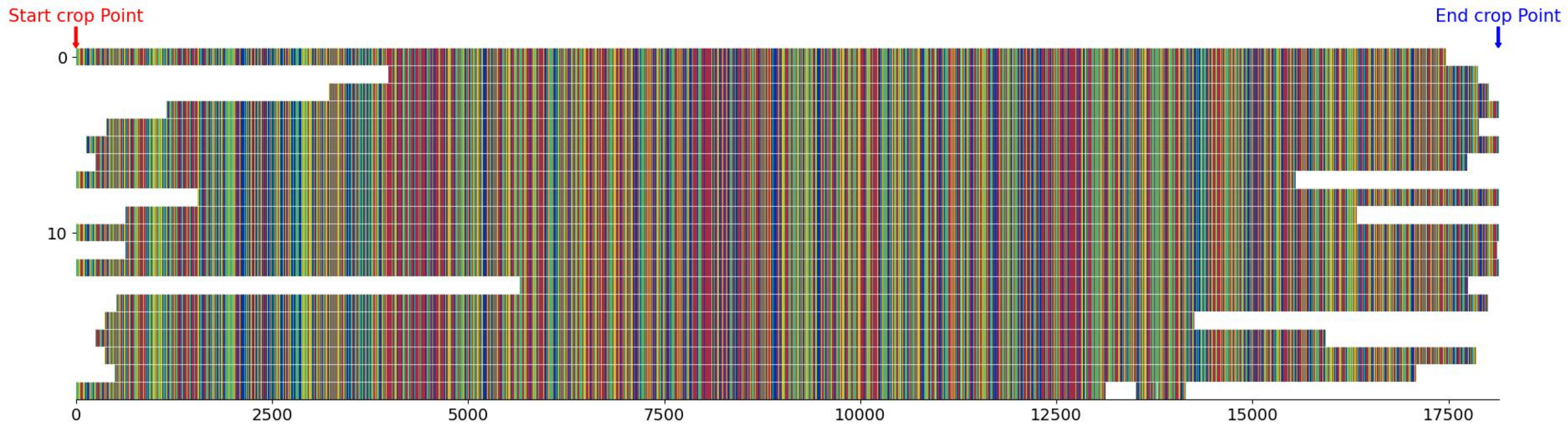


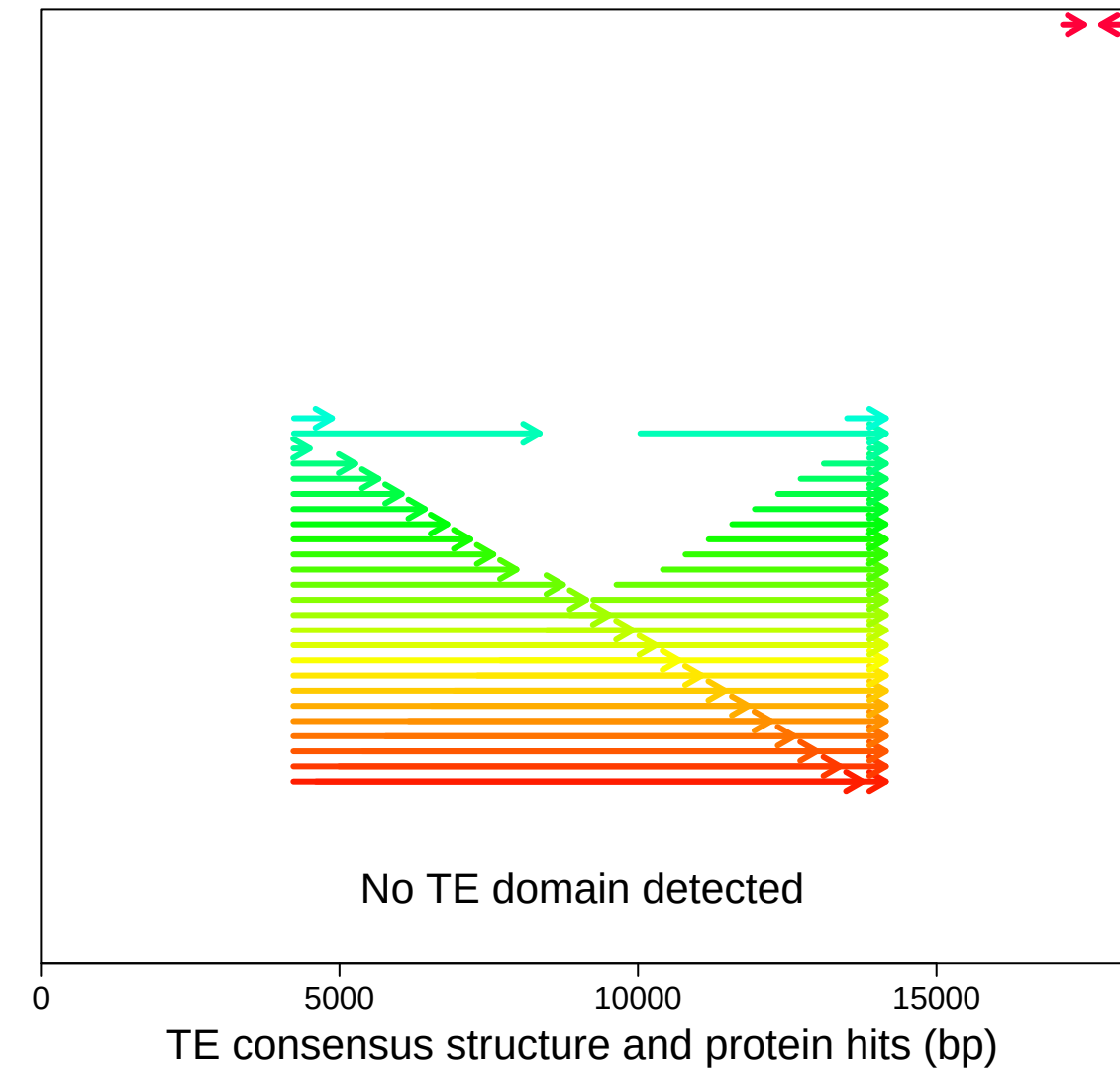
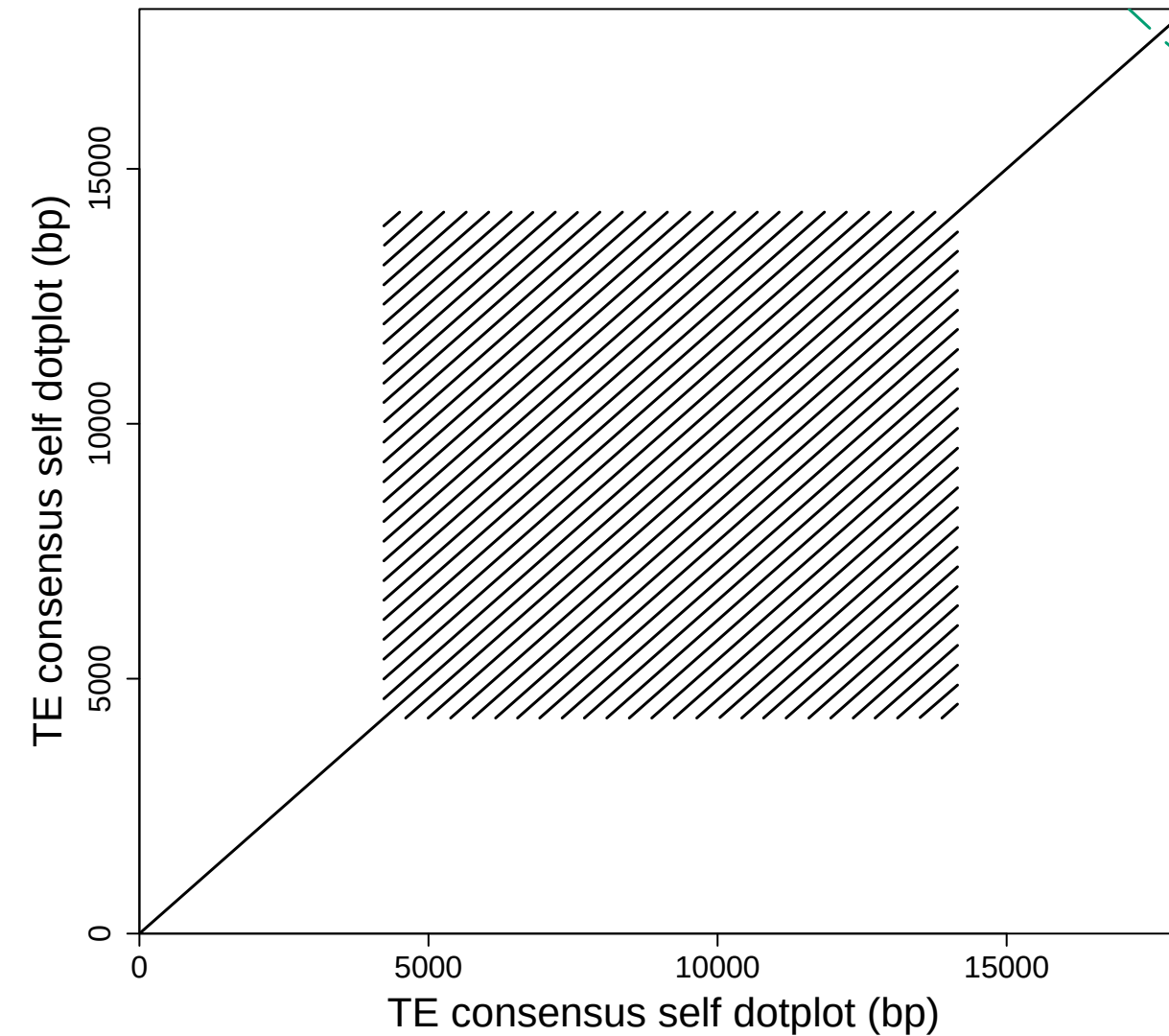
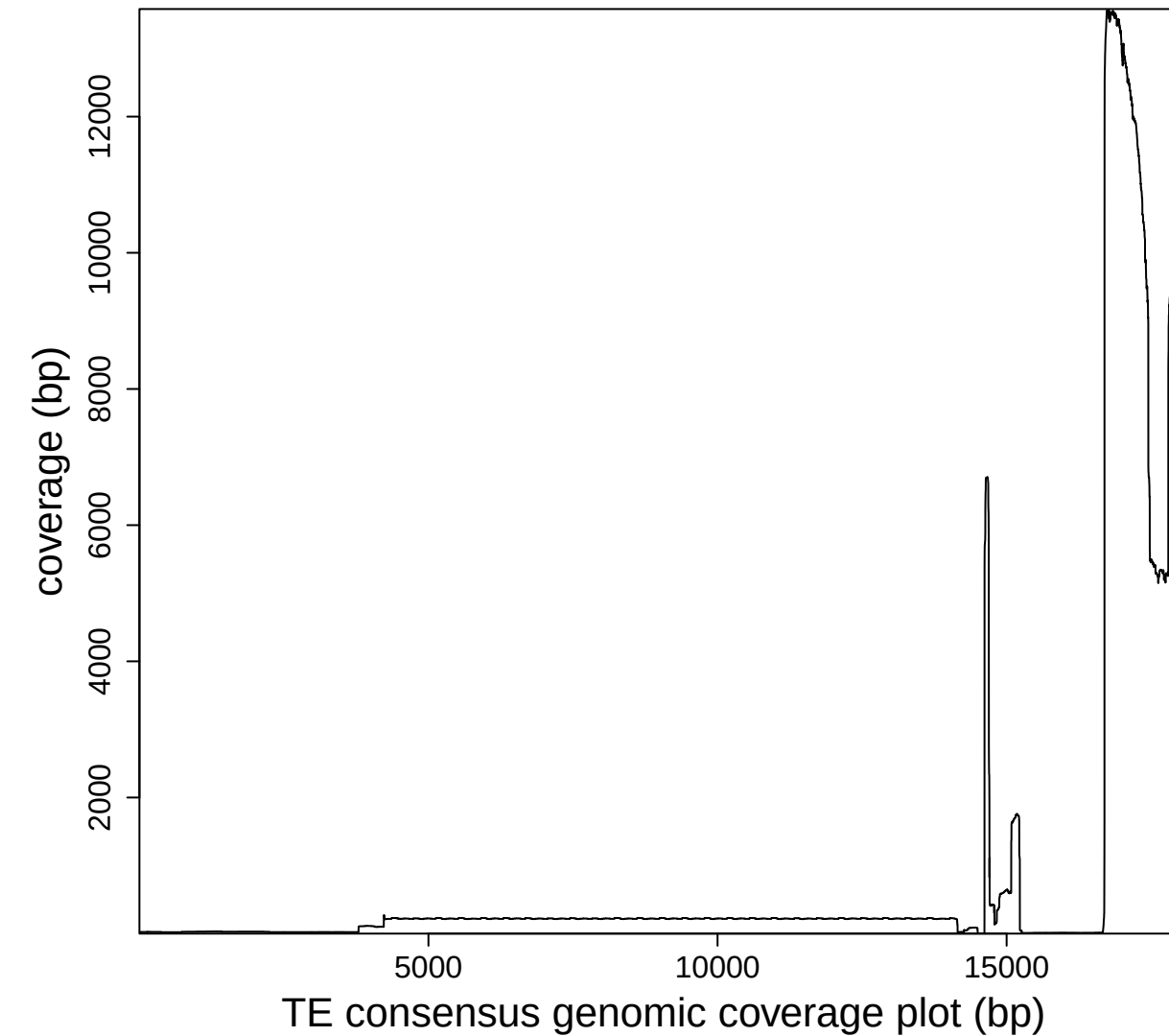
