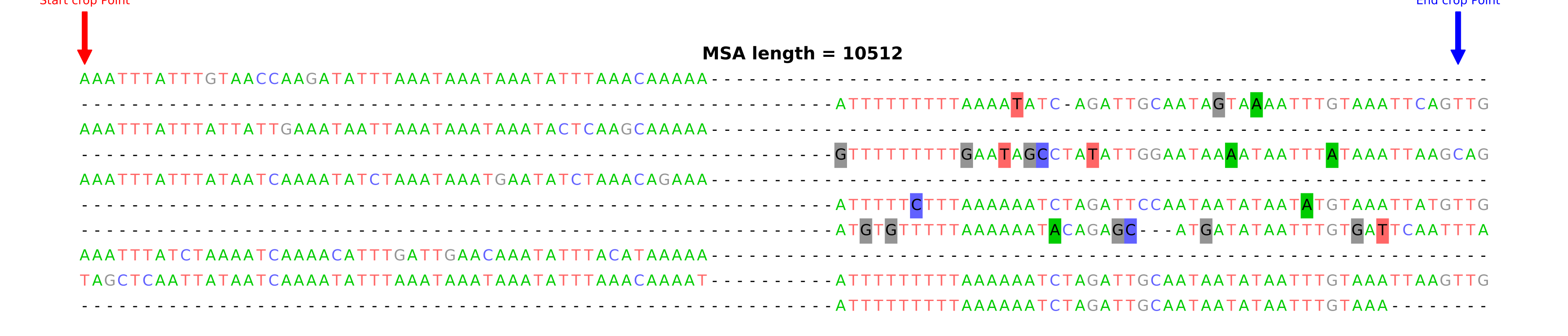
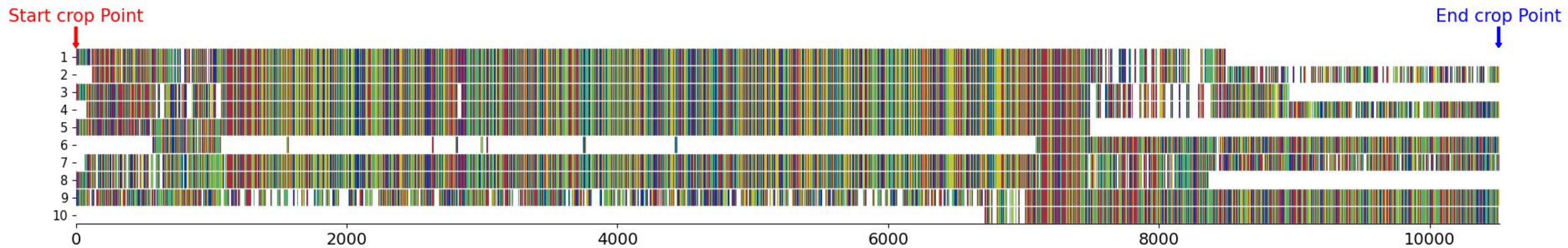
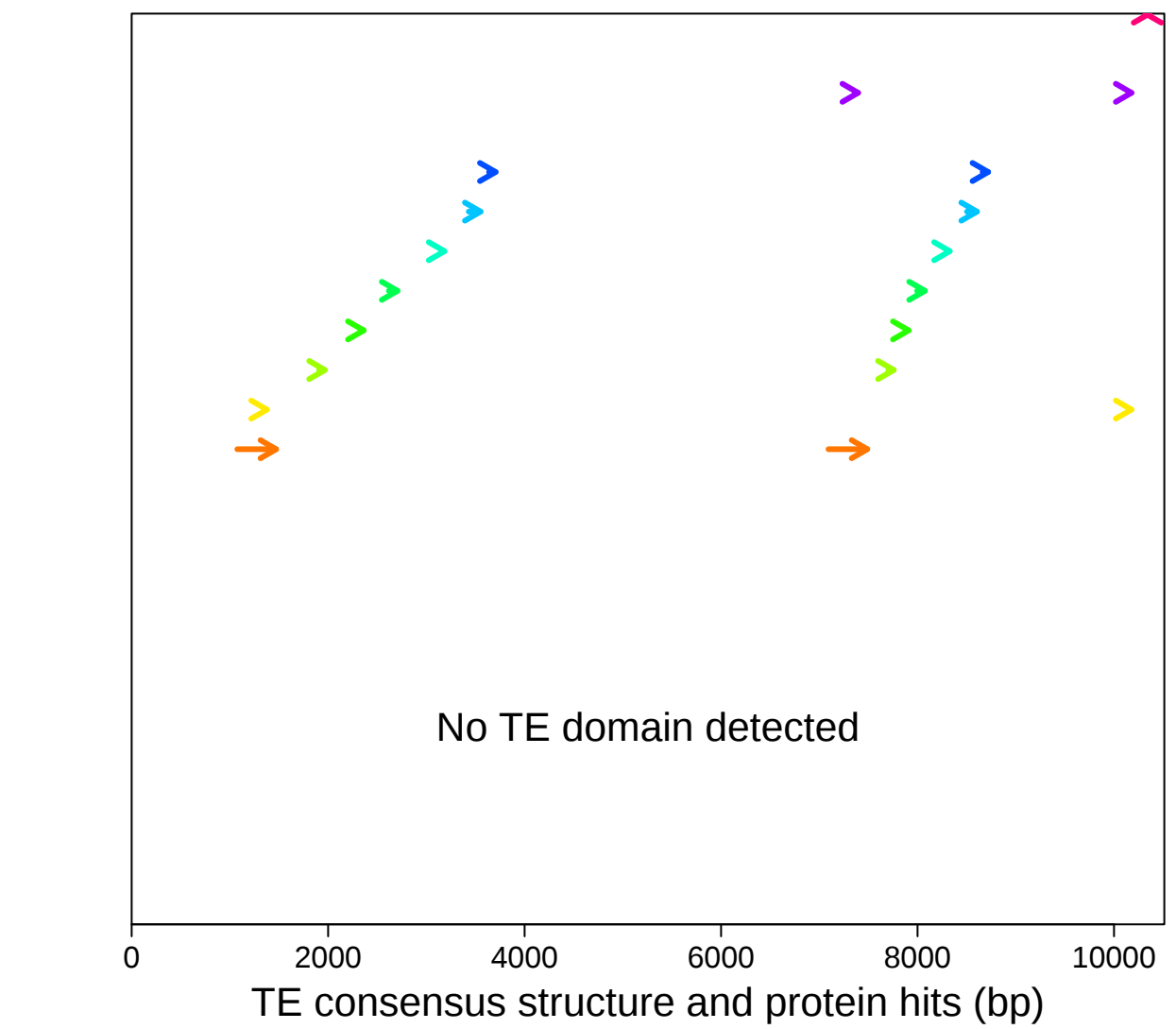
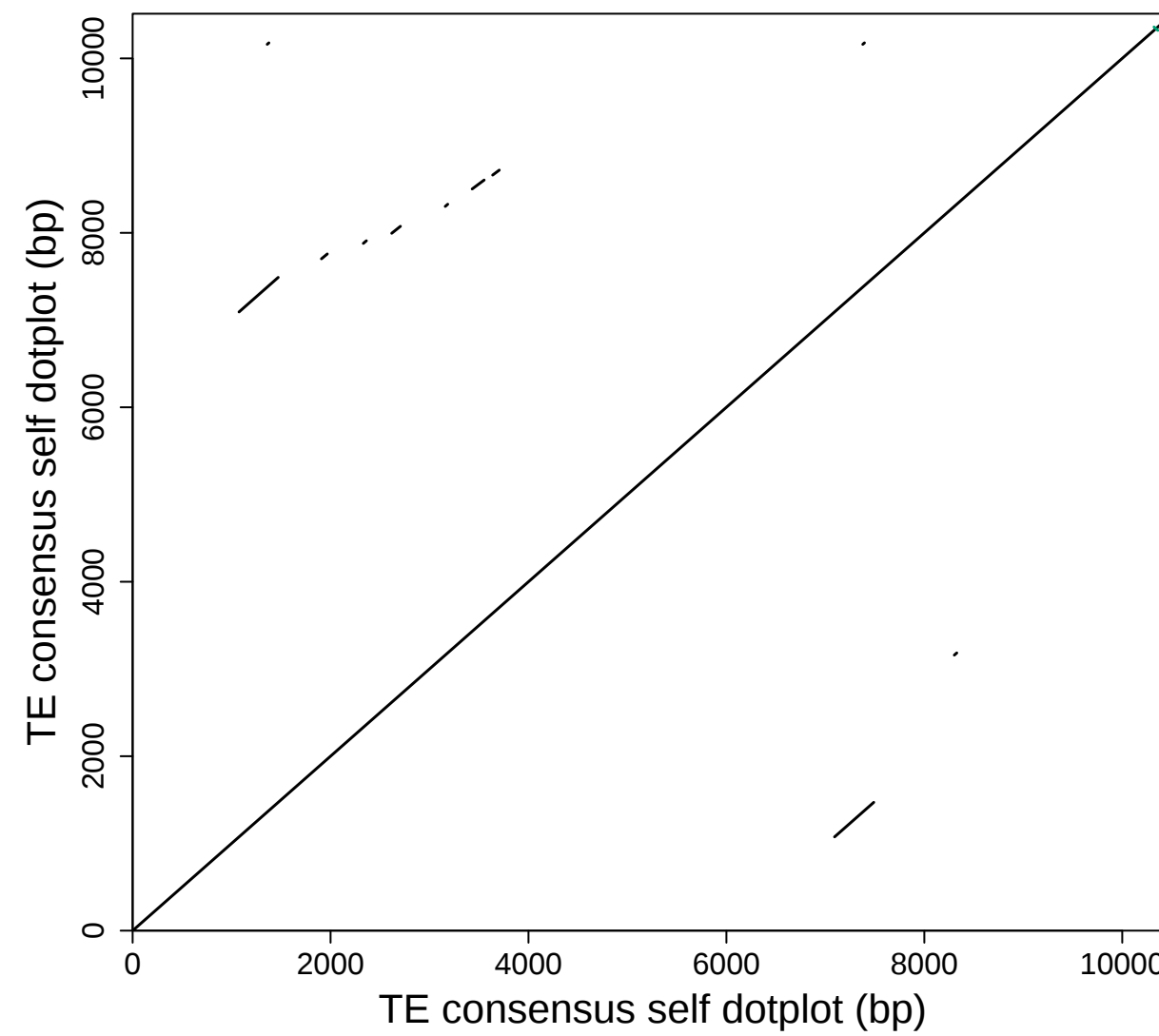
