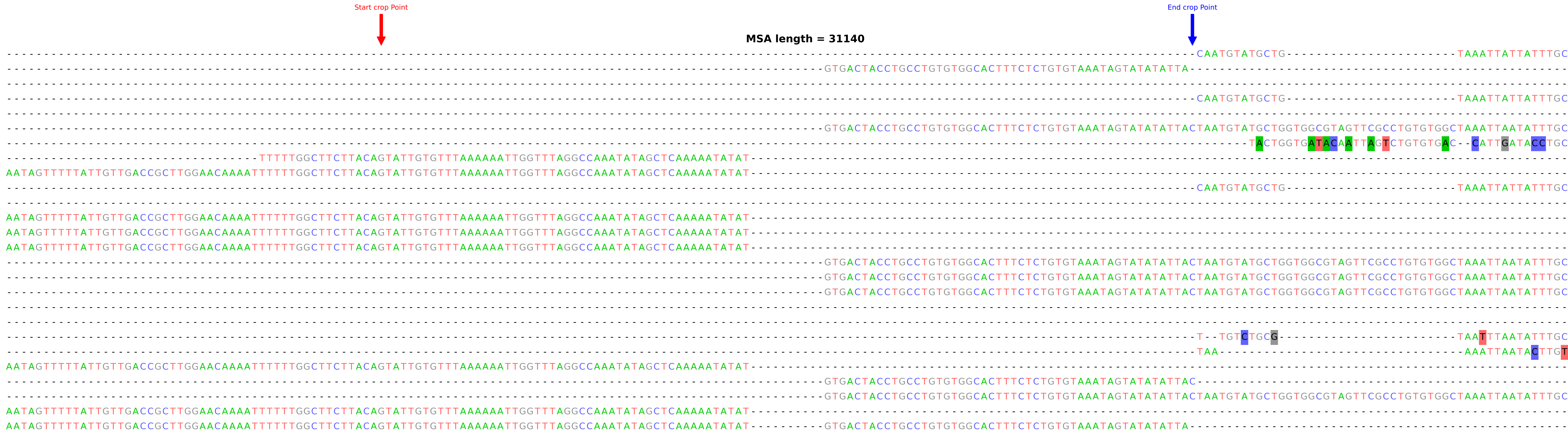
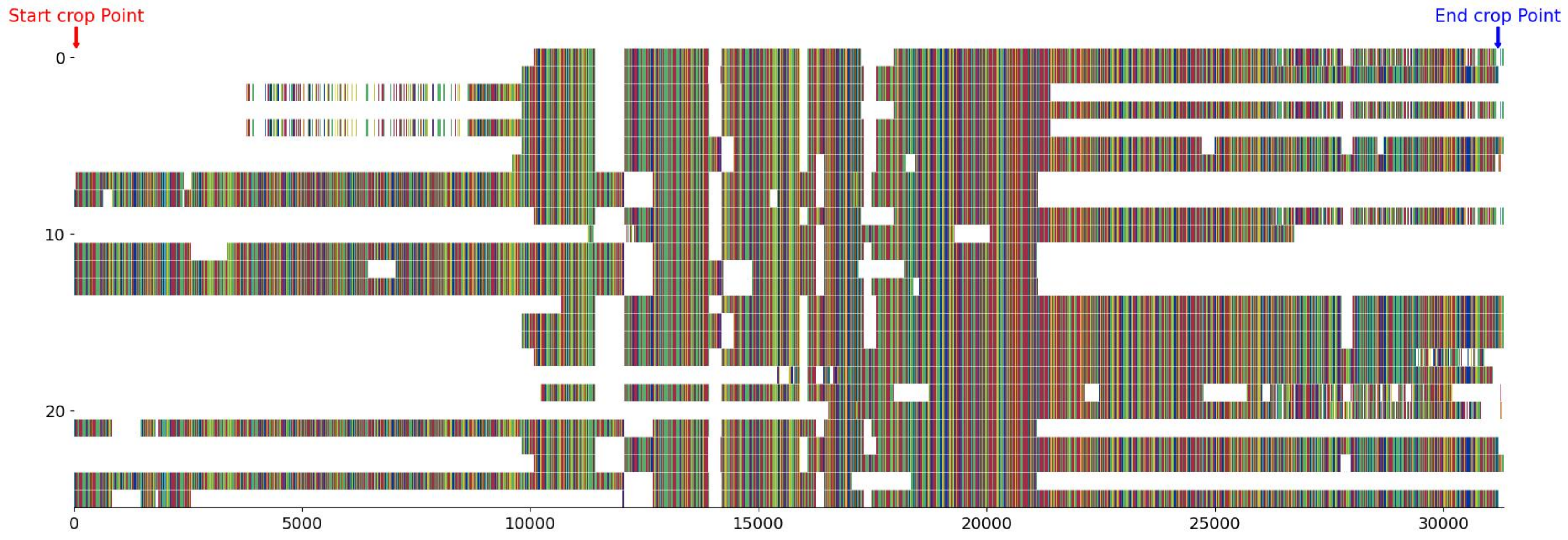
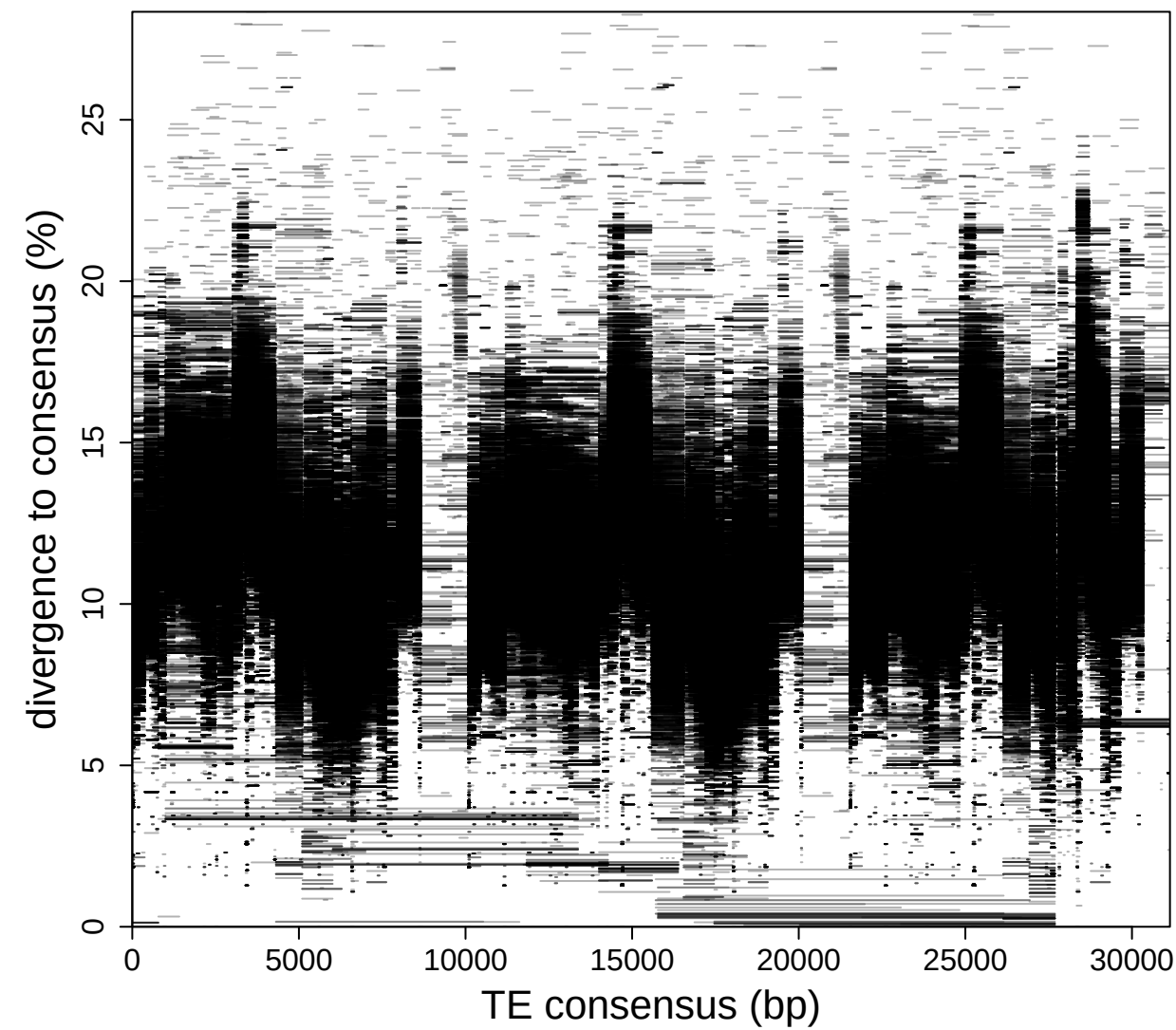