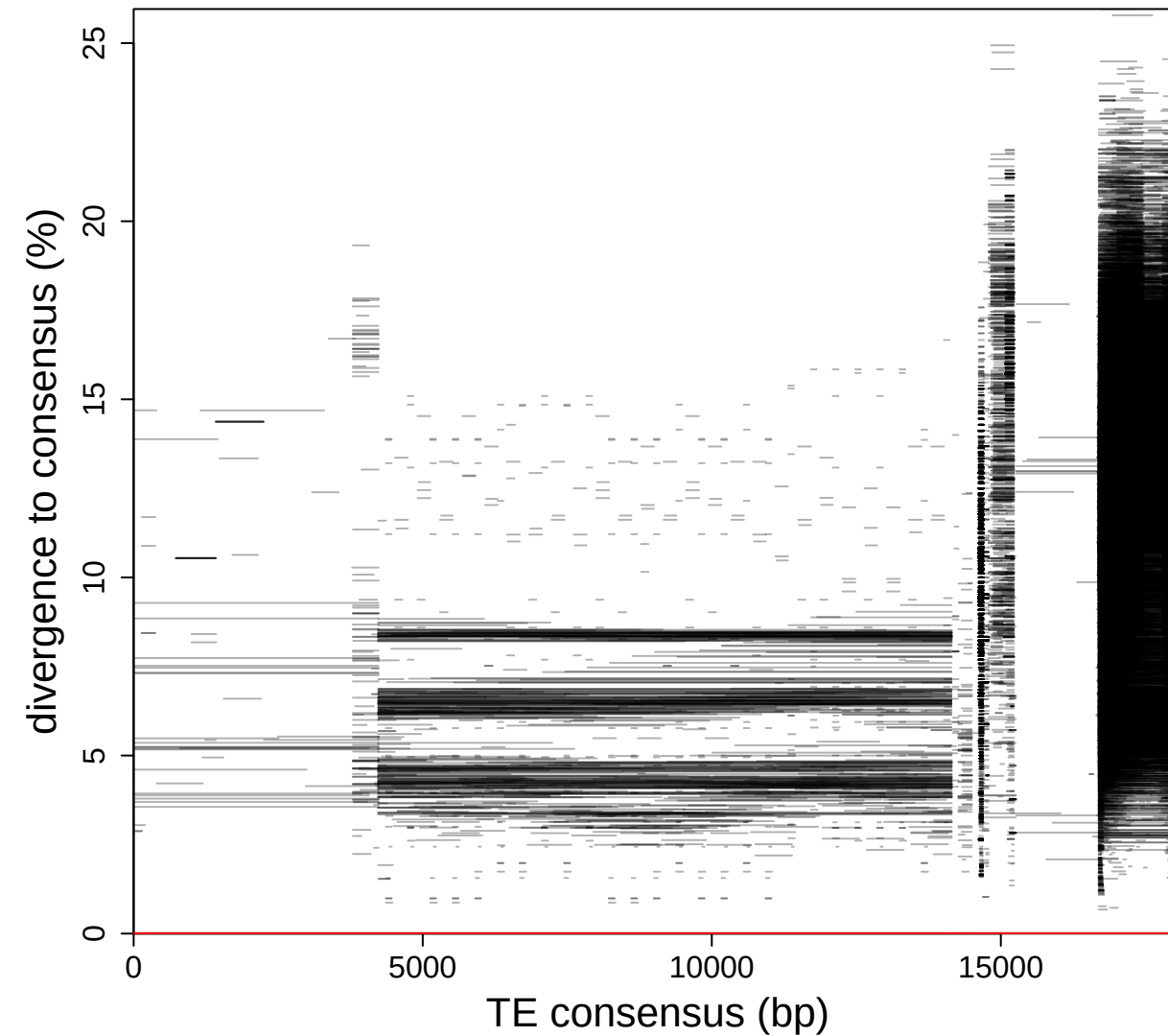


MSA length = 18134



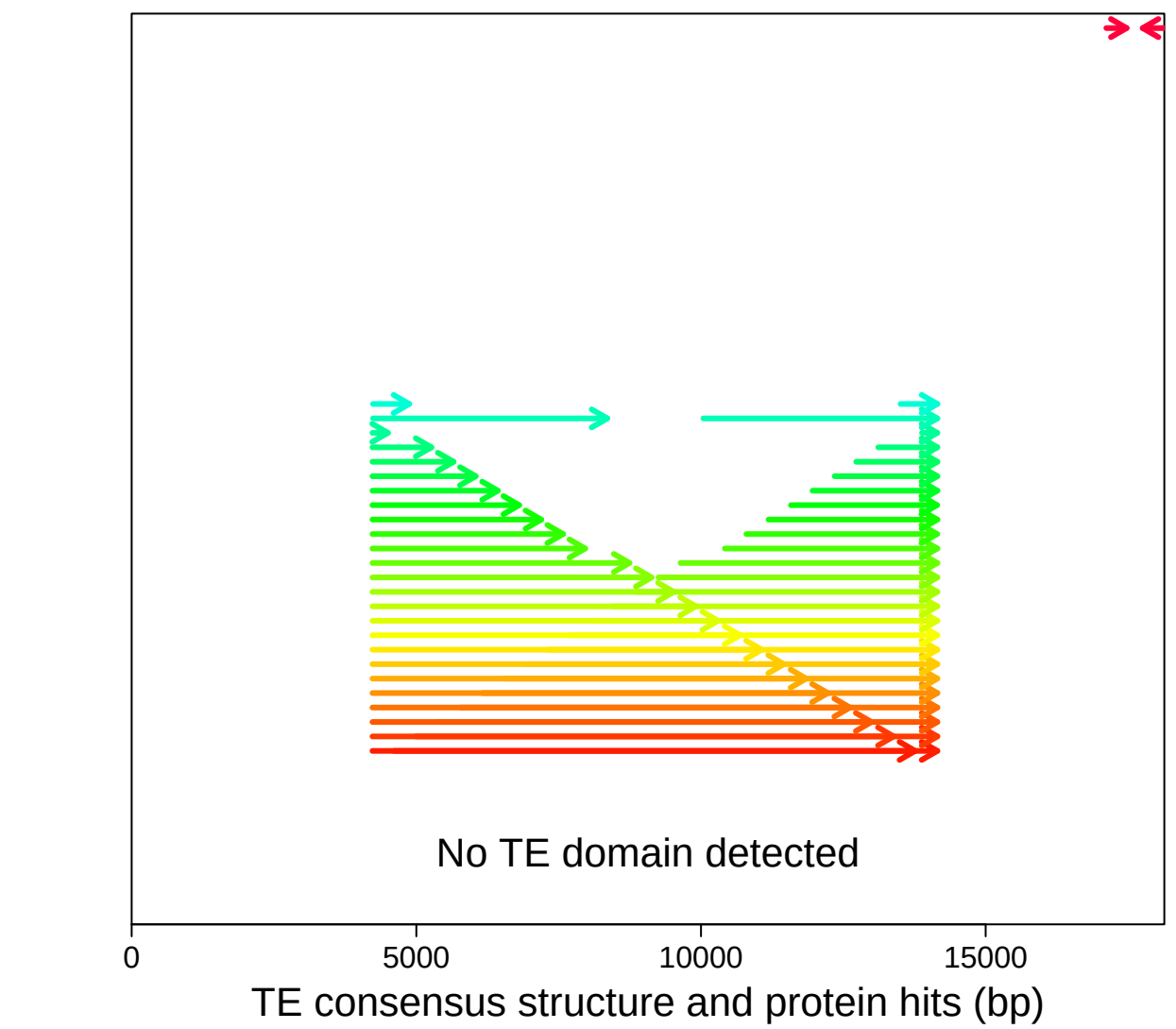
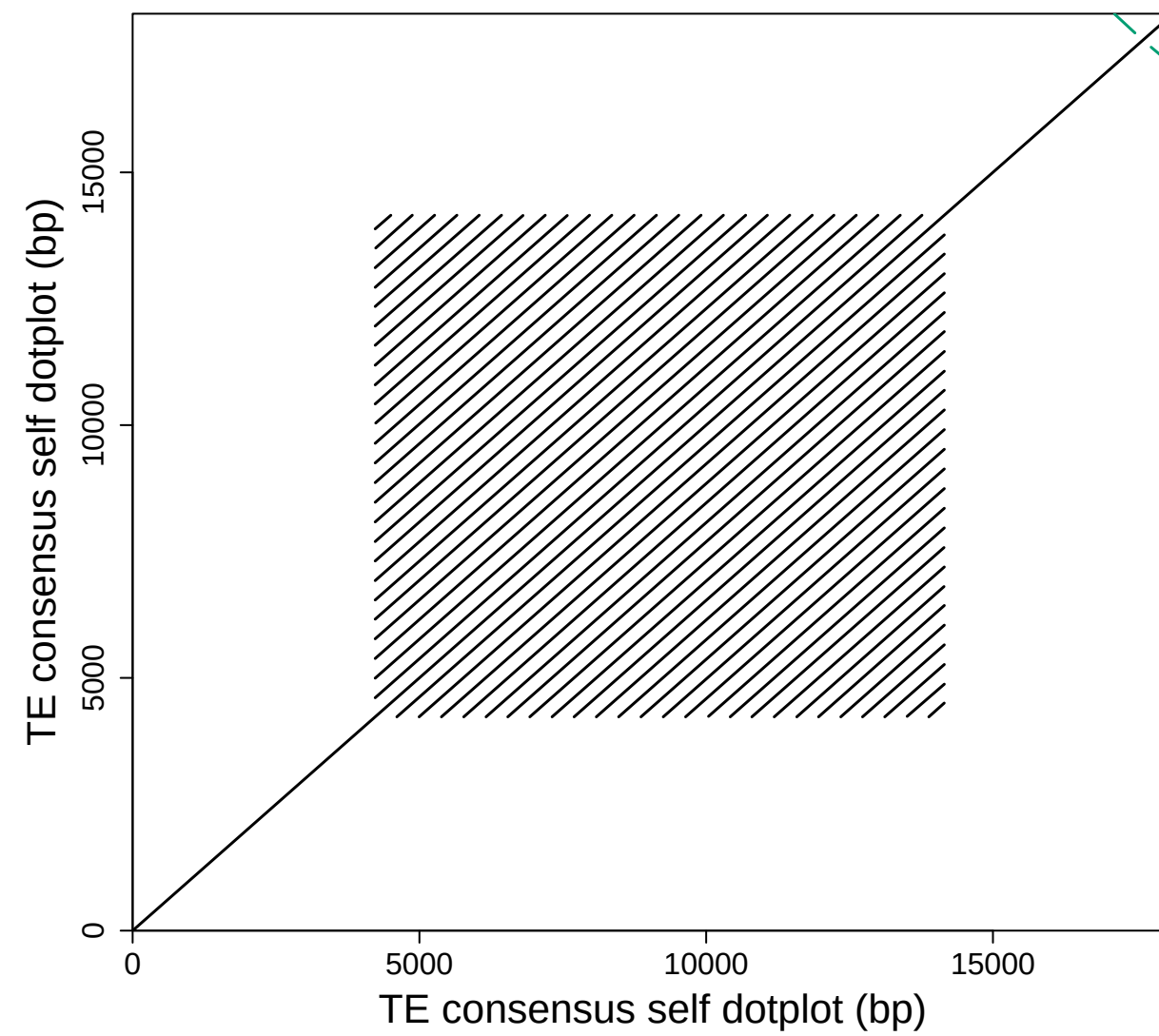
