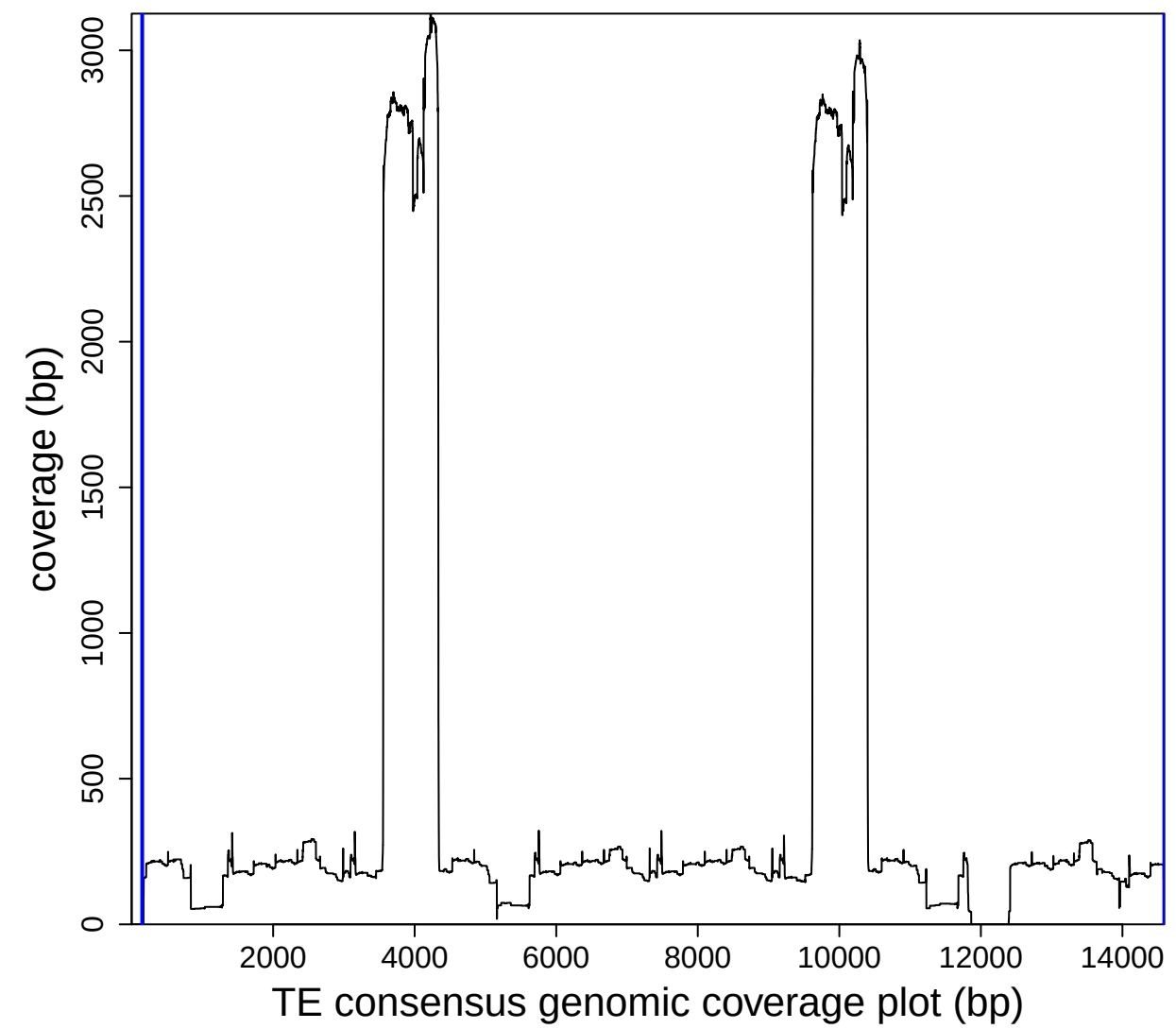
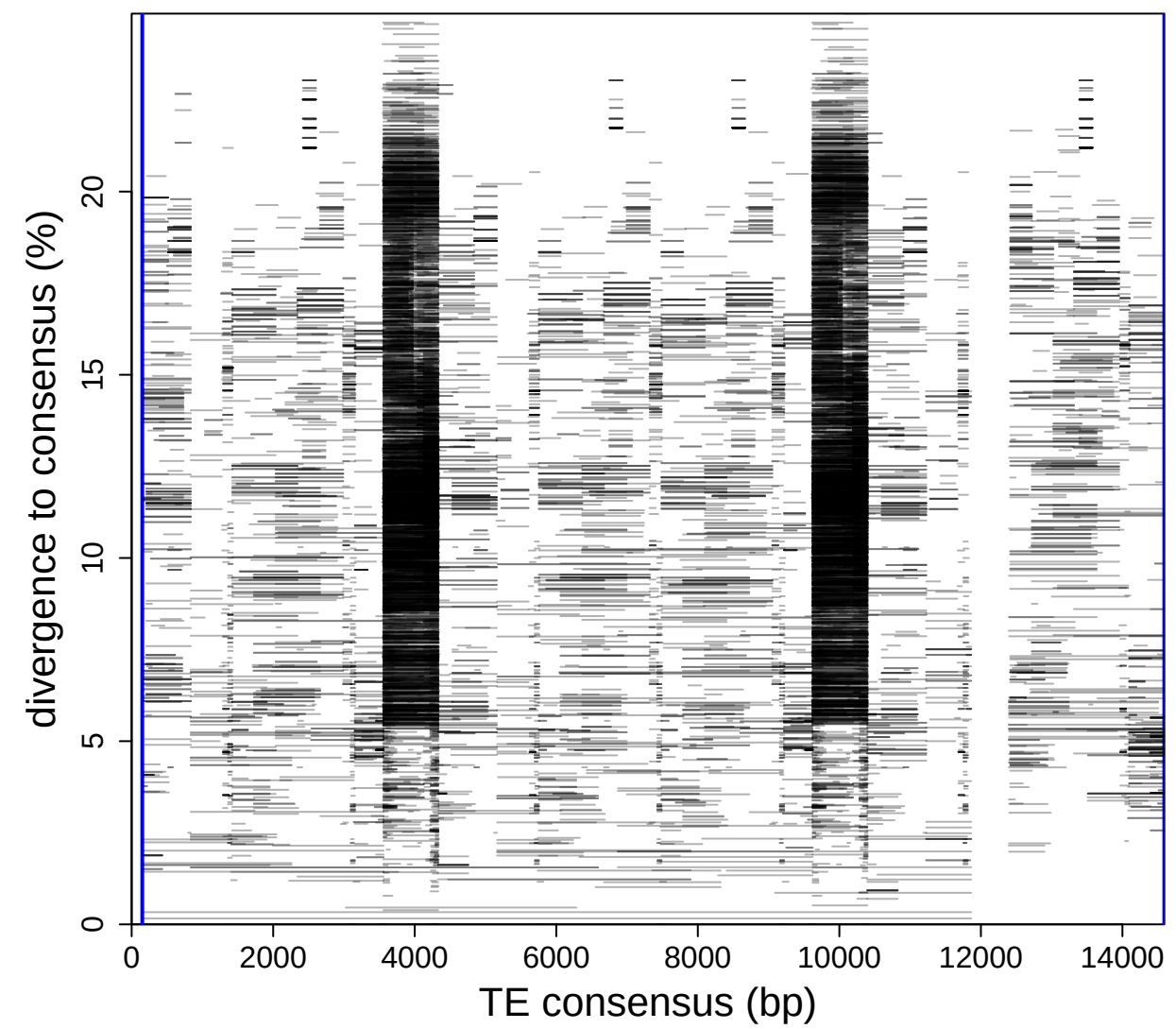
ed_fm_1.bed 0 0_n.bed_g_1.bed_fm_2.bed 