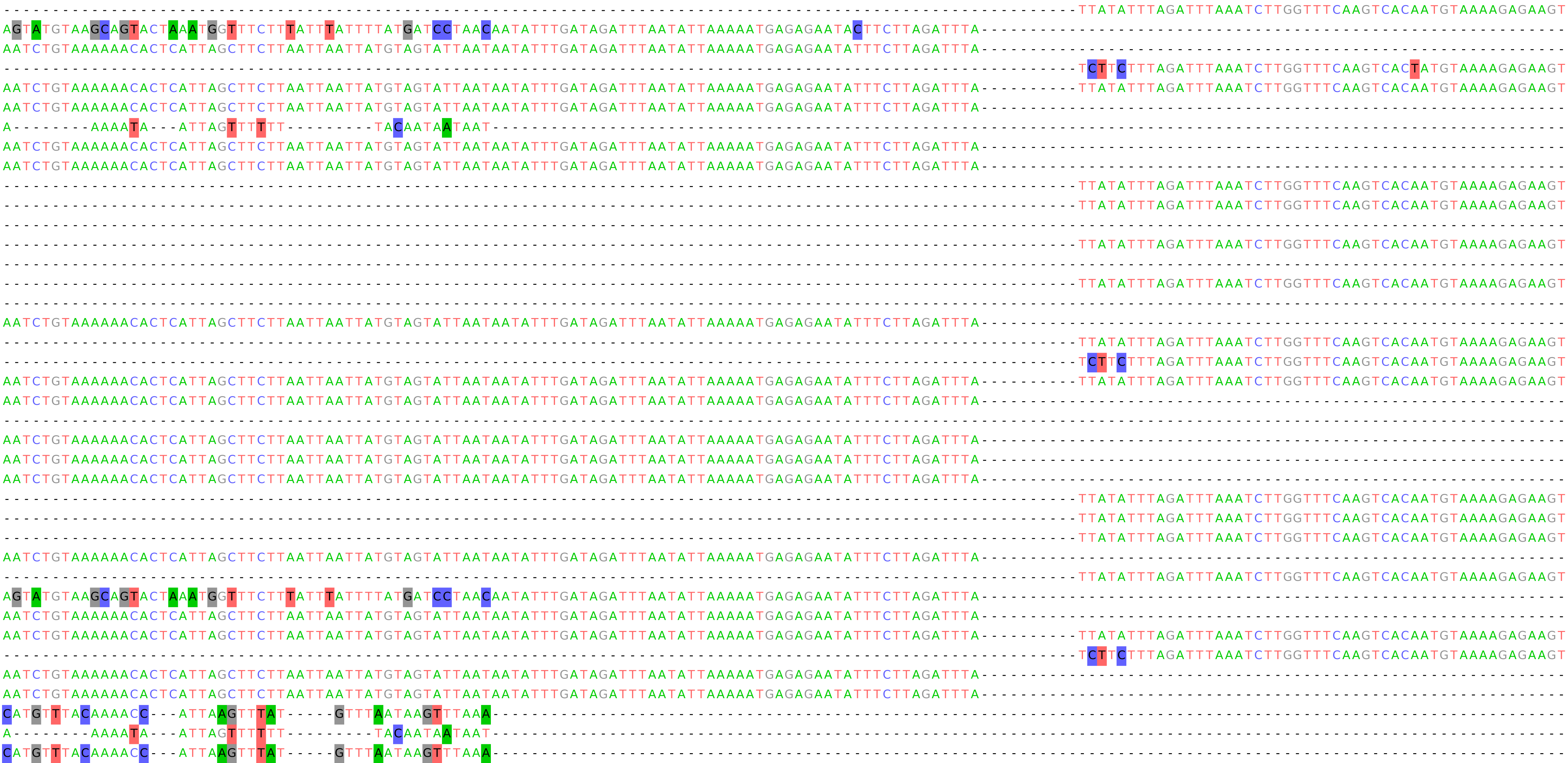
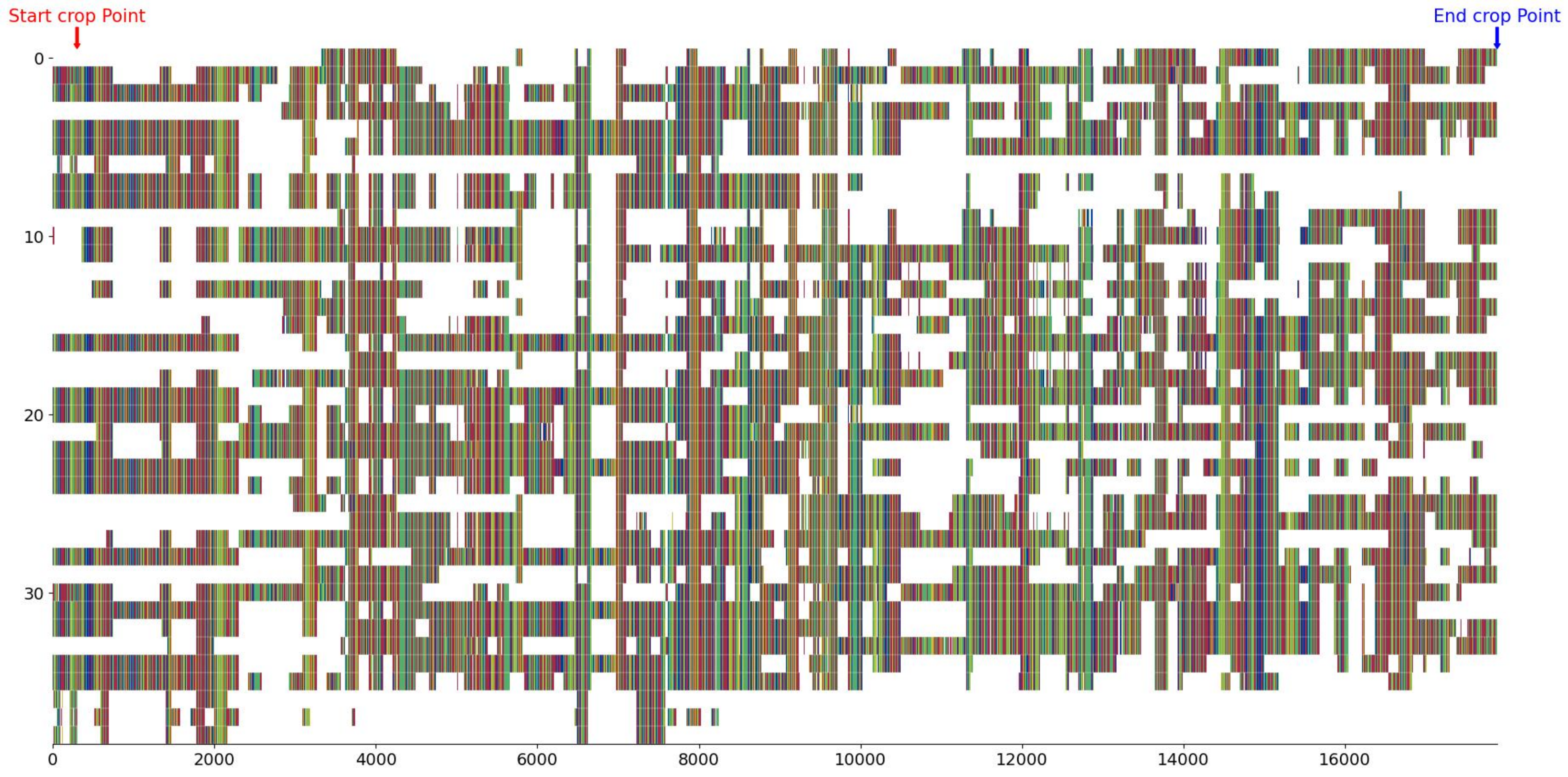


Start crop Point