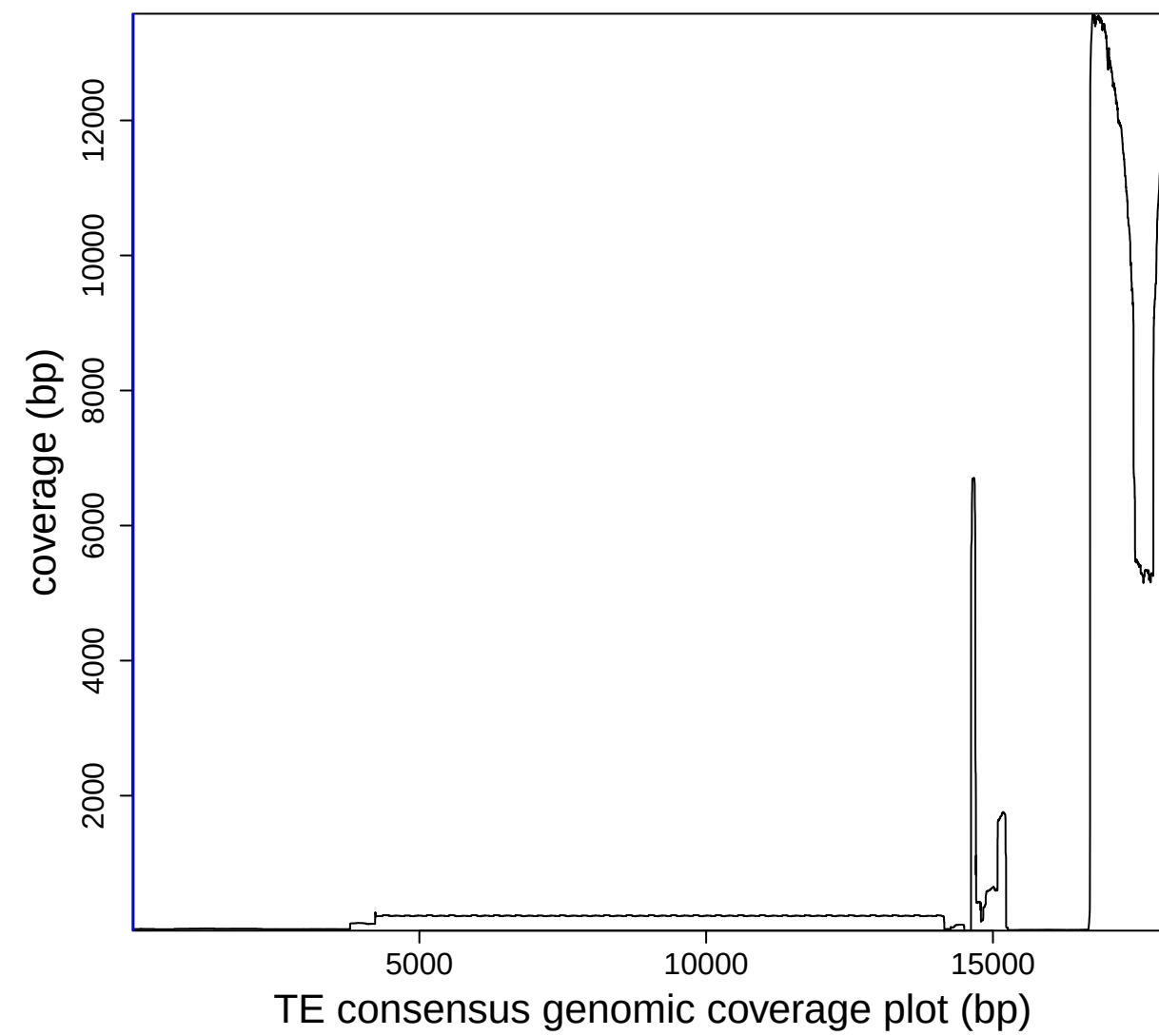
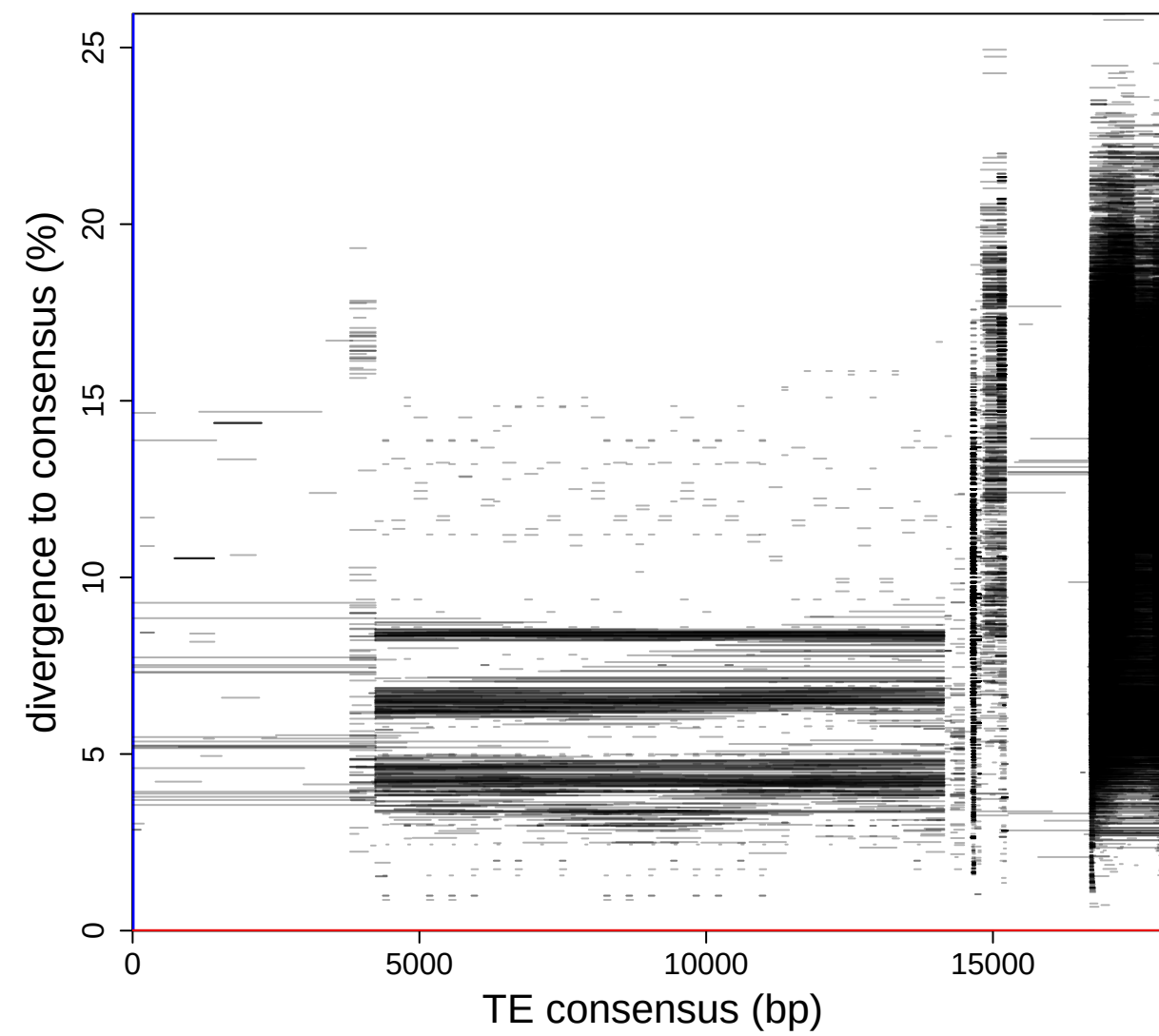
_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcln.fa al