0 0_bcIn.fa_aln.fa
size: 14427bp; fragments: 14973; full length: 0 (>=12984.3bp)

After TEtrimmer 14427 bp

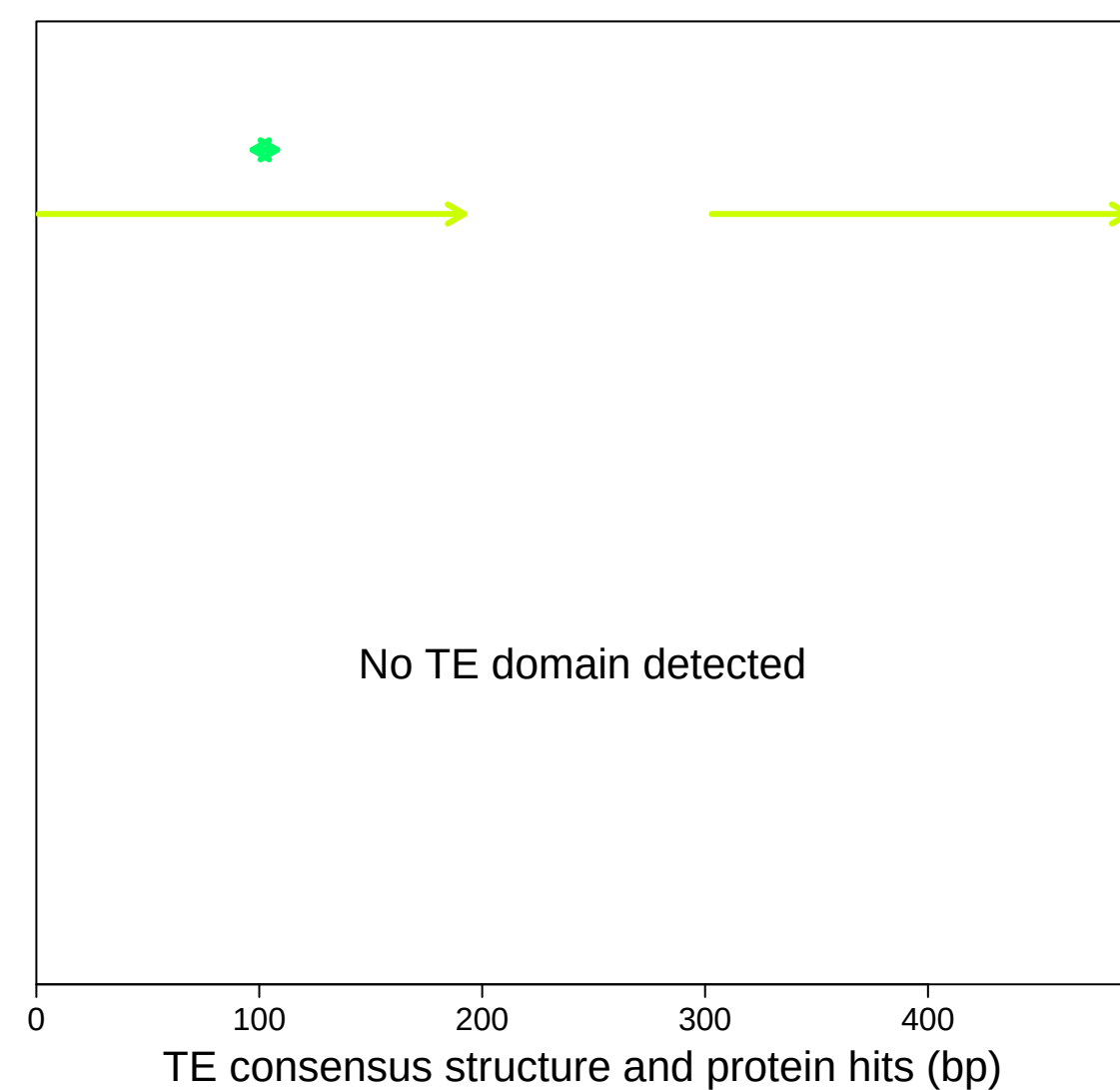
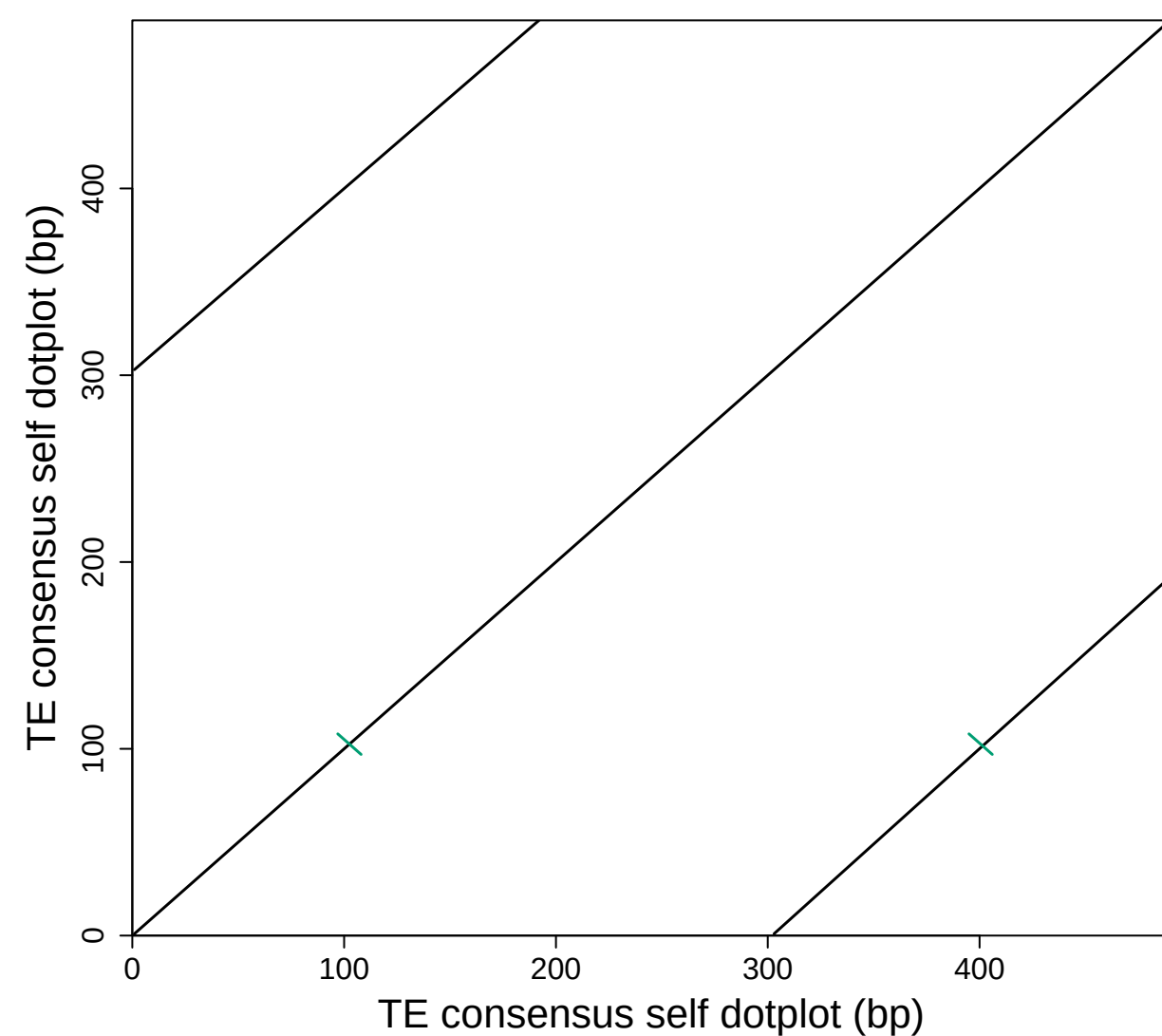
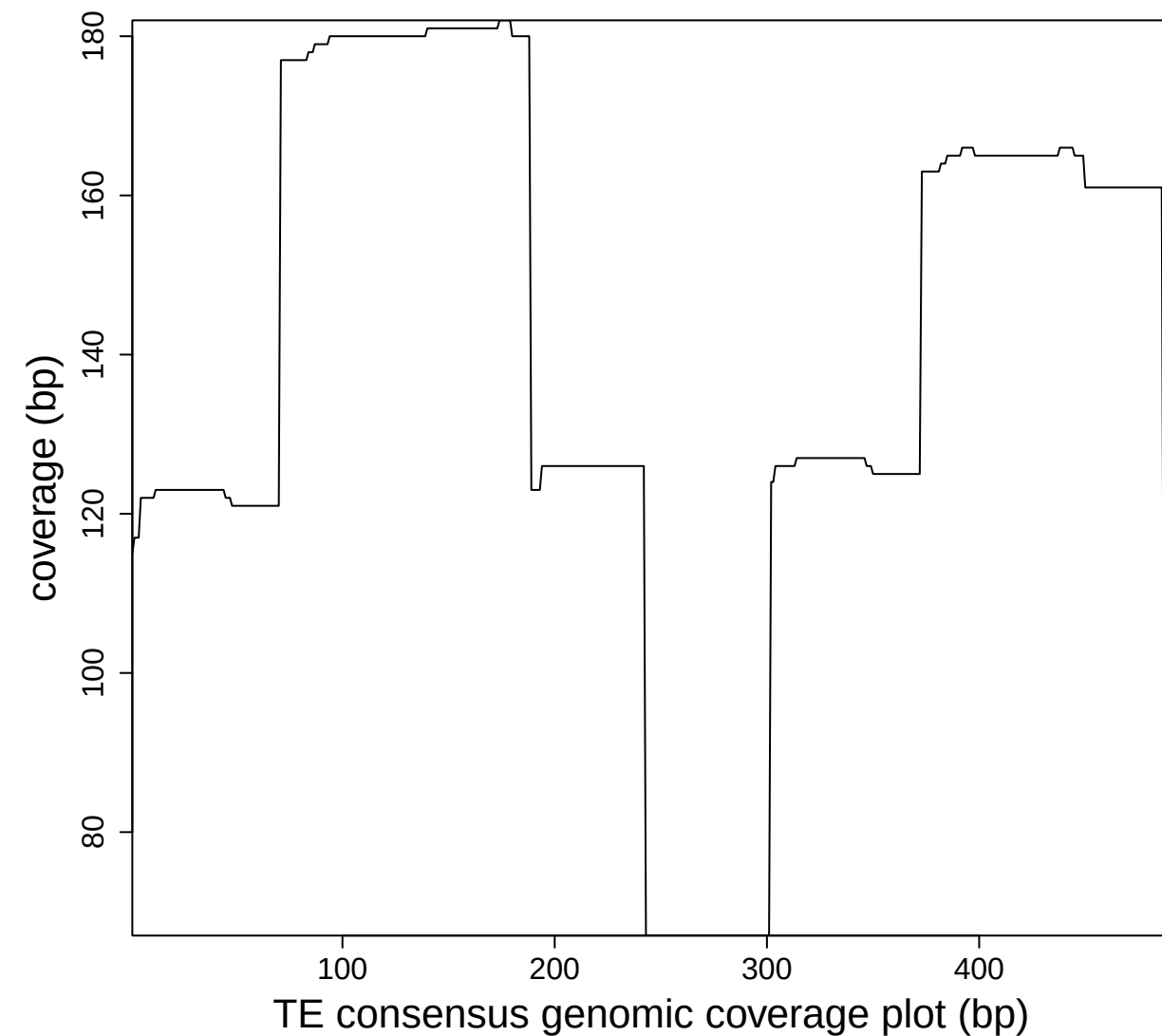