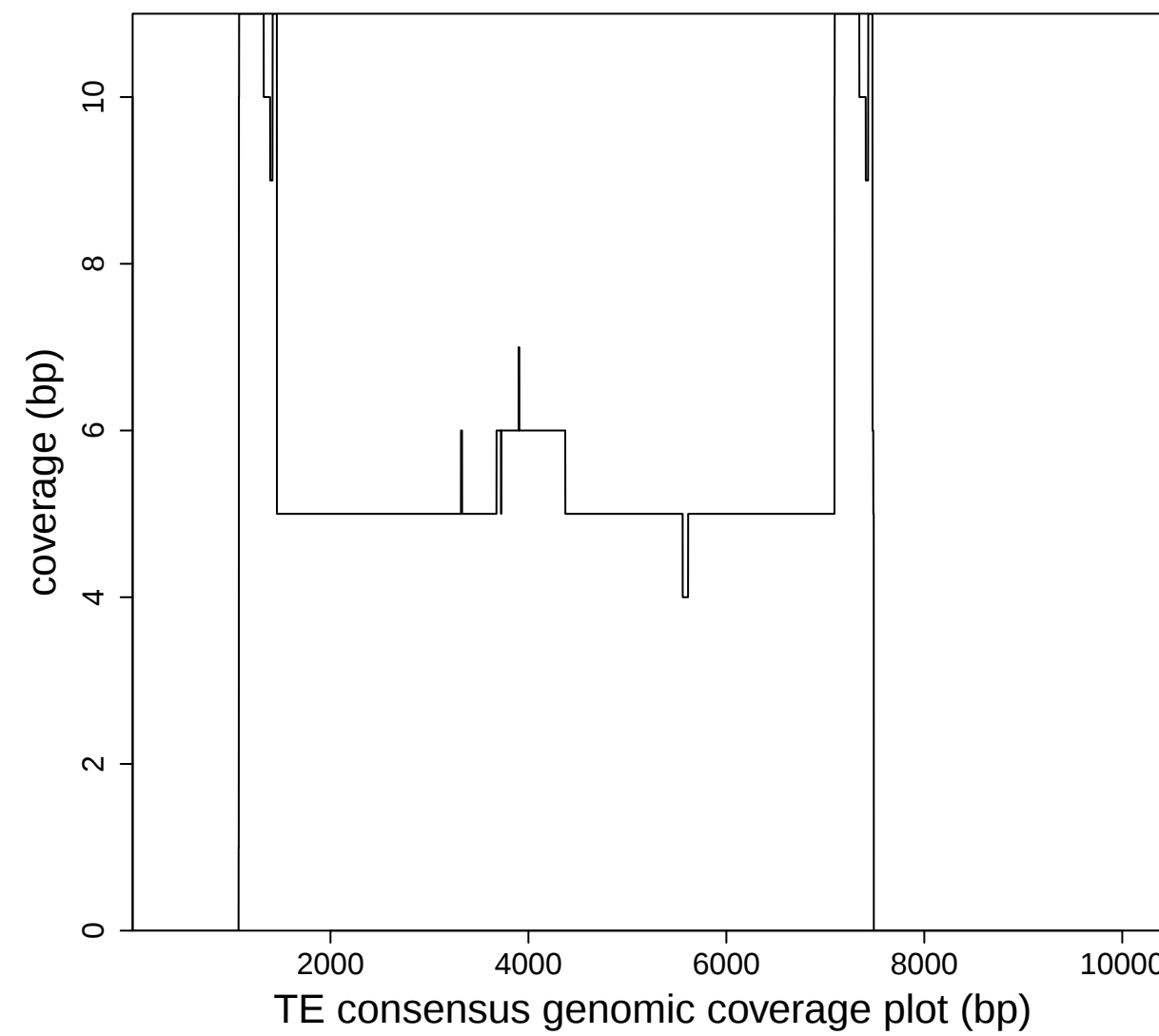
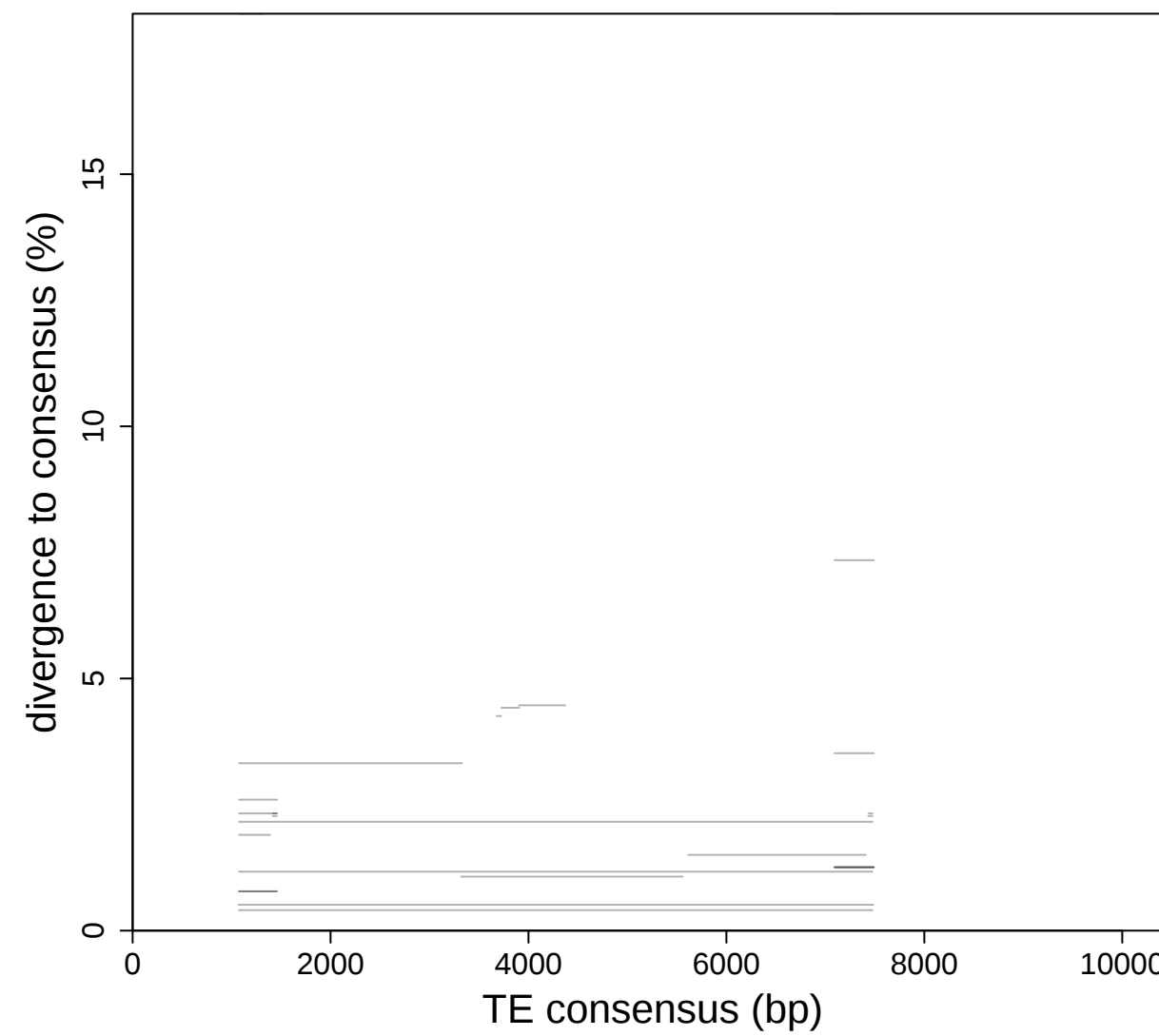






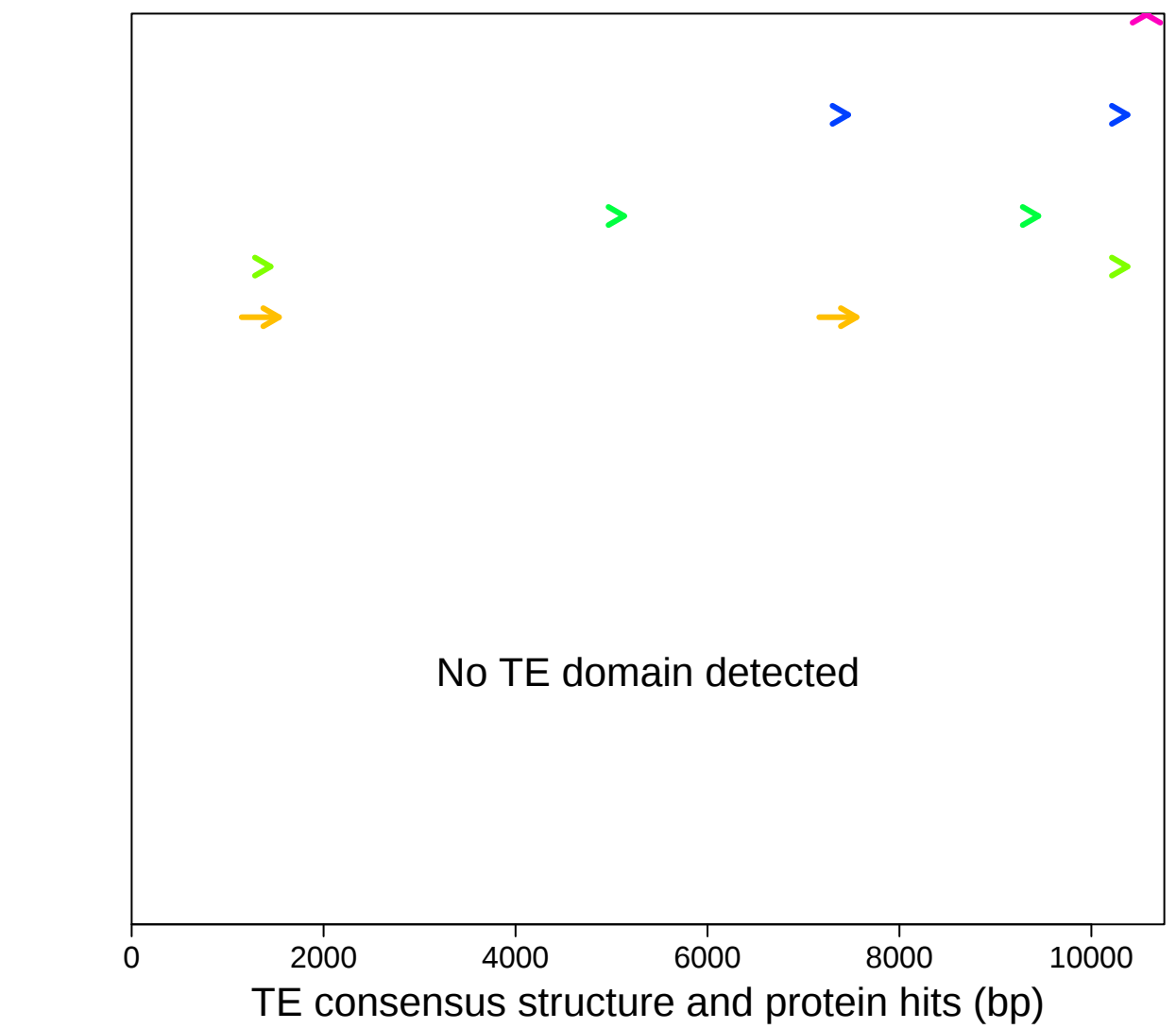
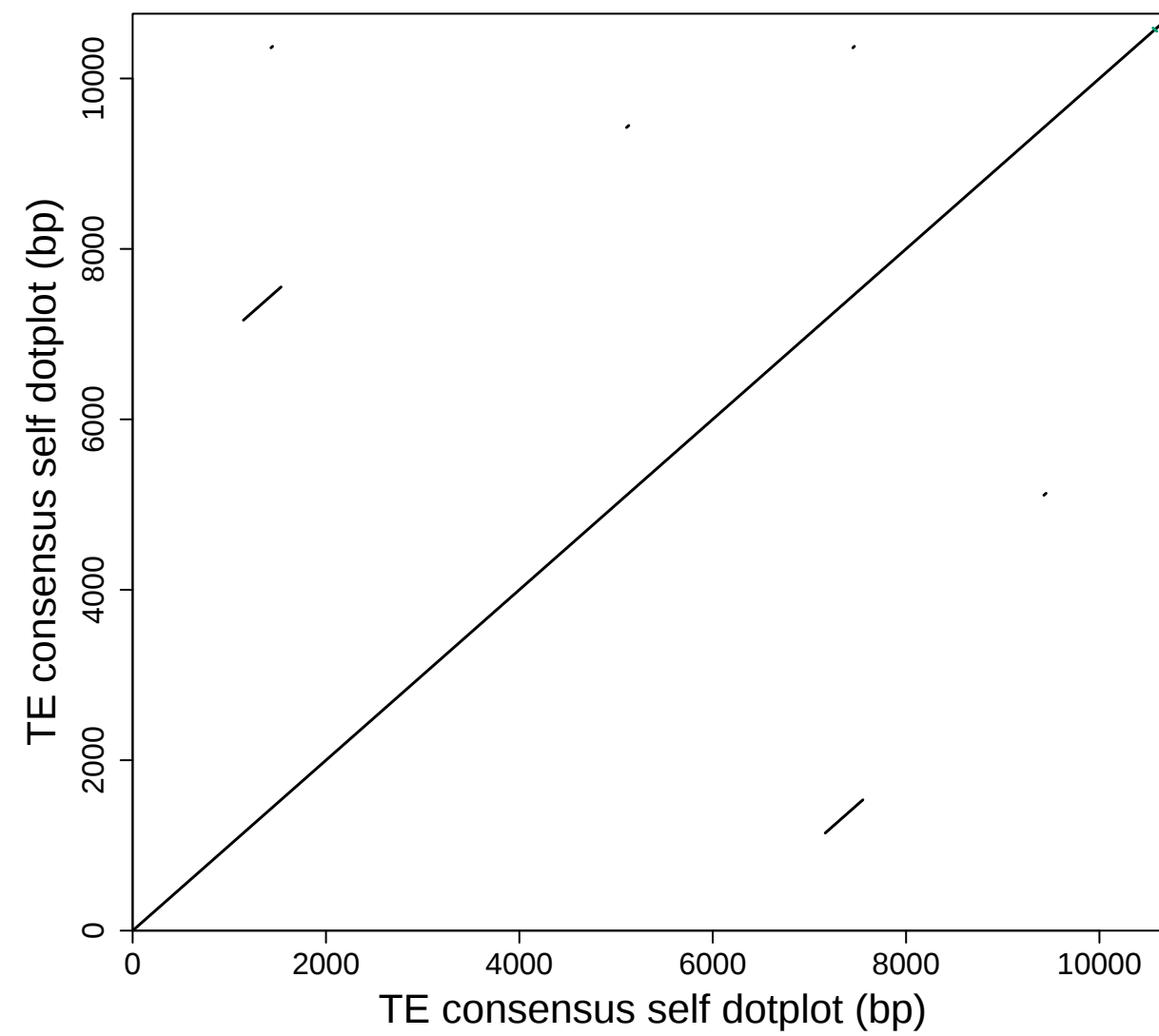