End crop Point

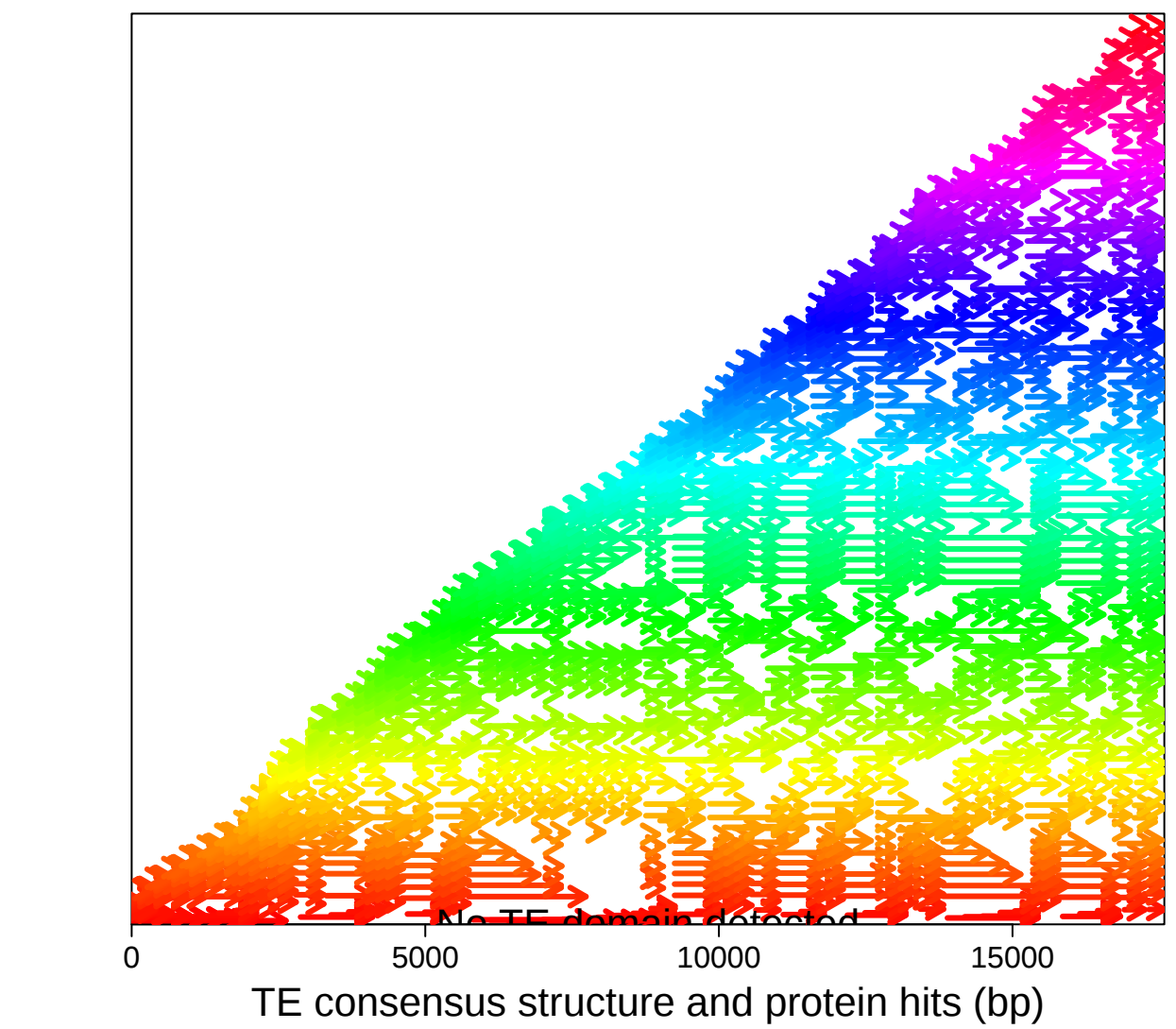
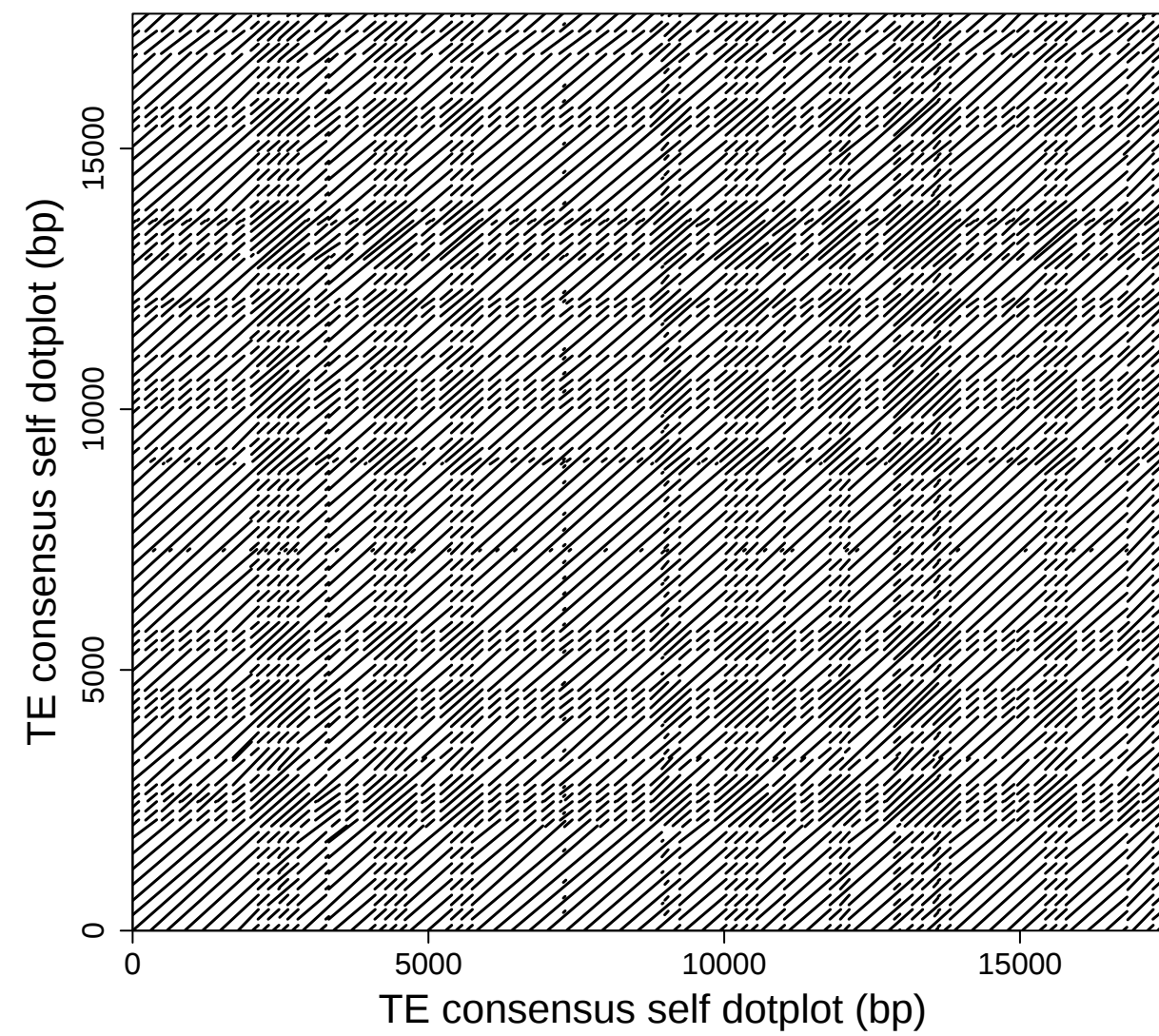