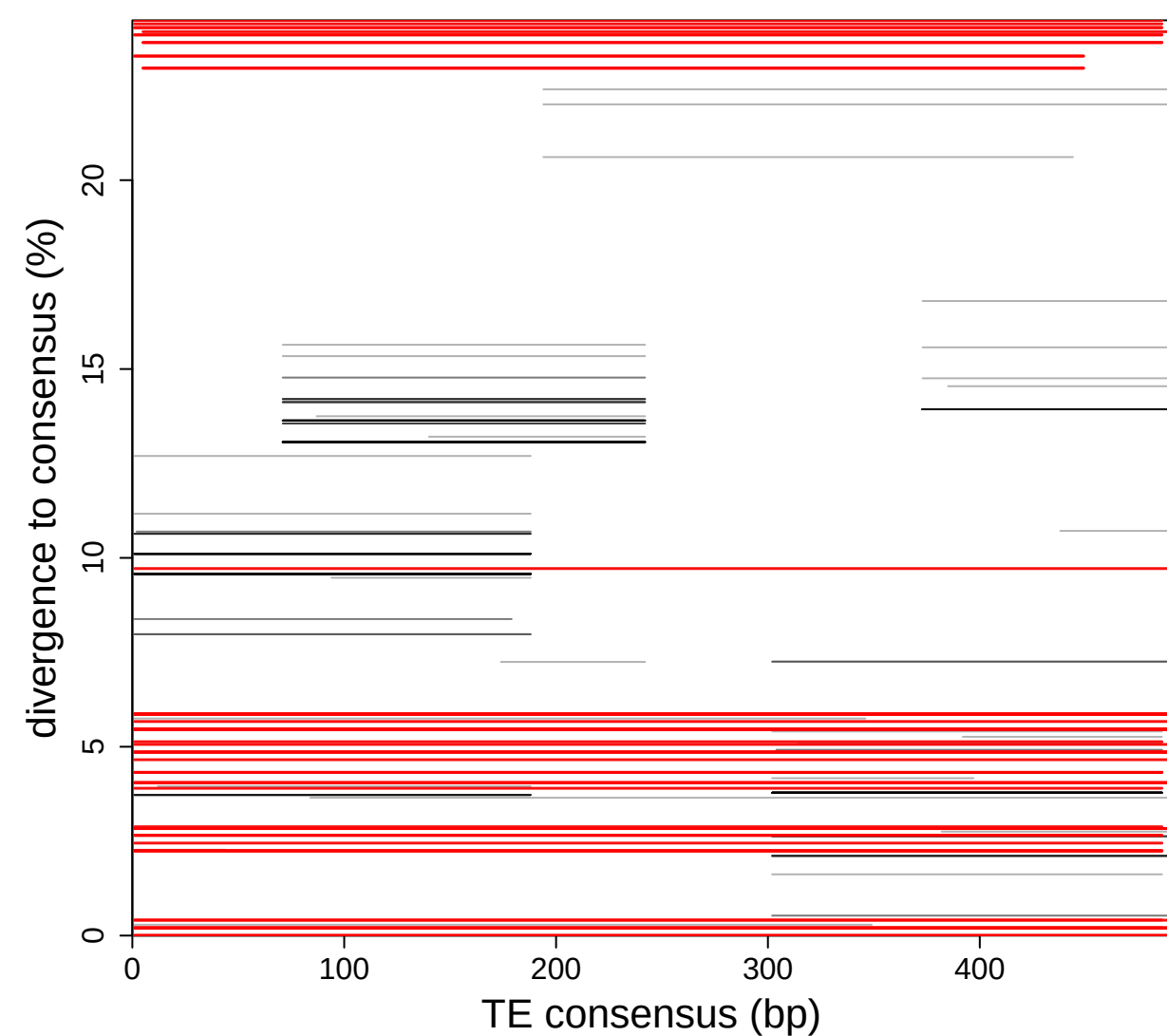


ed_uf.bed_g_2.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_
size: 14592bp; fragments: 