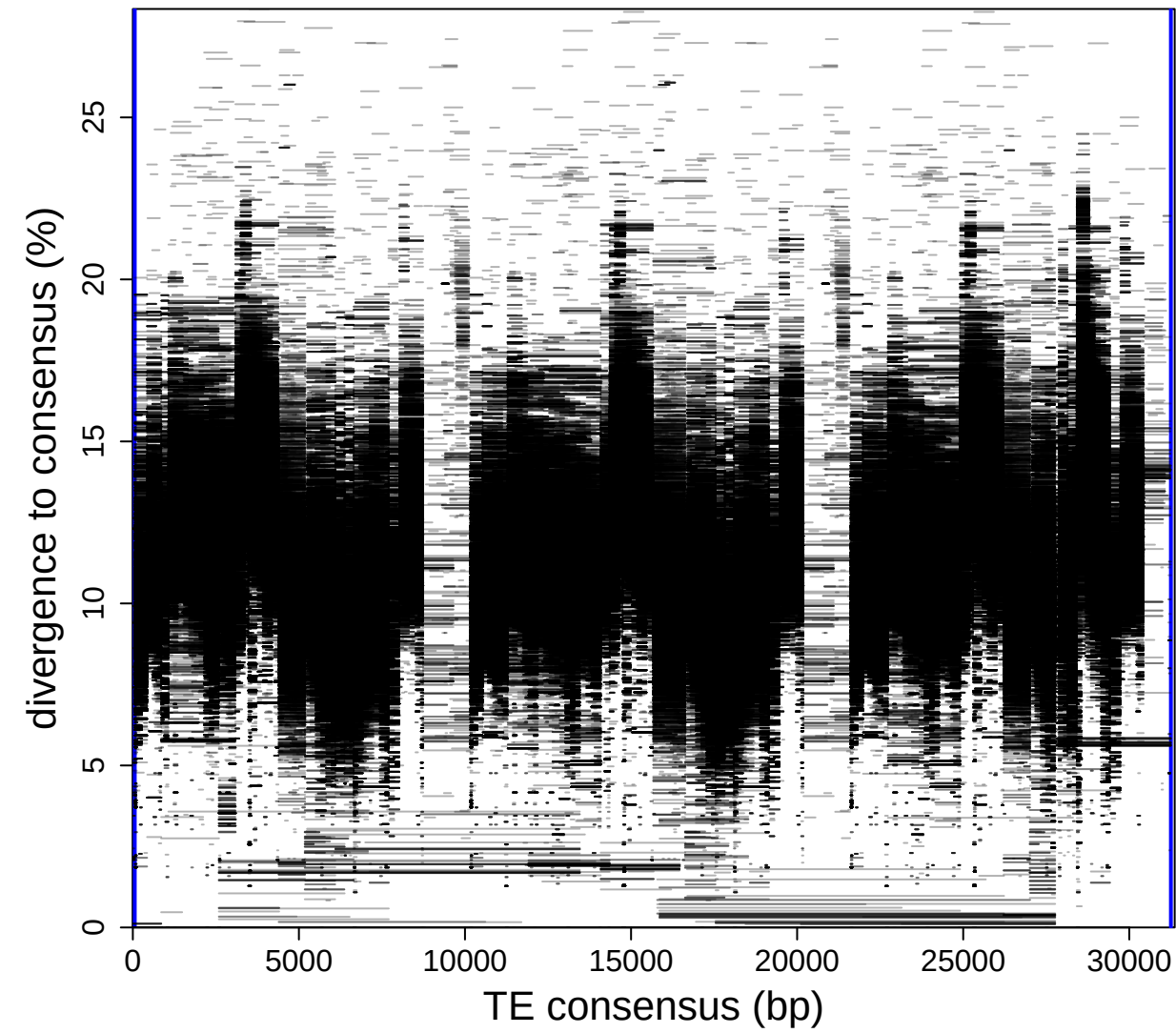




.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcIn.fa_aln.f
size: 31140bp; fragments: 843525; full length: 0 (>=28026bp)

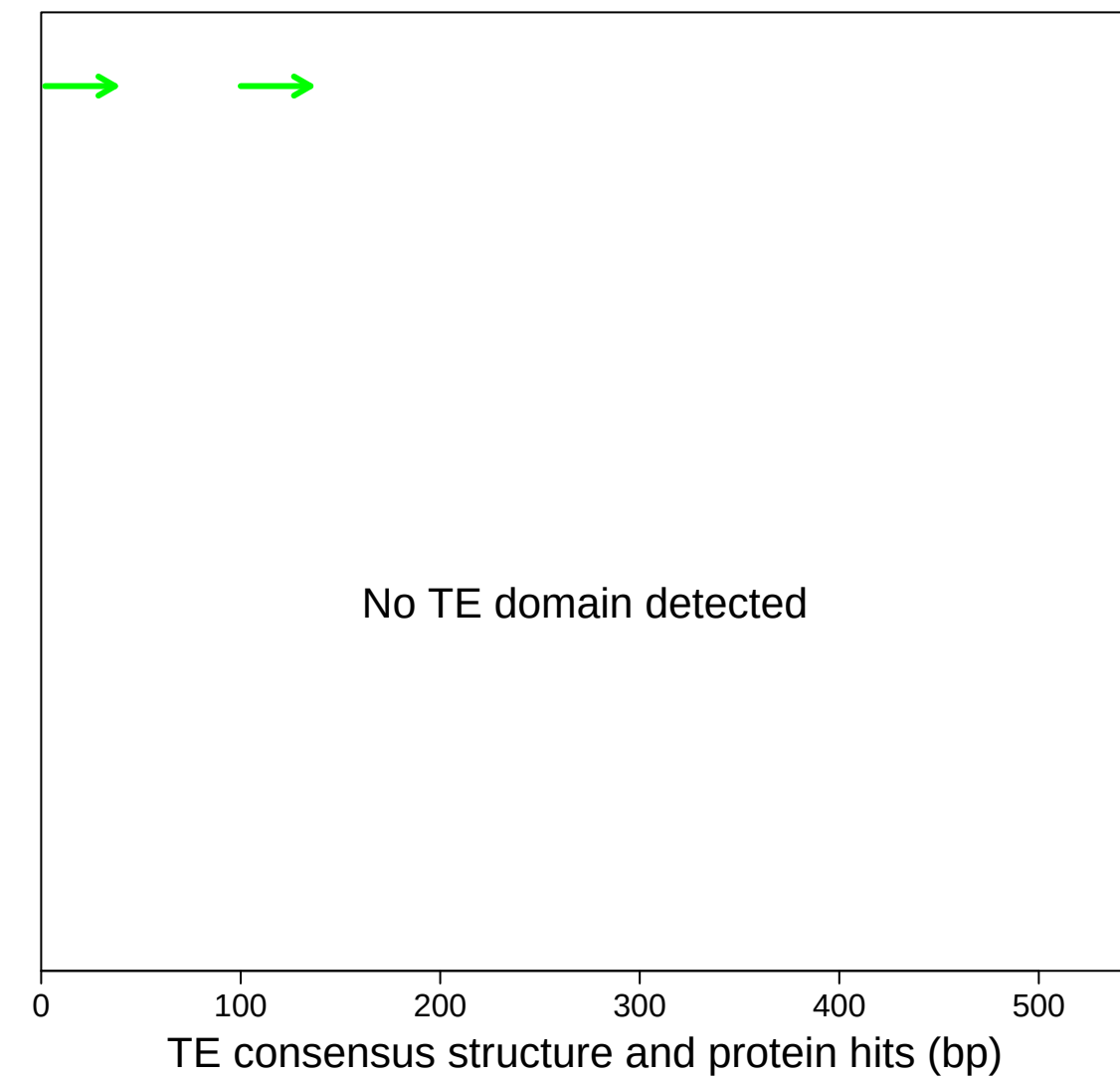