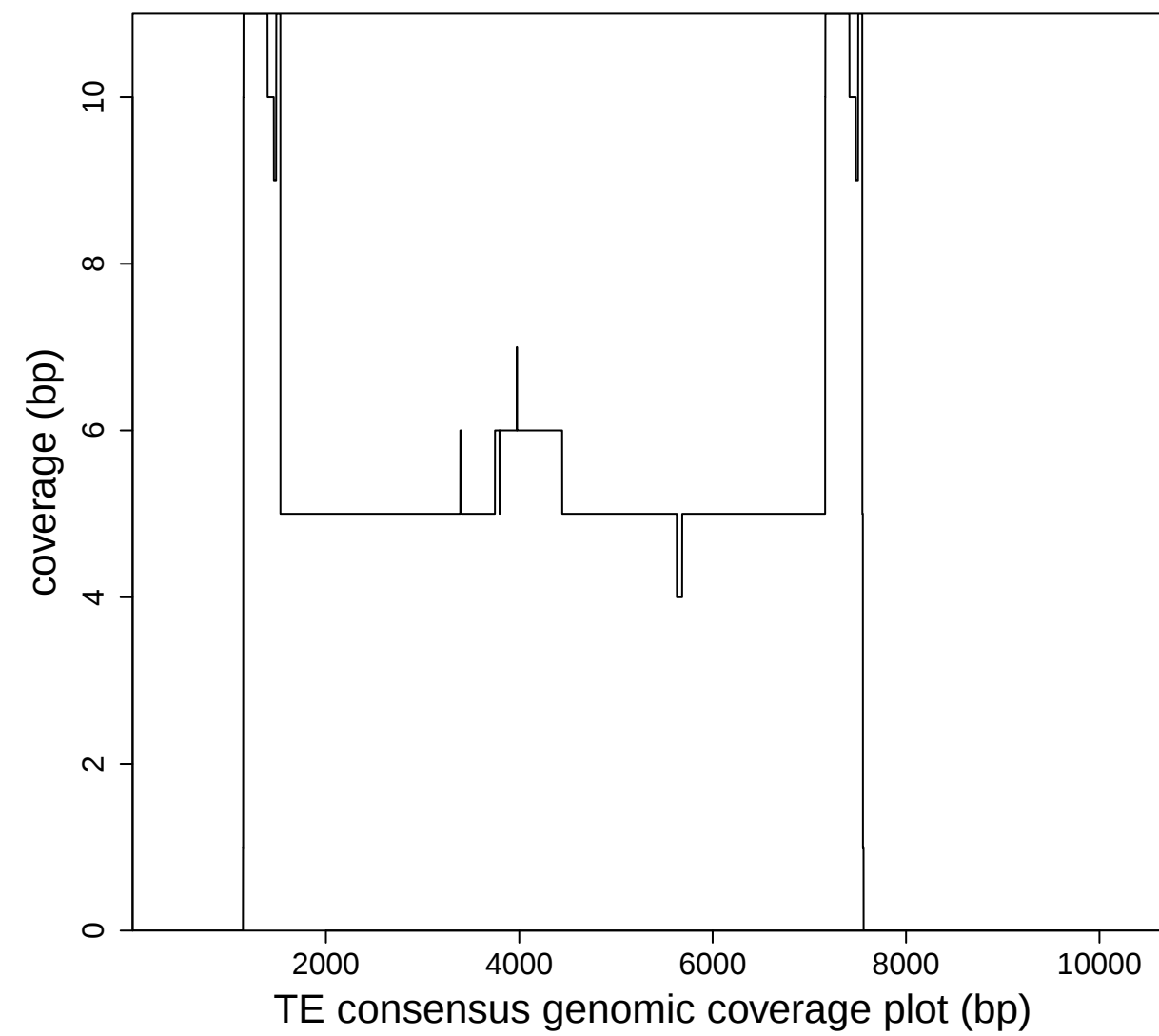
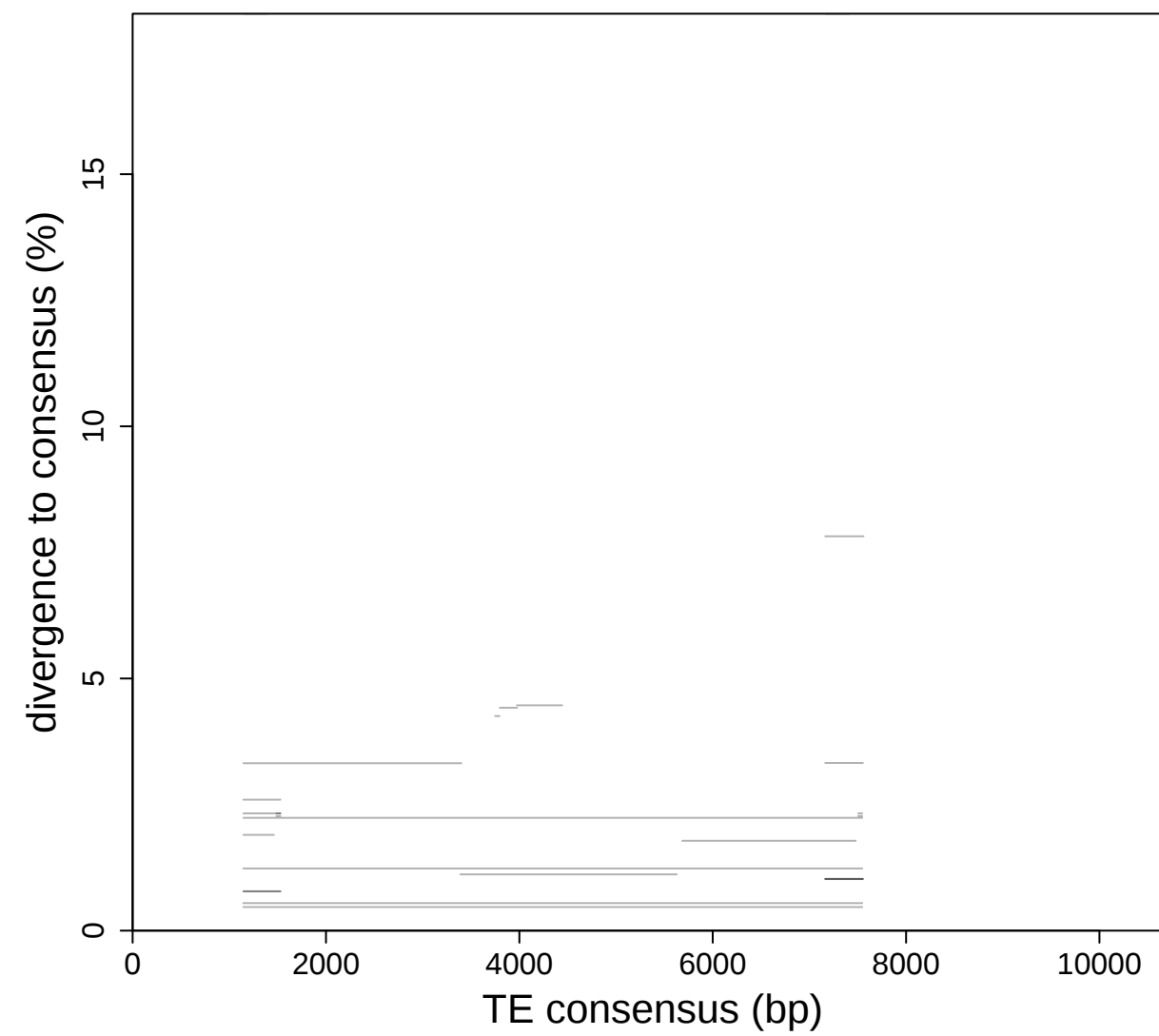
b.bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.fa