size: 18134bp; fragments: 38301; full length: 1 (>=16320.6bp)

After TEtrimmer 18134 bp

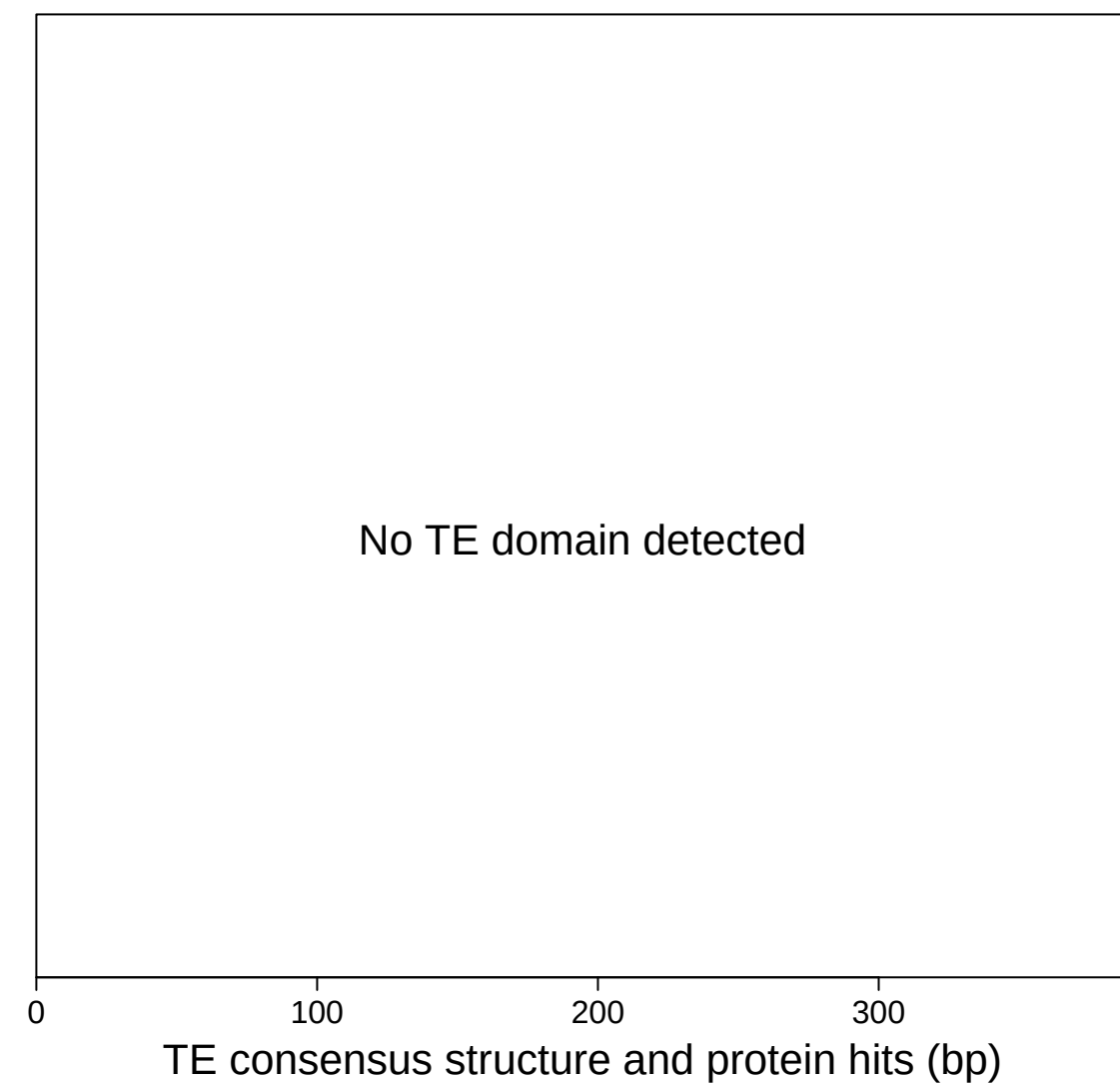
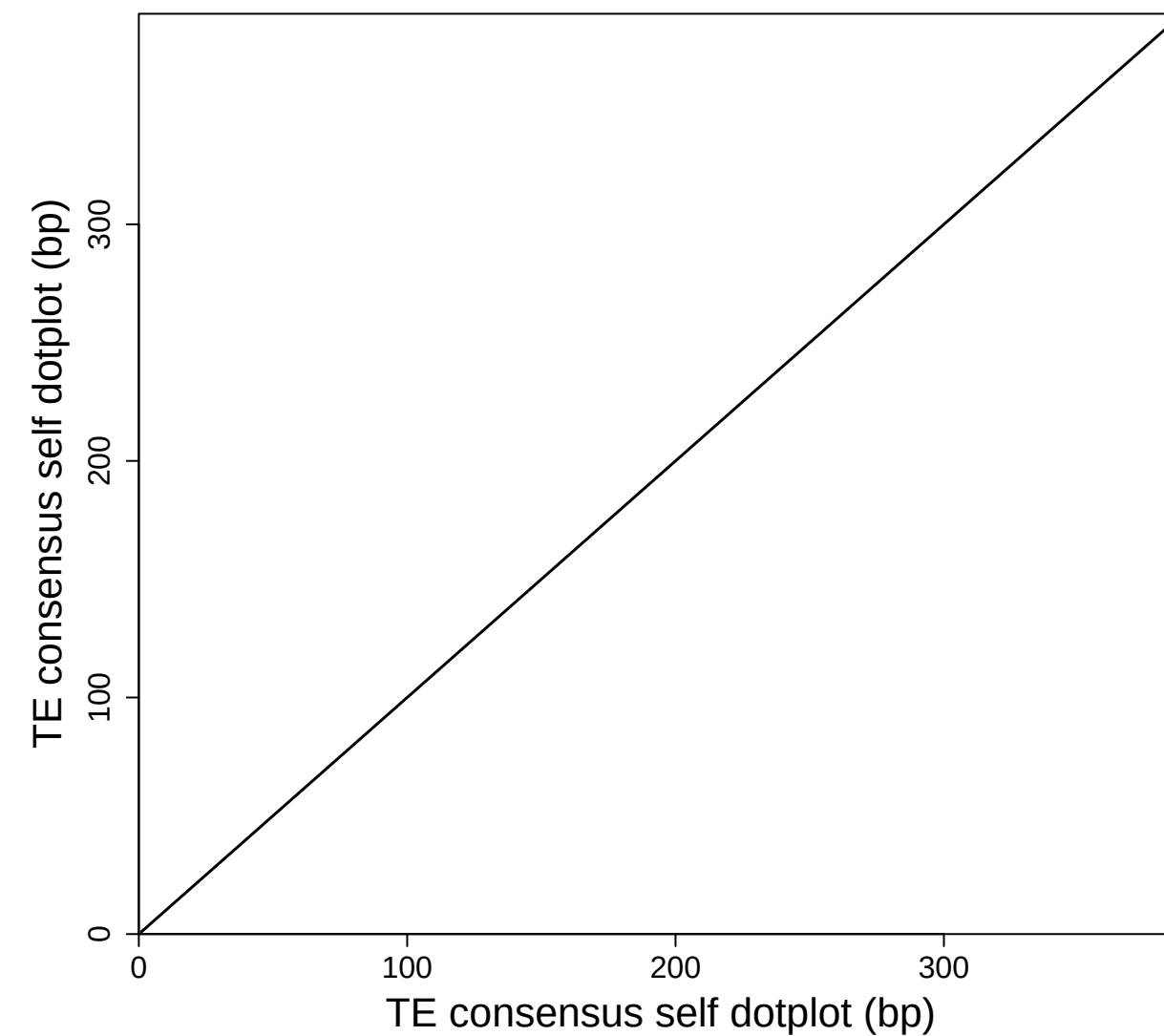