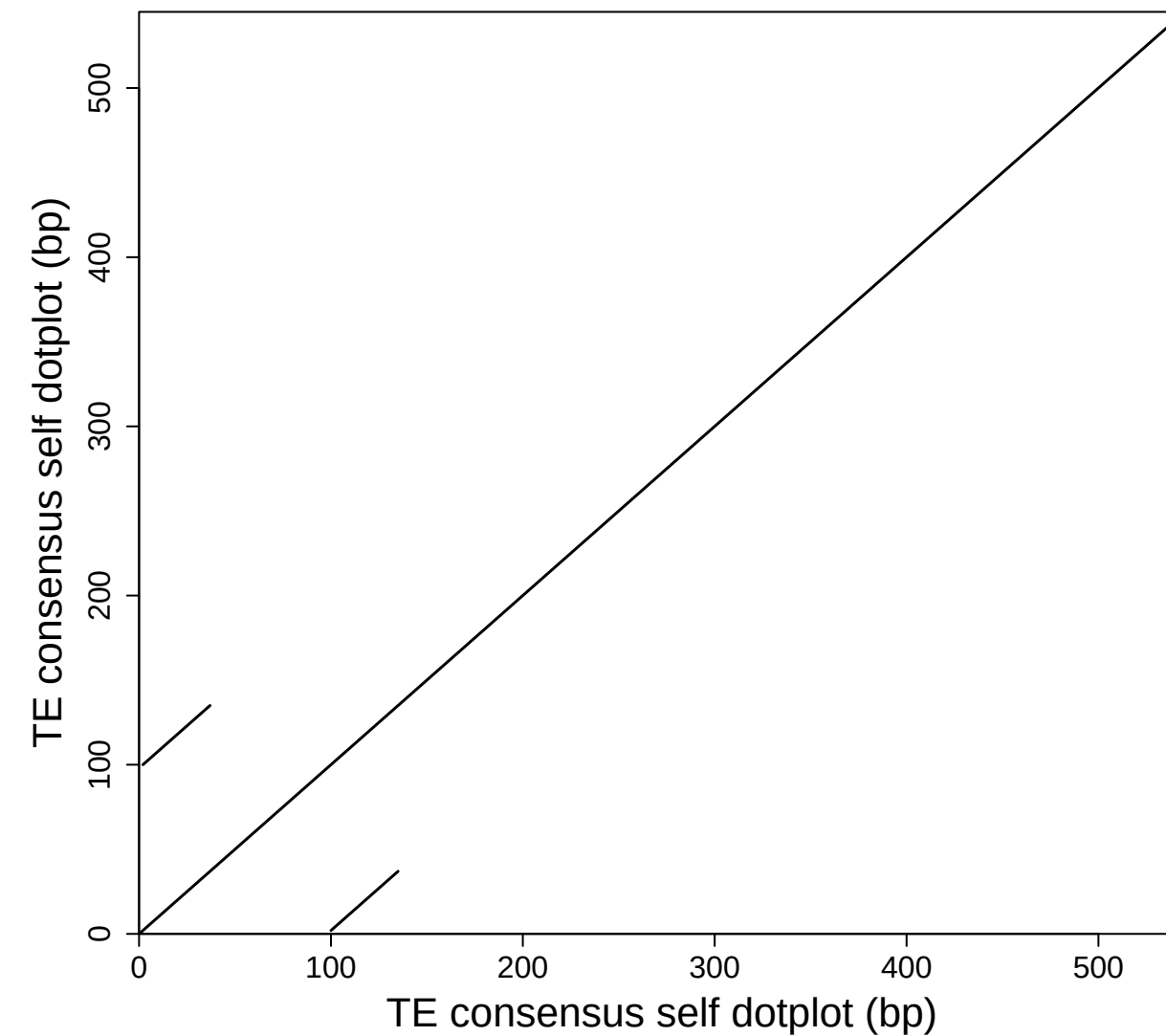
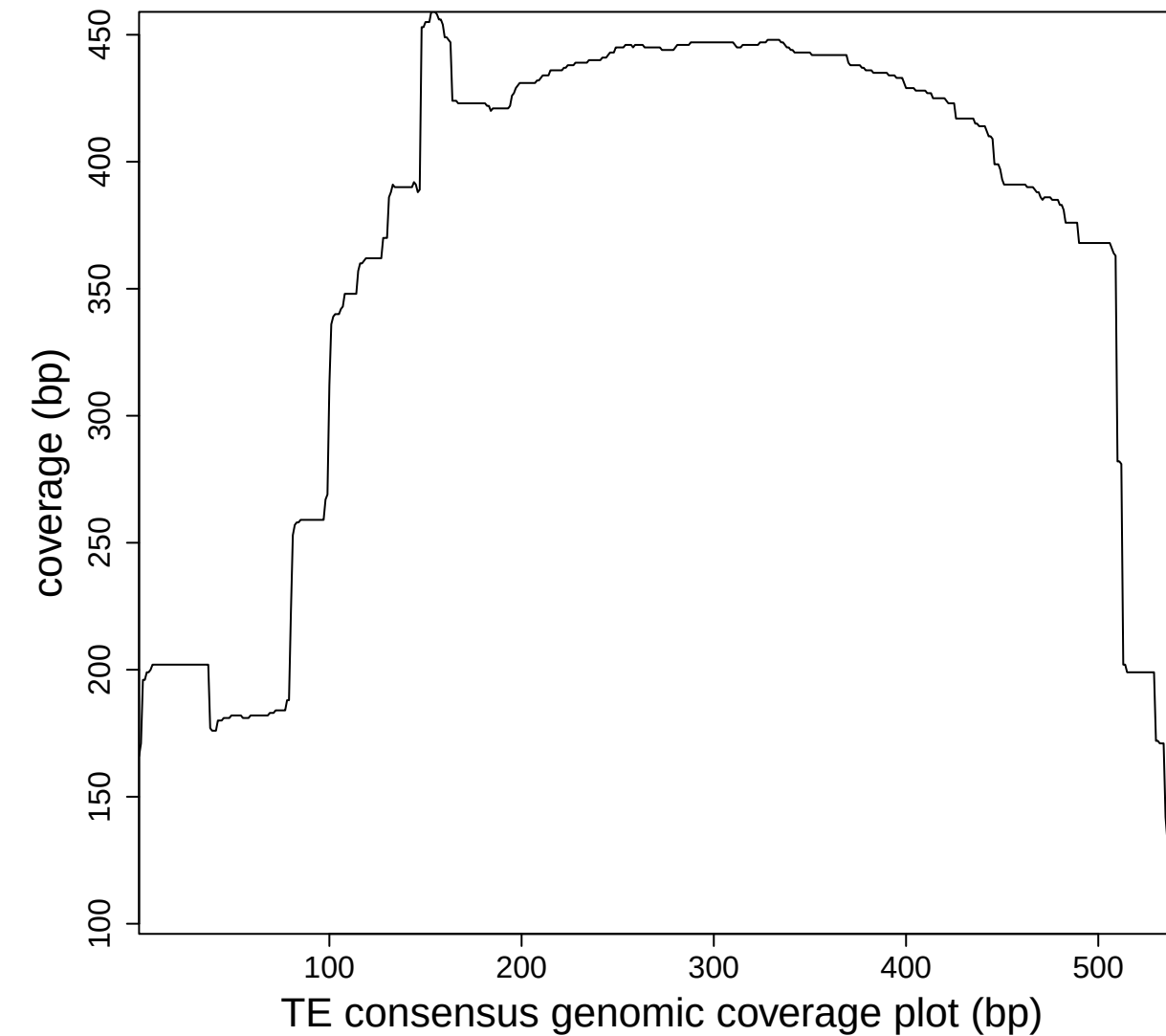