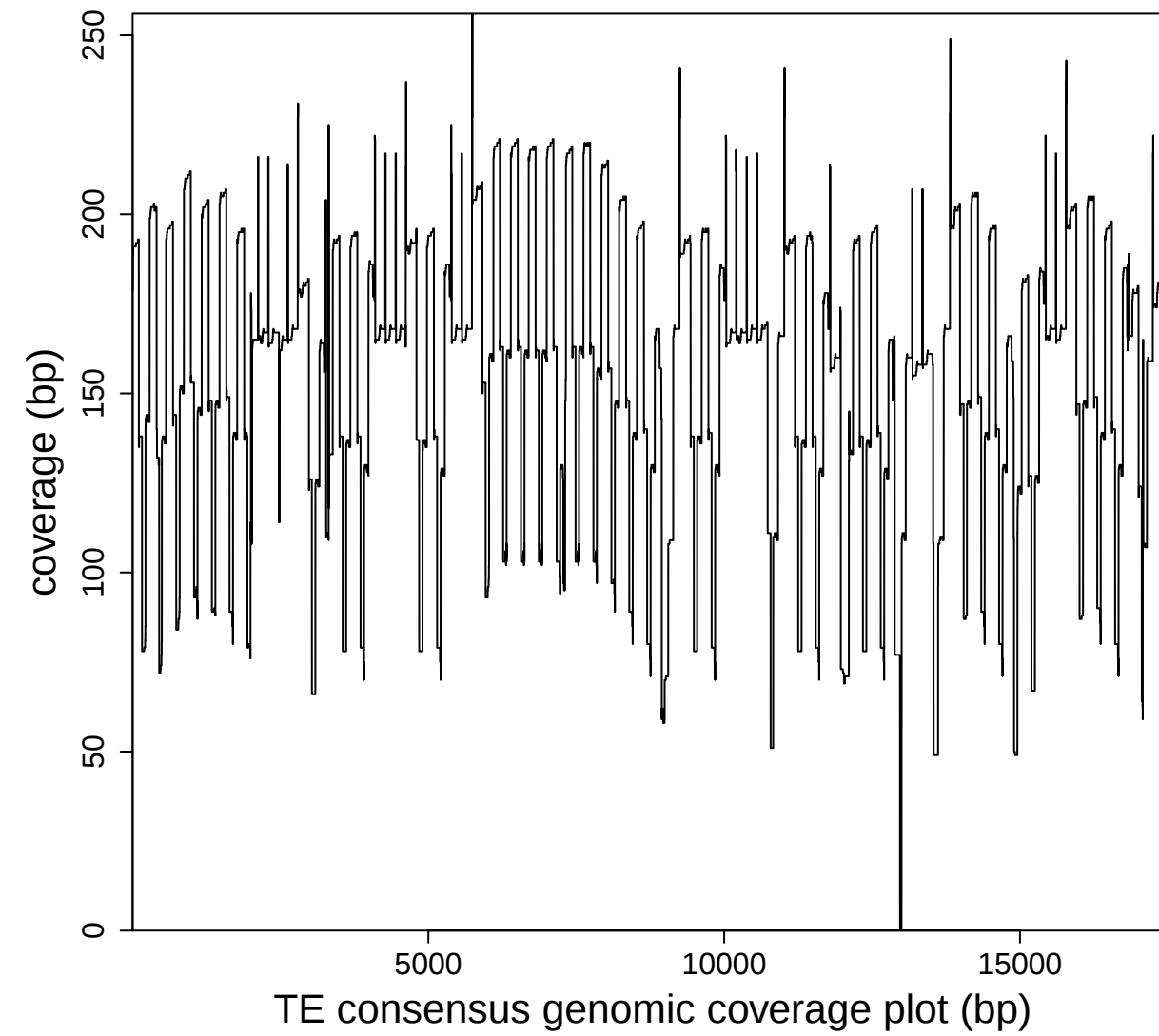
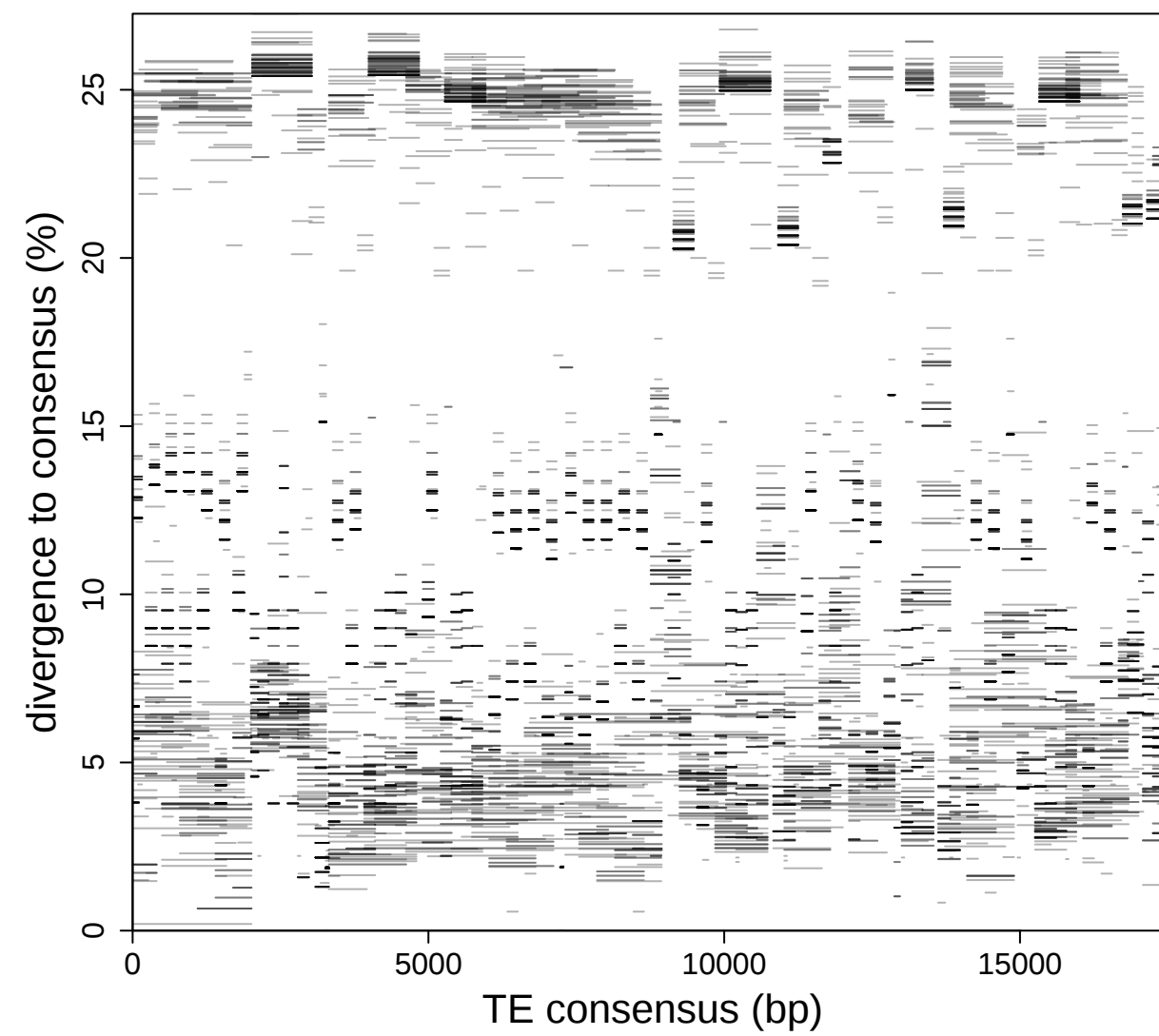
MSA length = 17585