14647; full length: 0 (>=13132.8bp)

After TEtrimmer Extended plot Blue lines are boundaries

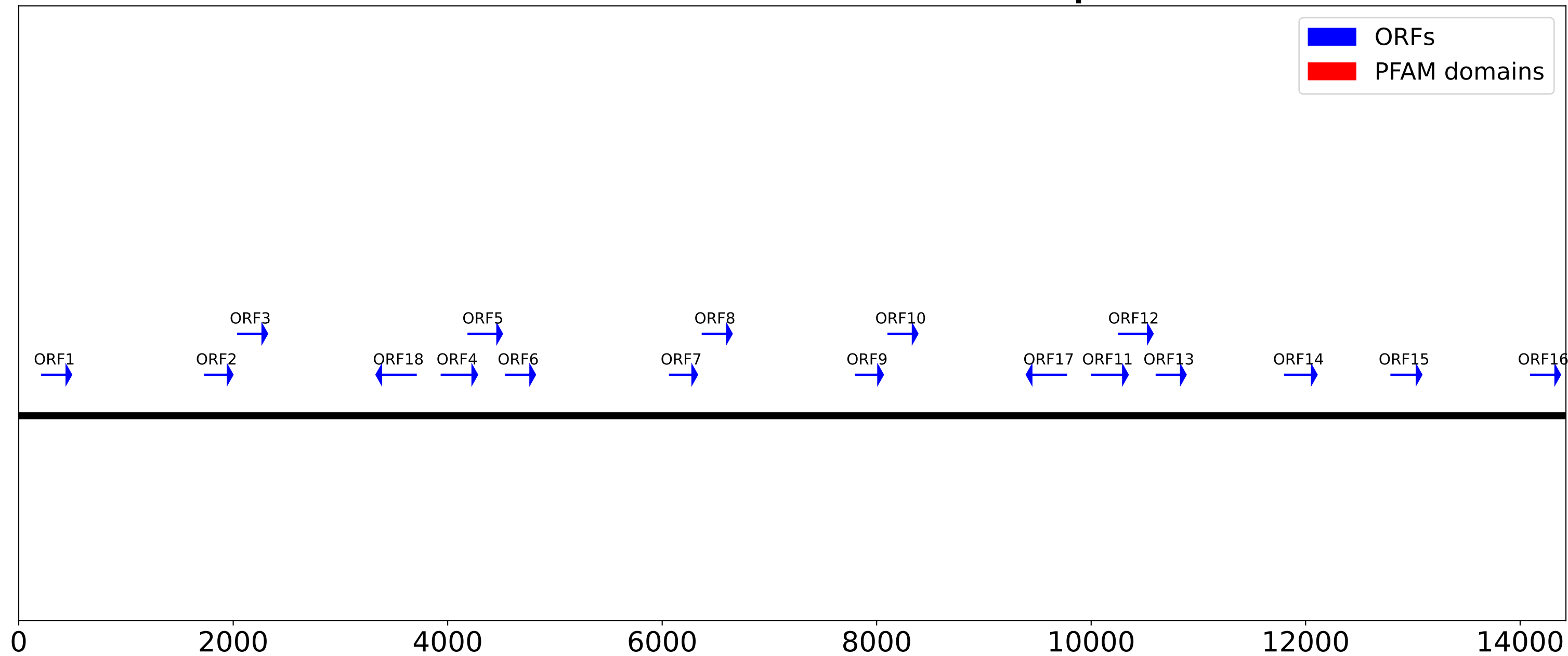