size: 10512bp; fragments: 26; full length: 0 (>=9460.8bp)

After TEtrimmer 10512 bp

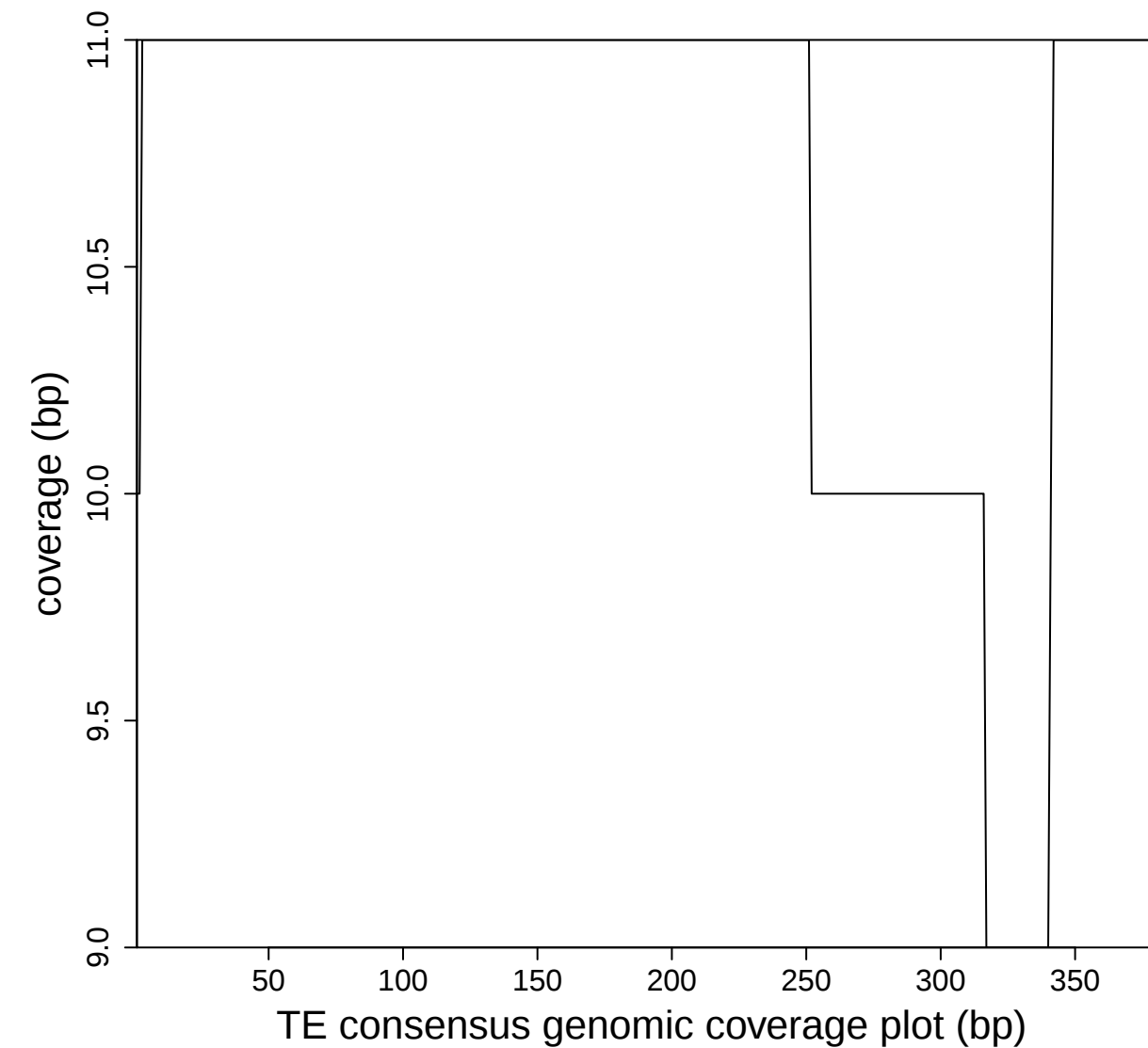
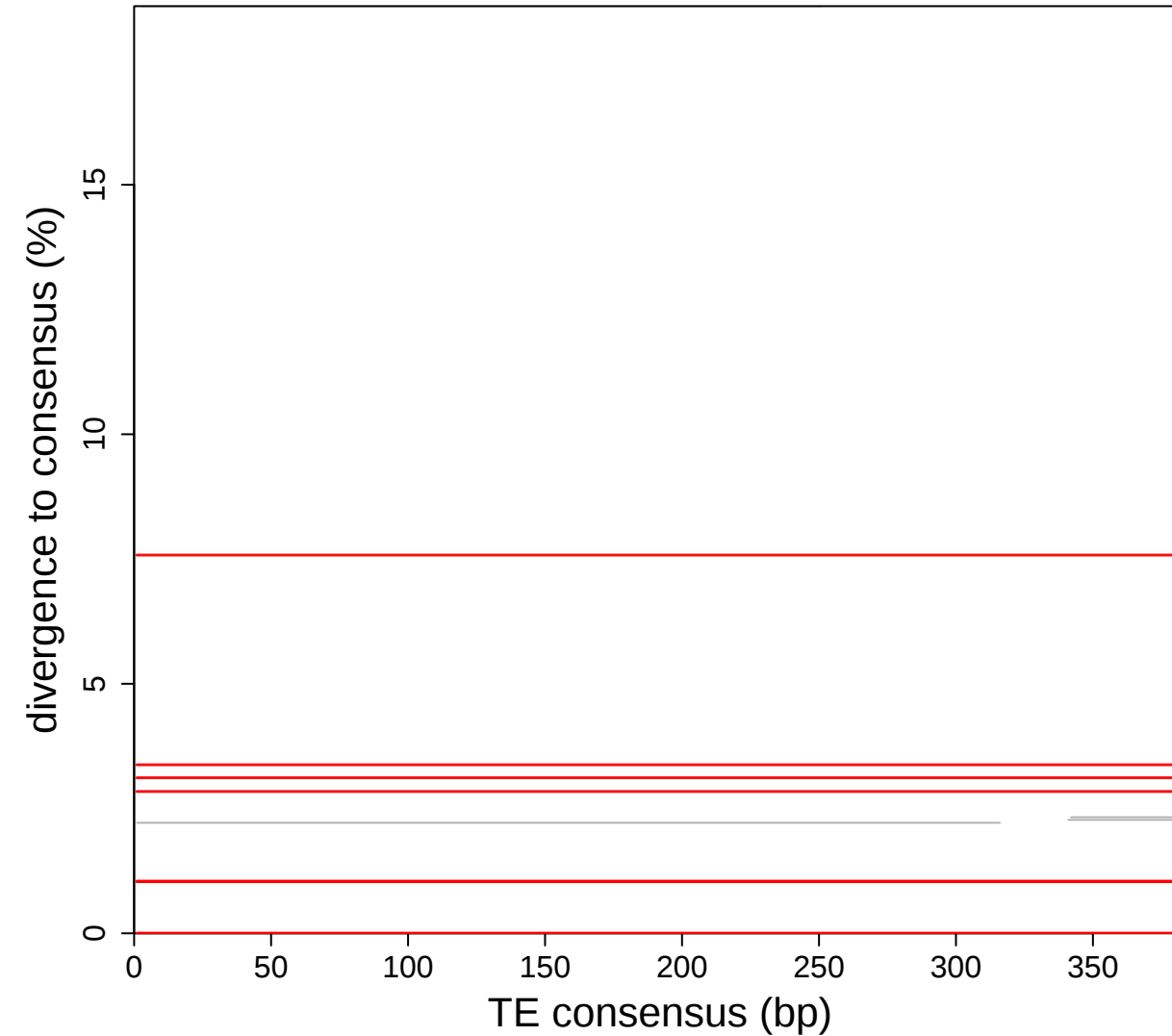