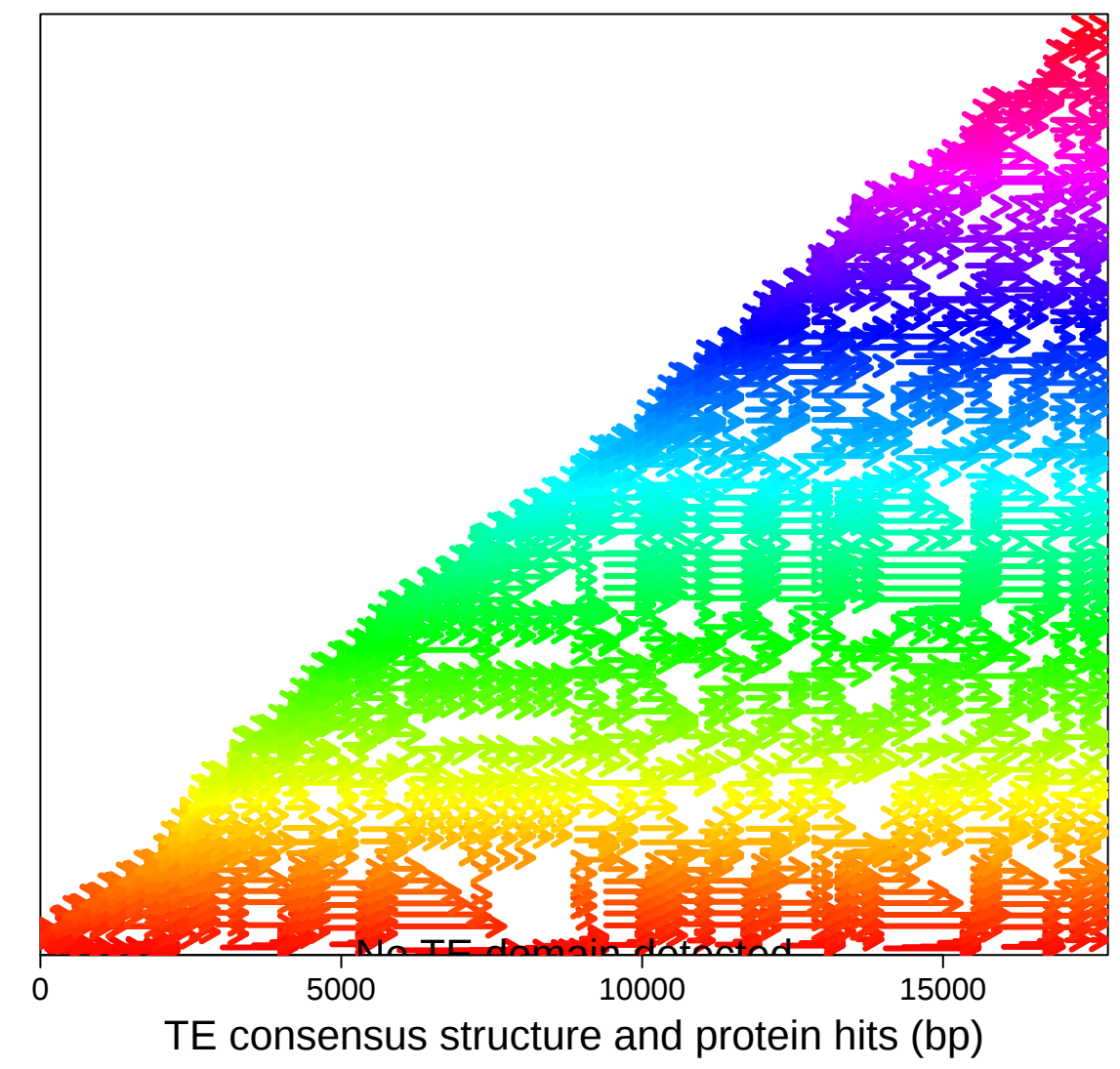
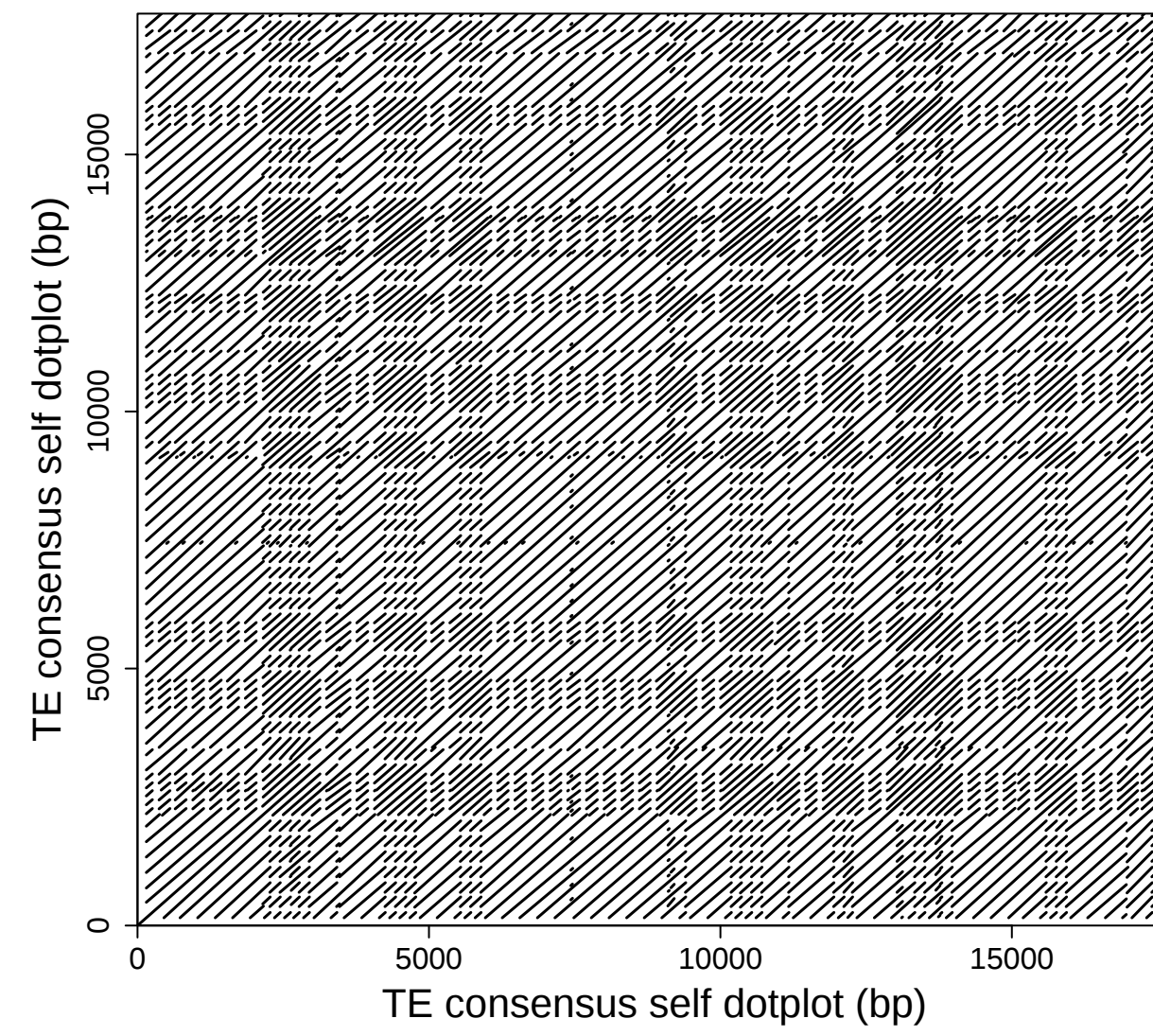
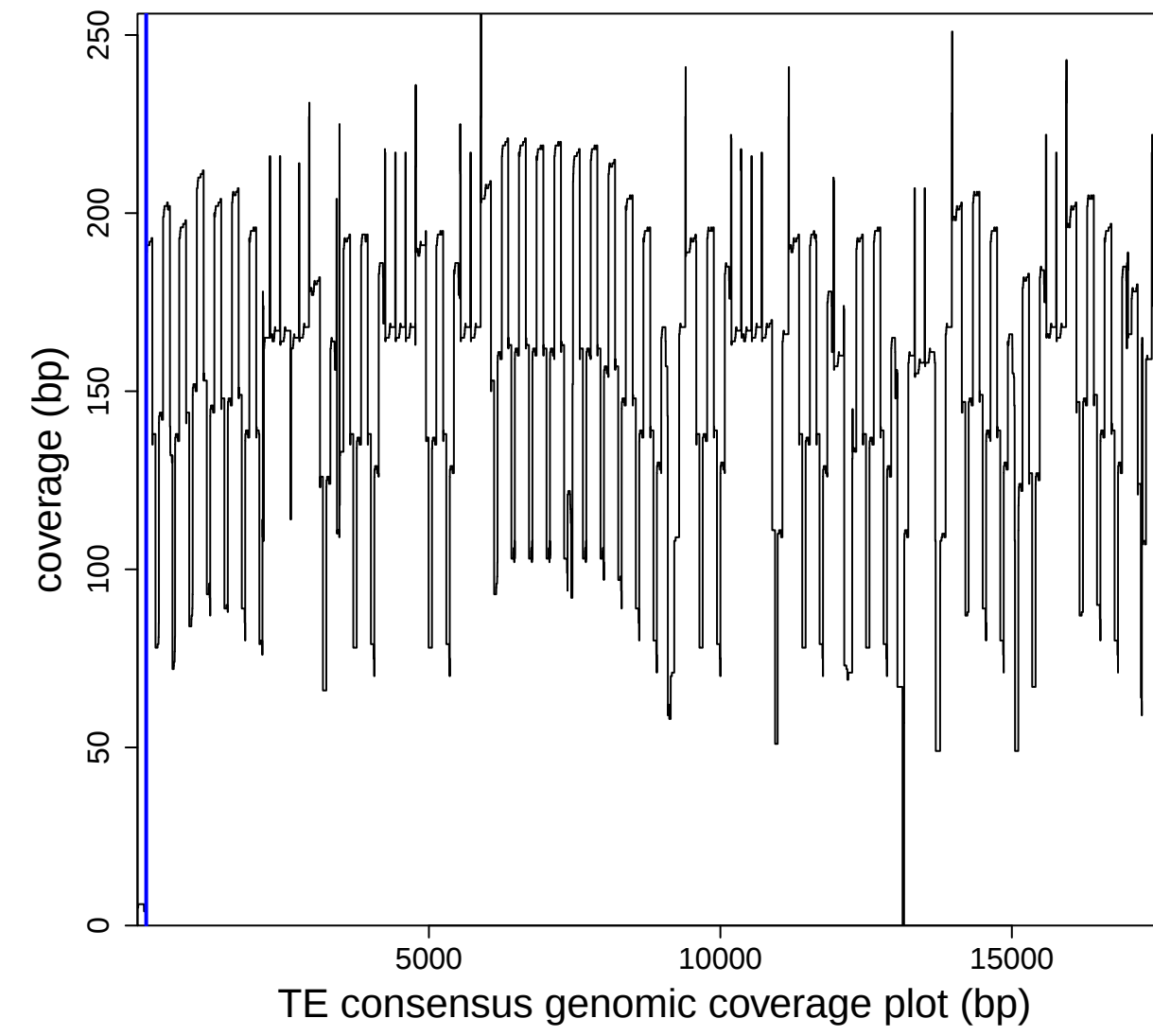