TE: ltr_1_family_68
size: 490bp; fragments: 289; full length: 61 (≥ 441 bp)

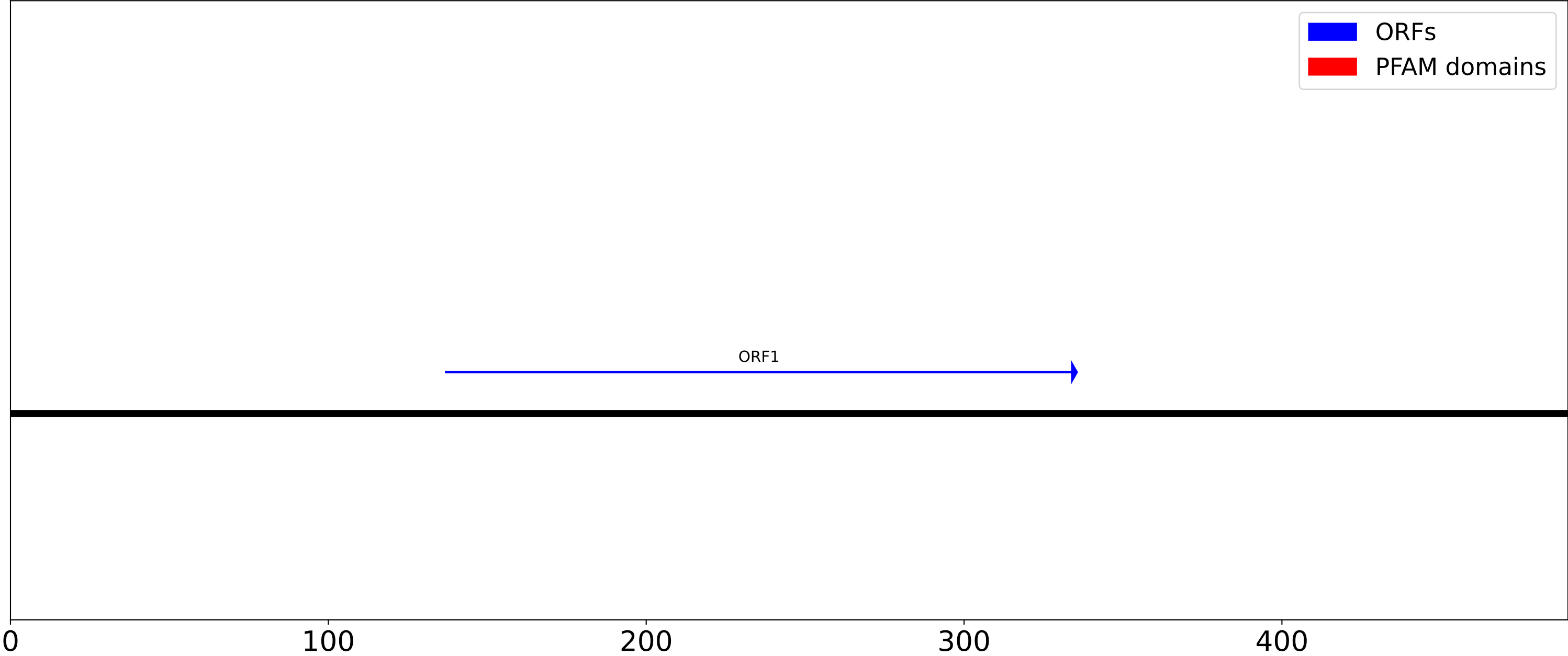