.fasta.b.bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa
size: 10760bp; fragments: 26; full length: 0 (>=9684bp)

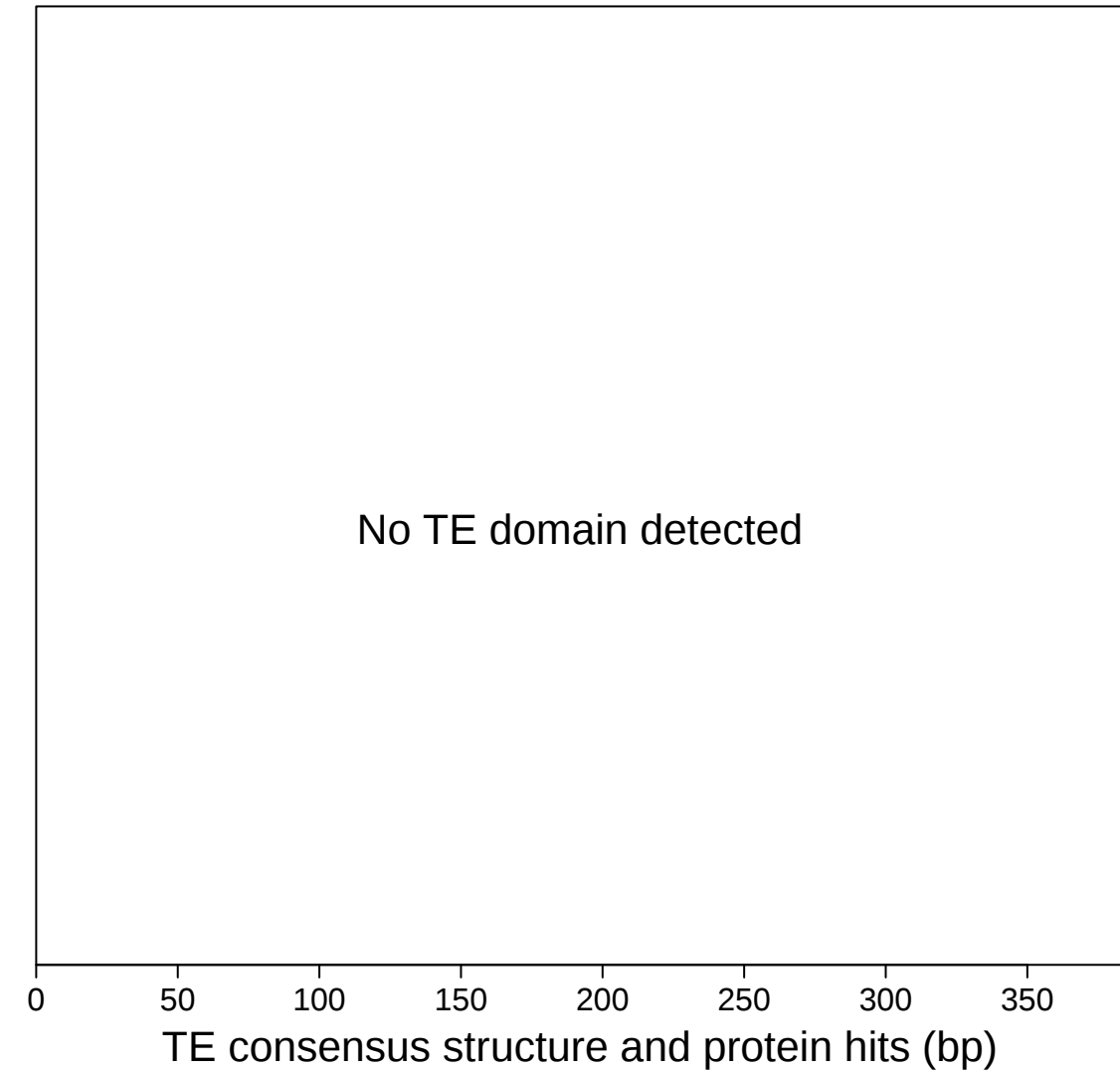