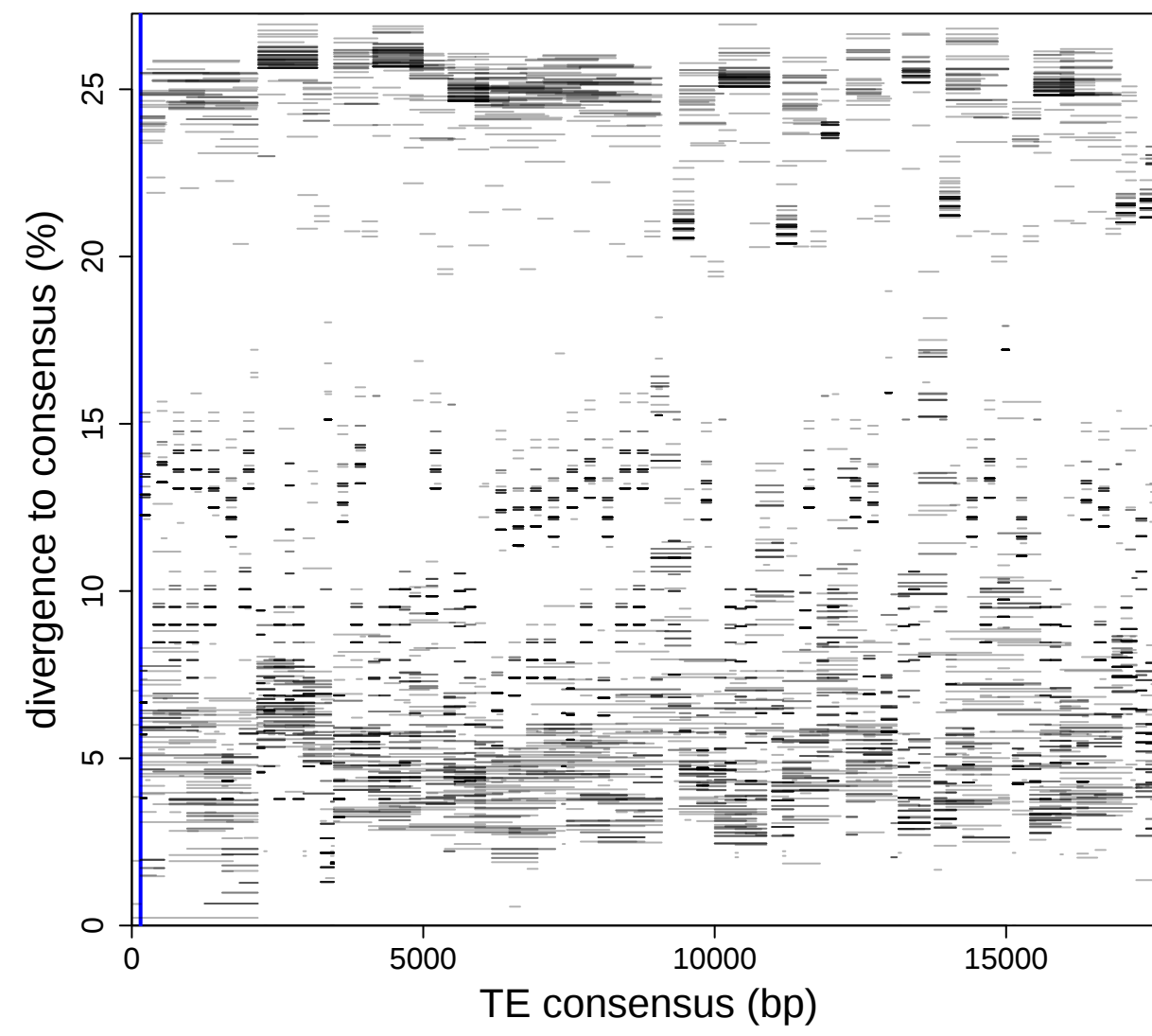
ed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.fa_ce.f