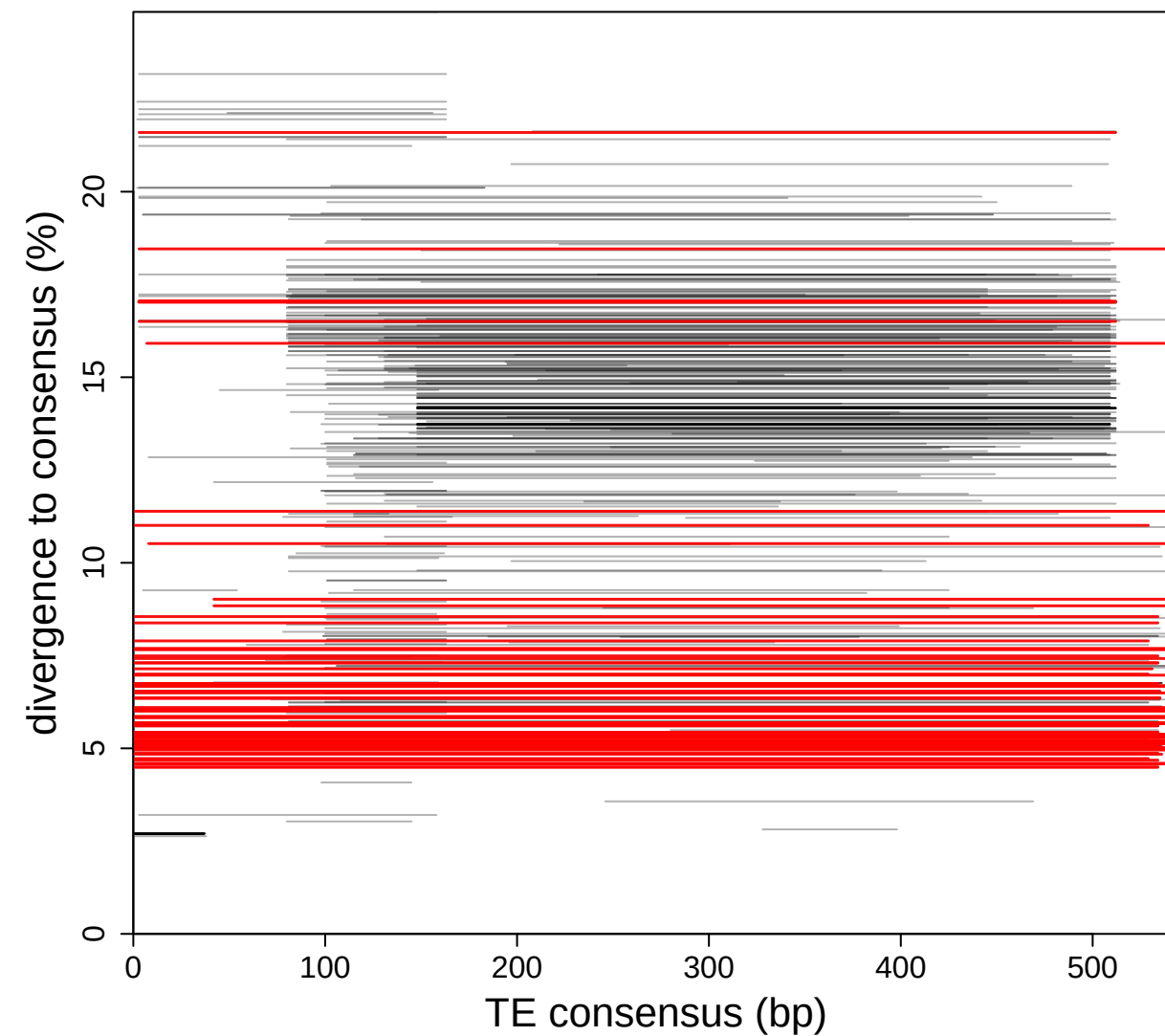


bed_uf.bed_g_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0