Before TEtrimmer 490 bp

After TEtrimmer ORF and PFAM domain plot



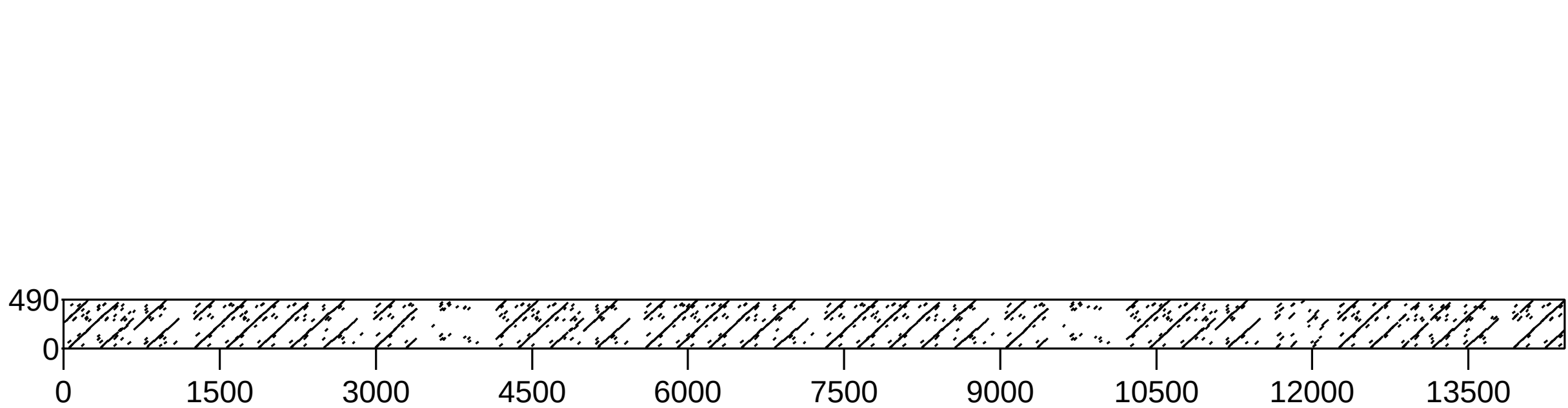
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

nsus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)