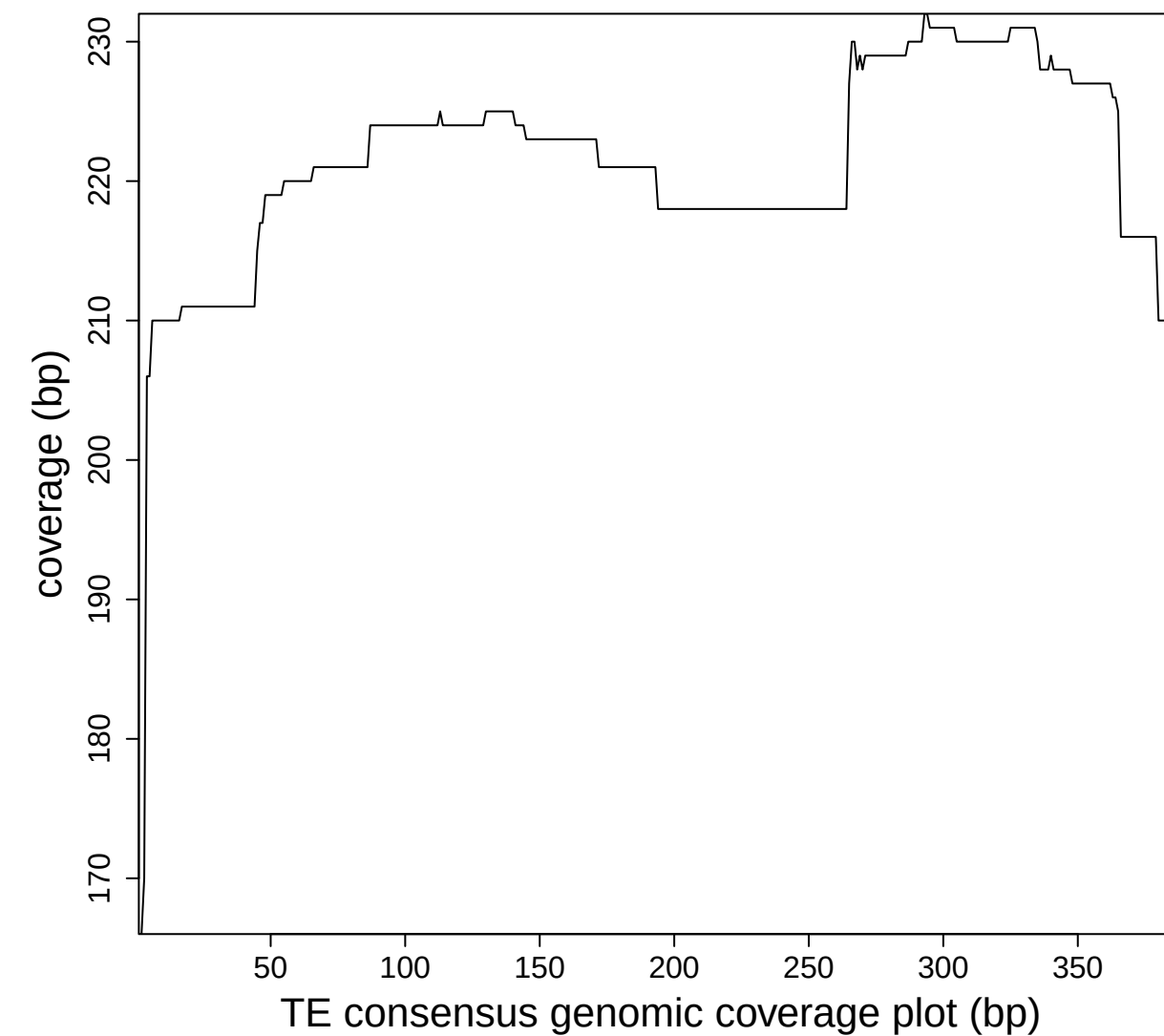
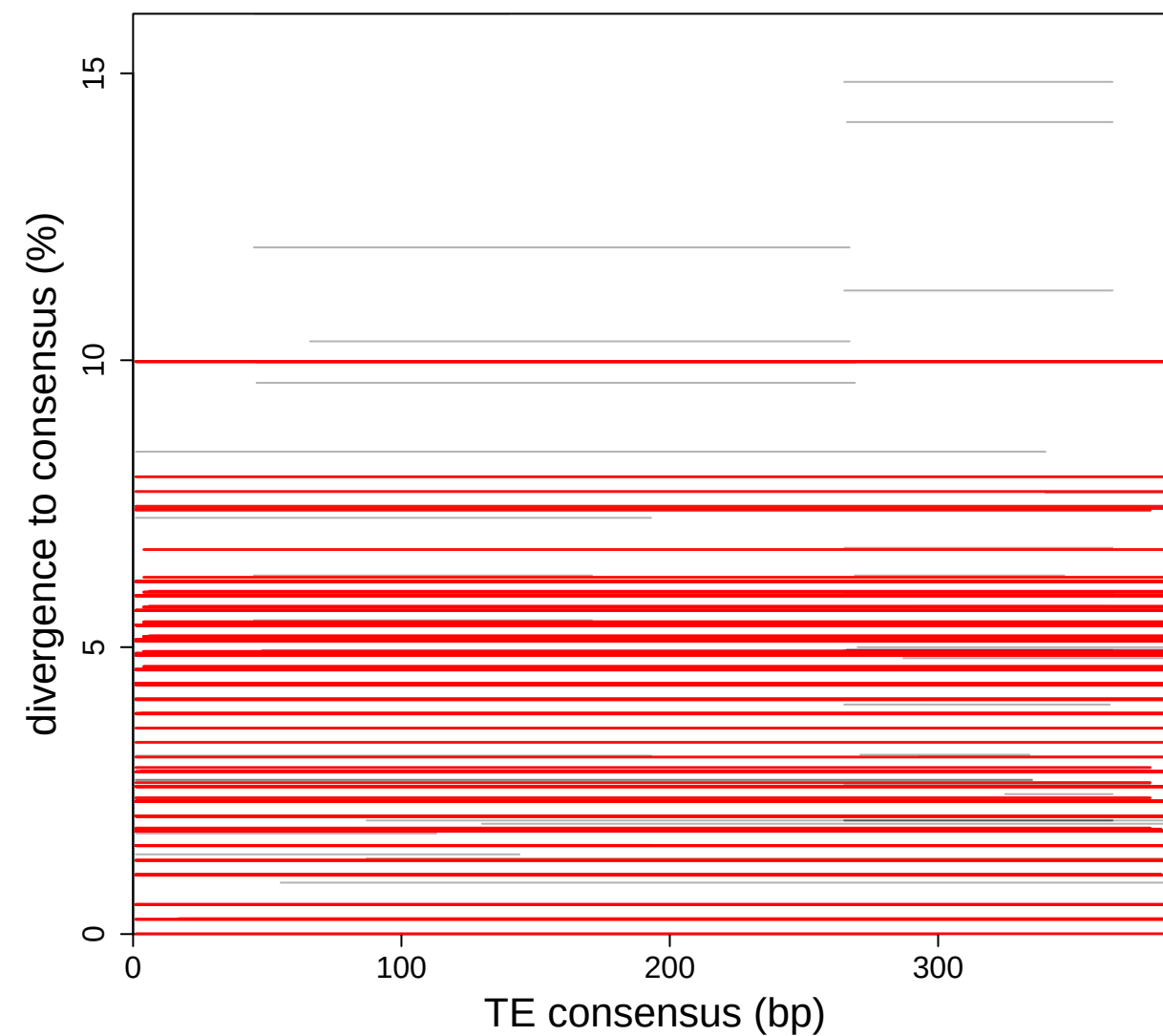


bed_g_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bclr