size: 31367bp; fragments: 846037; full length: 0 (≥ 28230.3 bp)

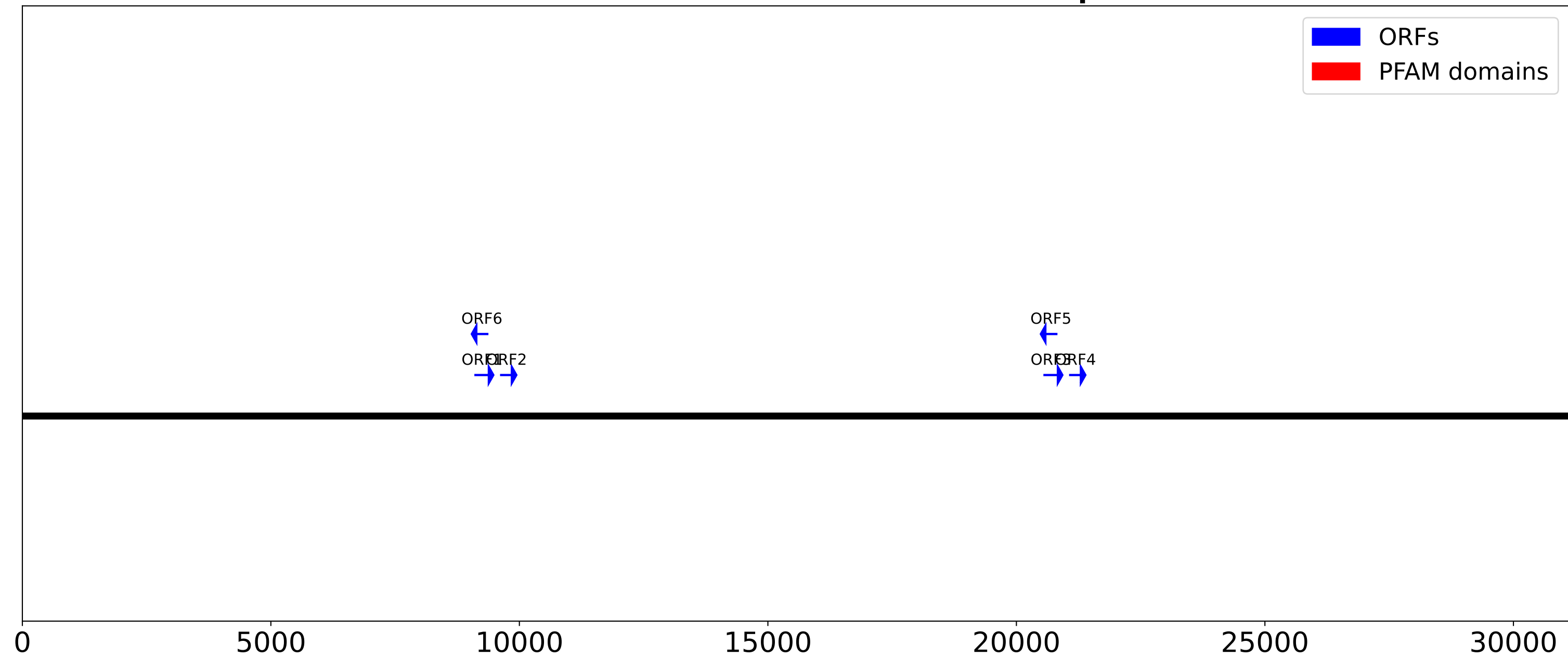


TE: rnd_4_family_3928
size: 545bp; fragments: 526; full length: 152 (≥ 490.5 bp)

Before TEtrimmer 545 bp



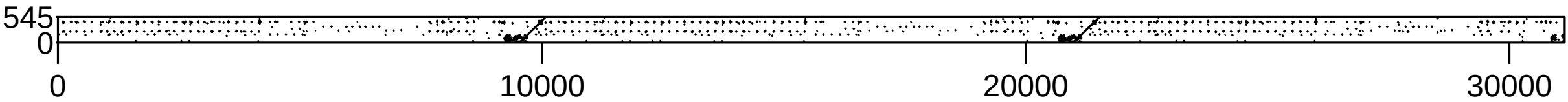
After TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)