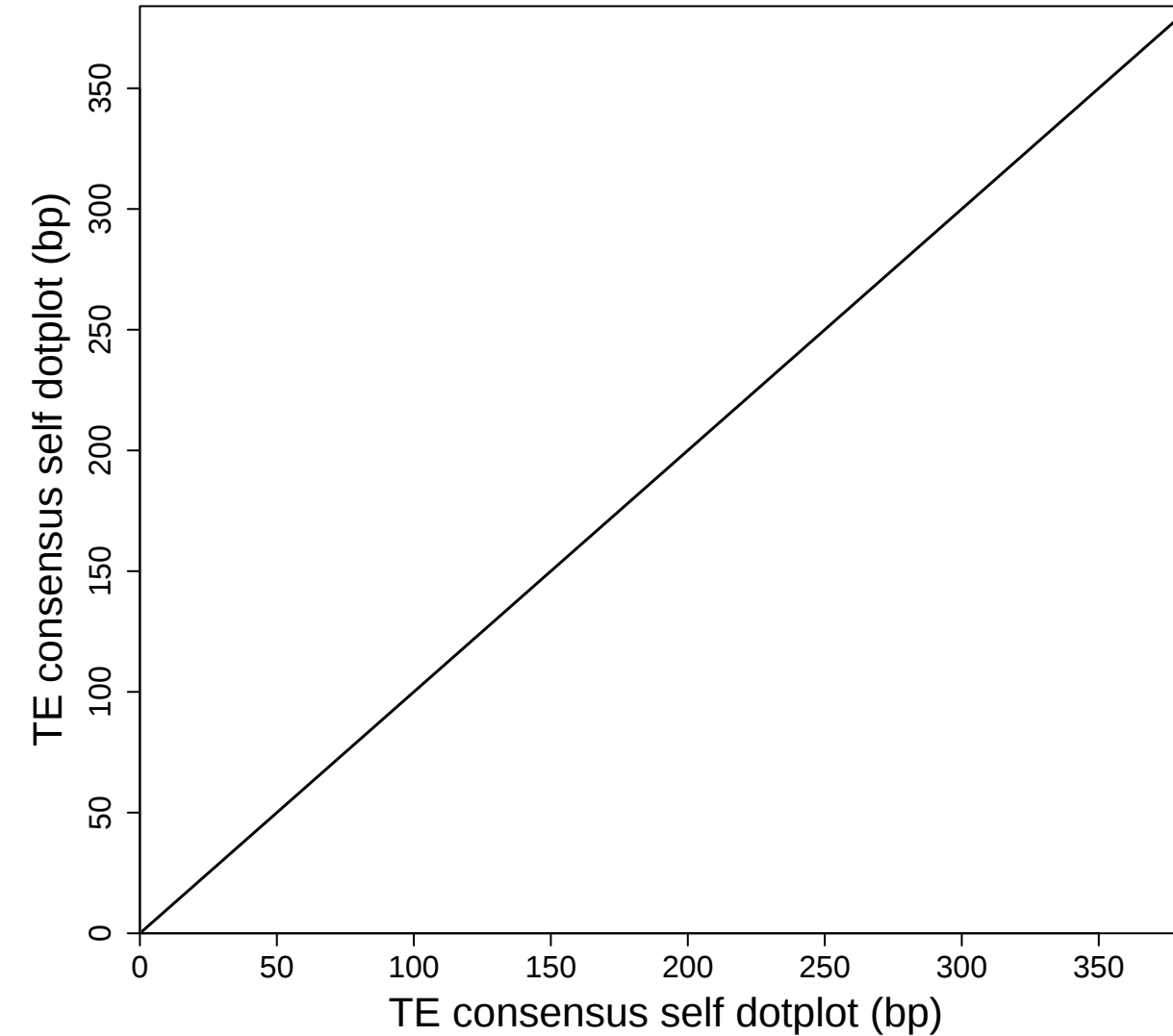
After TEtrimmer Extended plot Blue lines are boundaries



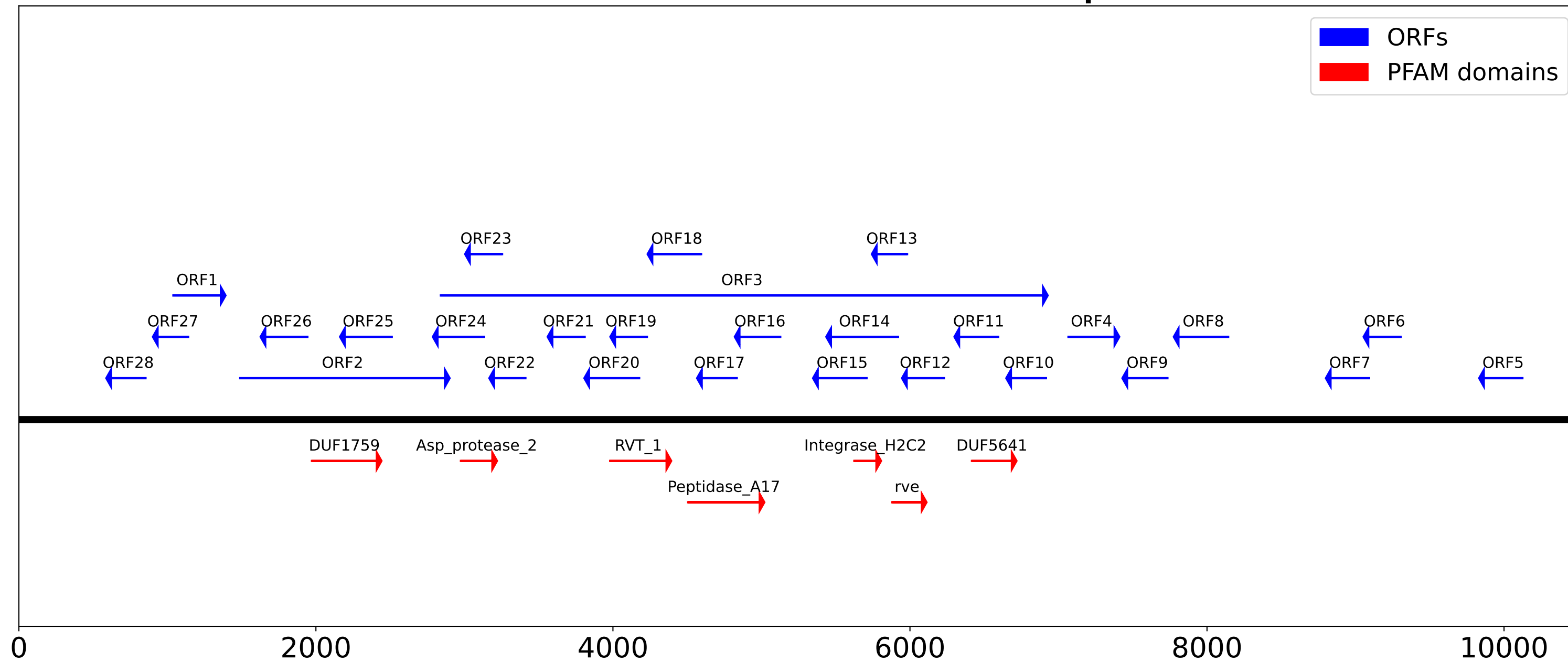
TE: ltr_1_family_136