size: 18138bp; fragments: 38301; full length: 1 (>=16324.2bp)

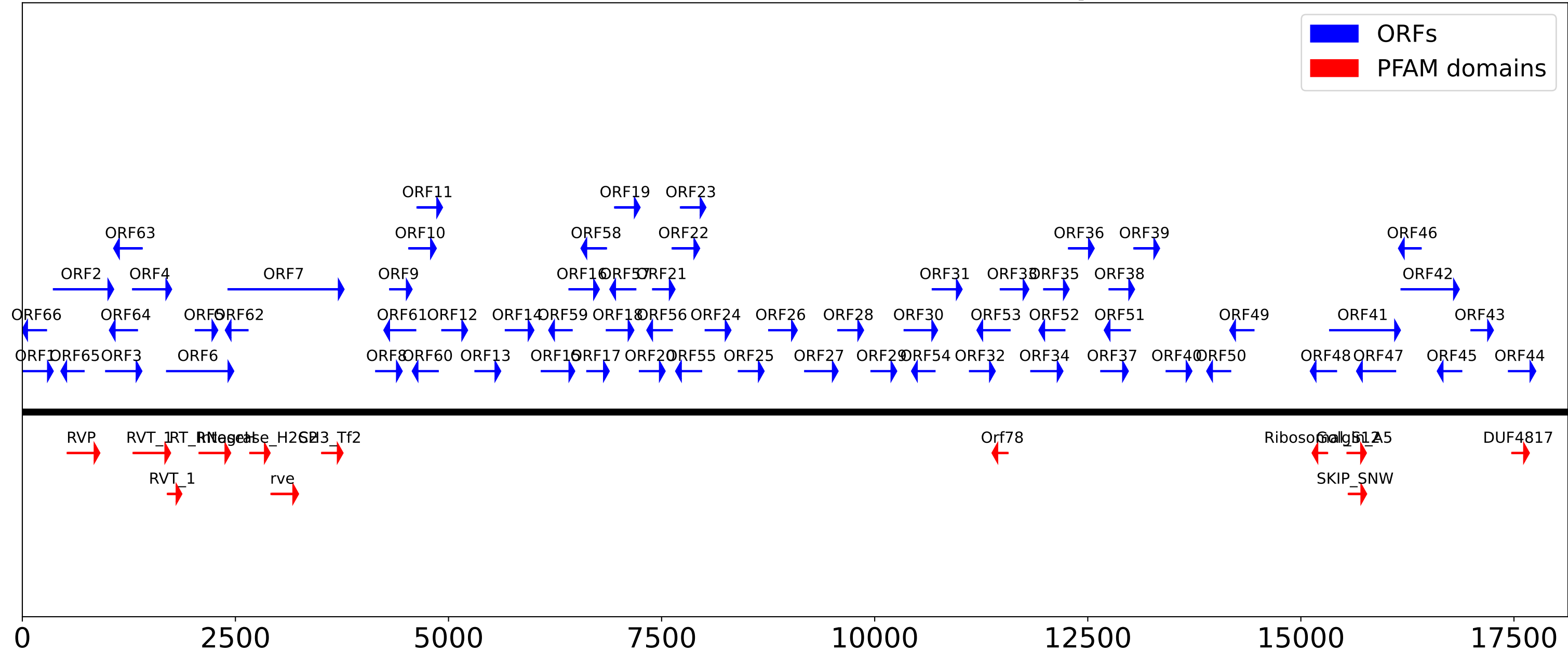
After TEtrimmer Extended plot Blue lines are boundaries



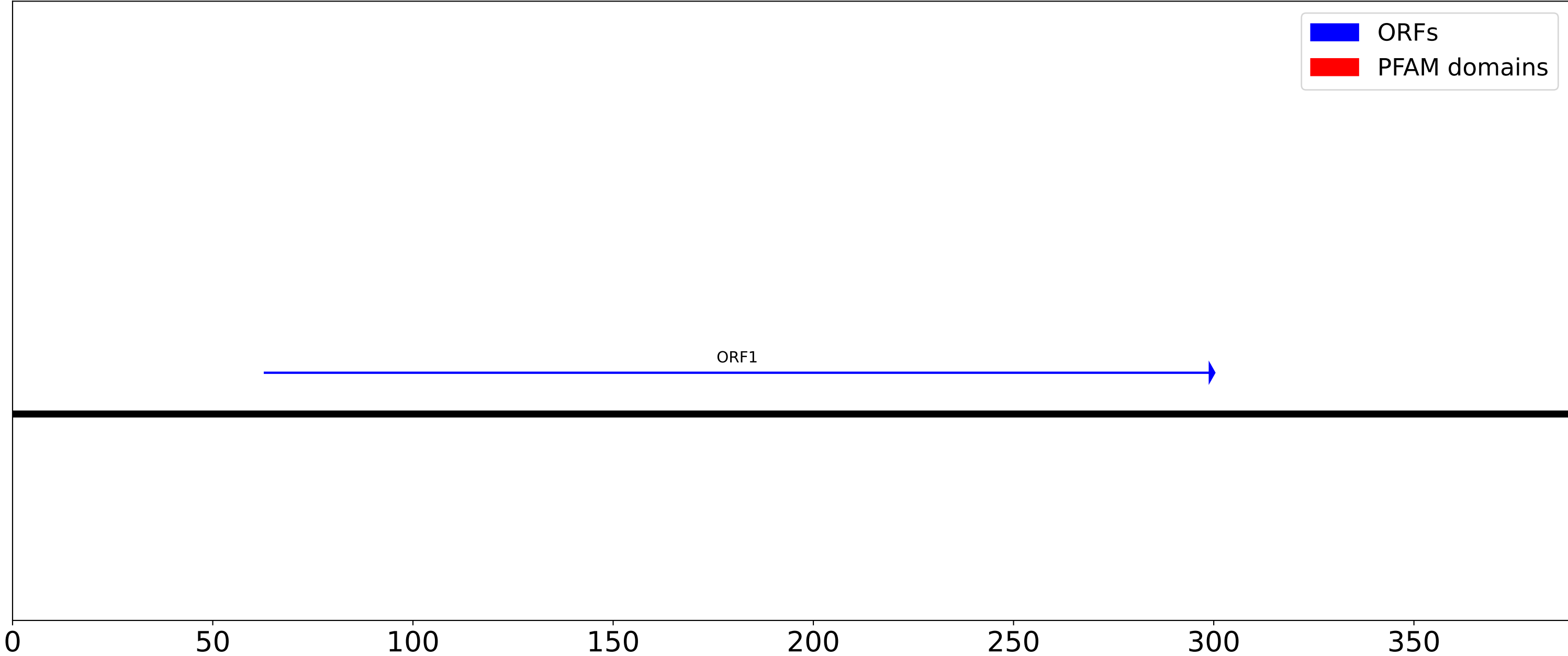
TE: ltr_1_family_105
size: 389bp; fragments: 246; full length: 201 (≥ 350.1 bp)



After TETrimmer ORF and PFAM domain plot



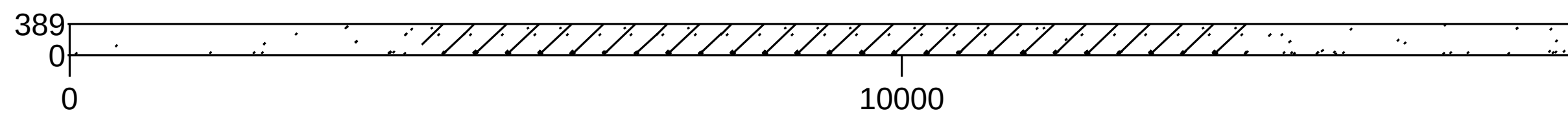
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TETrimmer (bp)



TE consensus after TETrimmer (bp)