size: 384bp; fragments: 13; full length: 9 (≥ 345.6 bp)



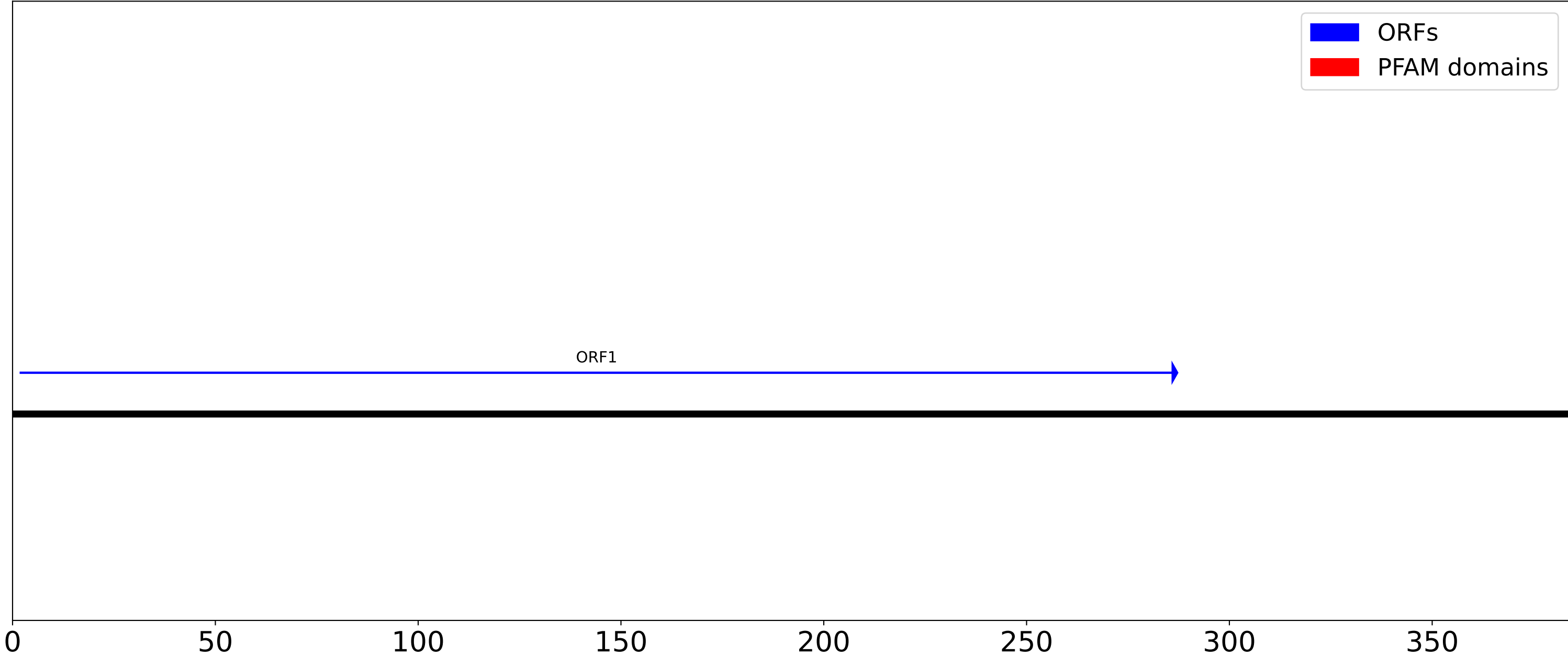
Before TEtrimmer 384 bp



After TETrimmer ORF and PFAM domain plot



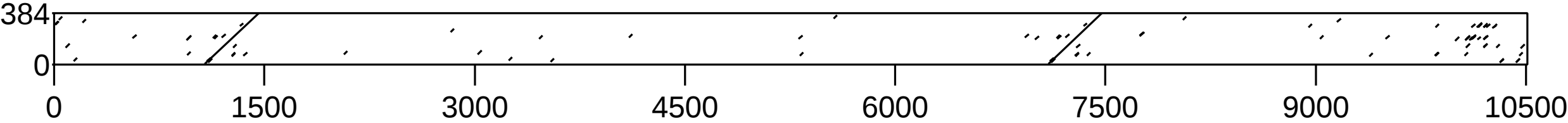
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

nsus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)