size: 17585bp; fragments: 7554; full length: 0 (>=15826.5bp)

After TEtrimmer 17585 bp

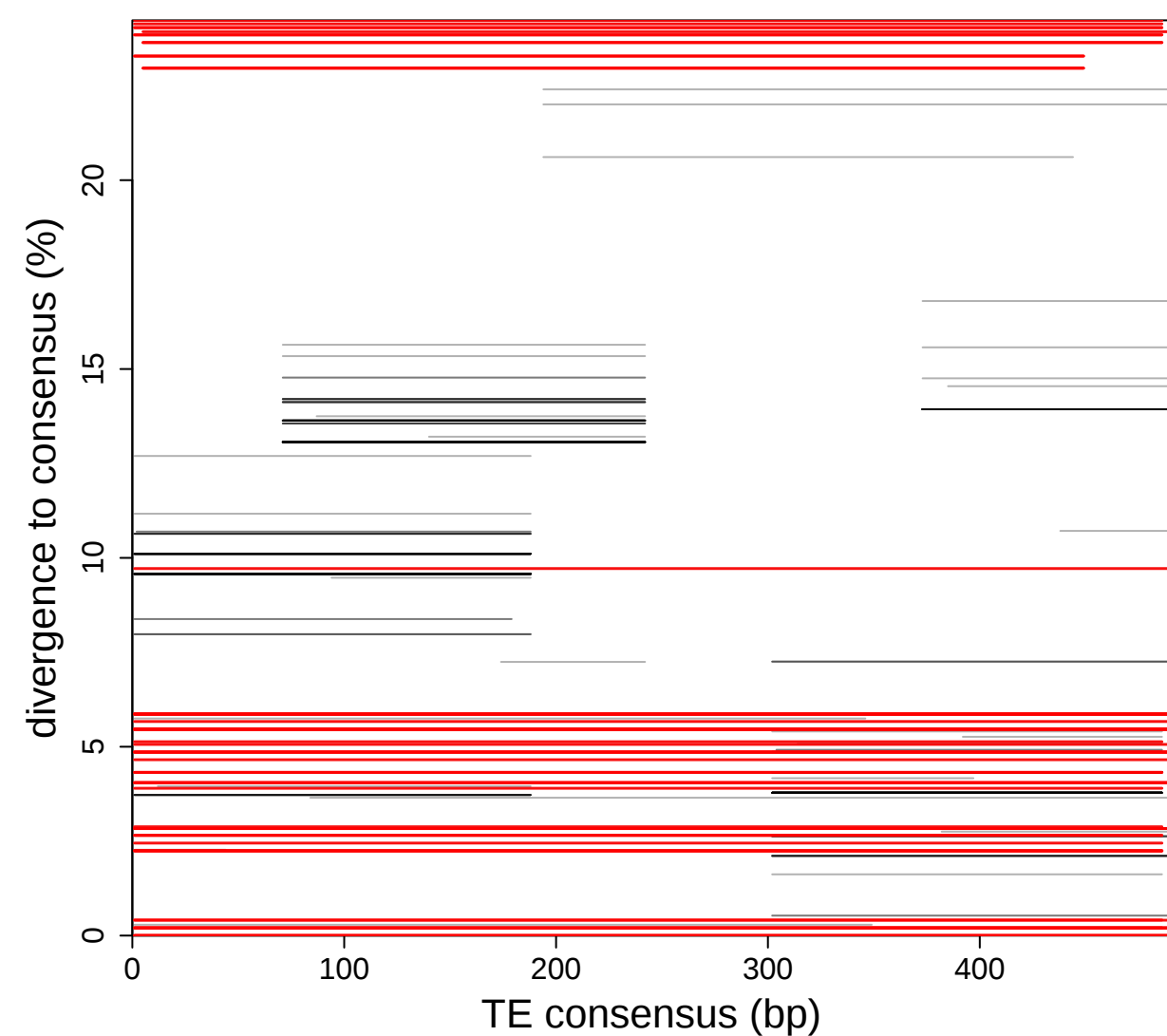


ily_68.fasta.b.bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa
size: 17740bp; fragments: 7537; full length: 0 (>=15966bp)

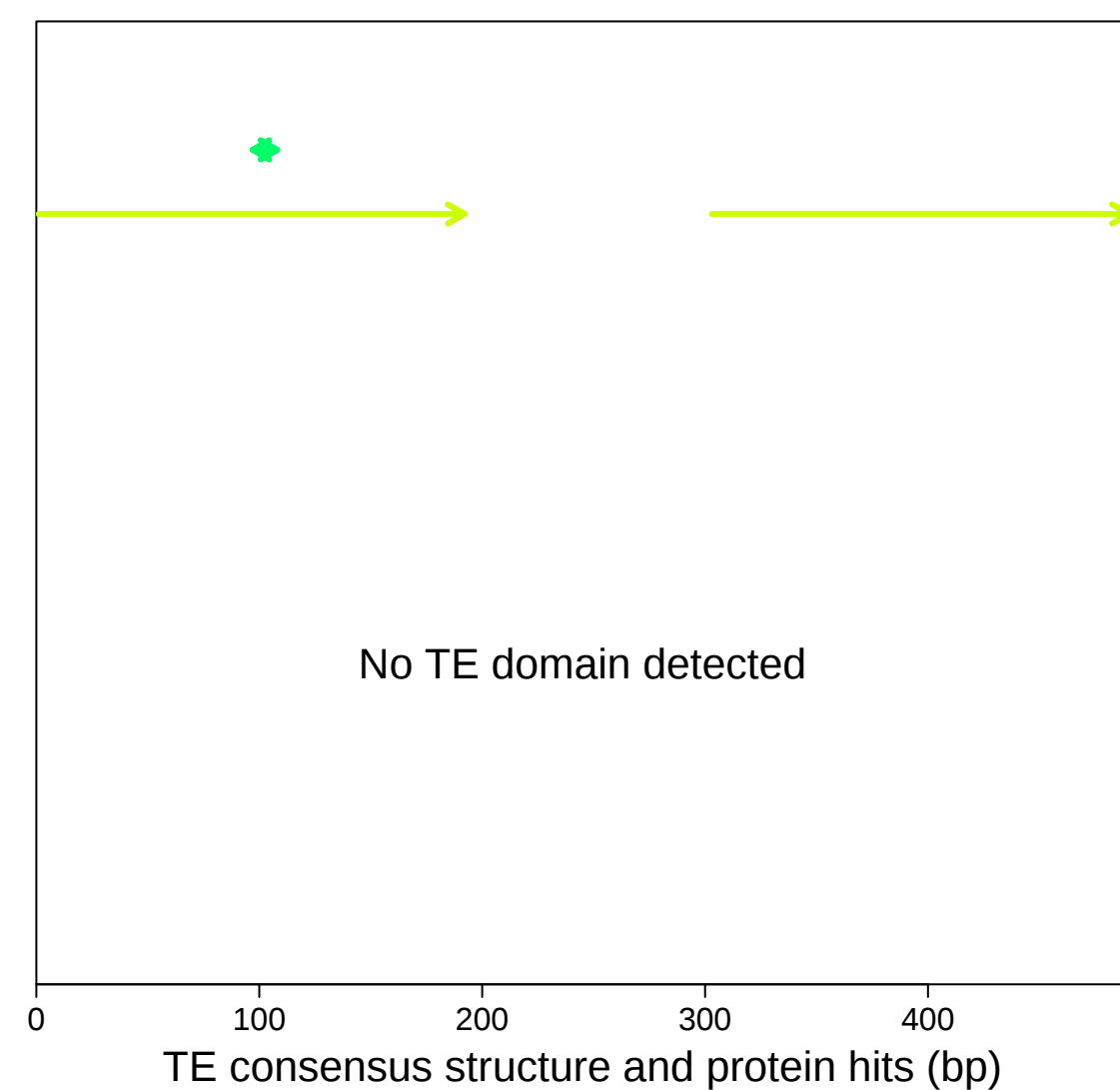
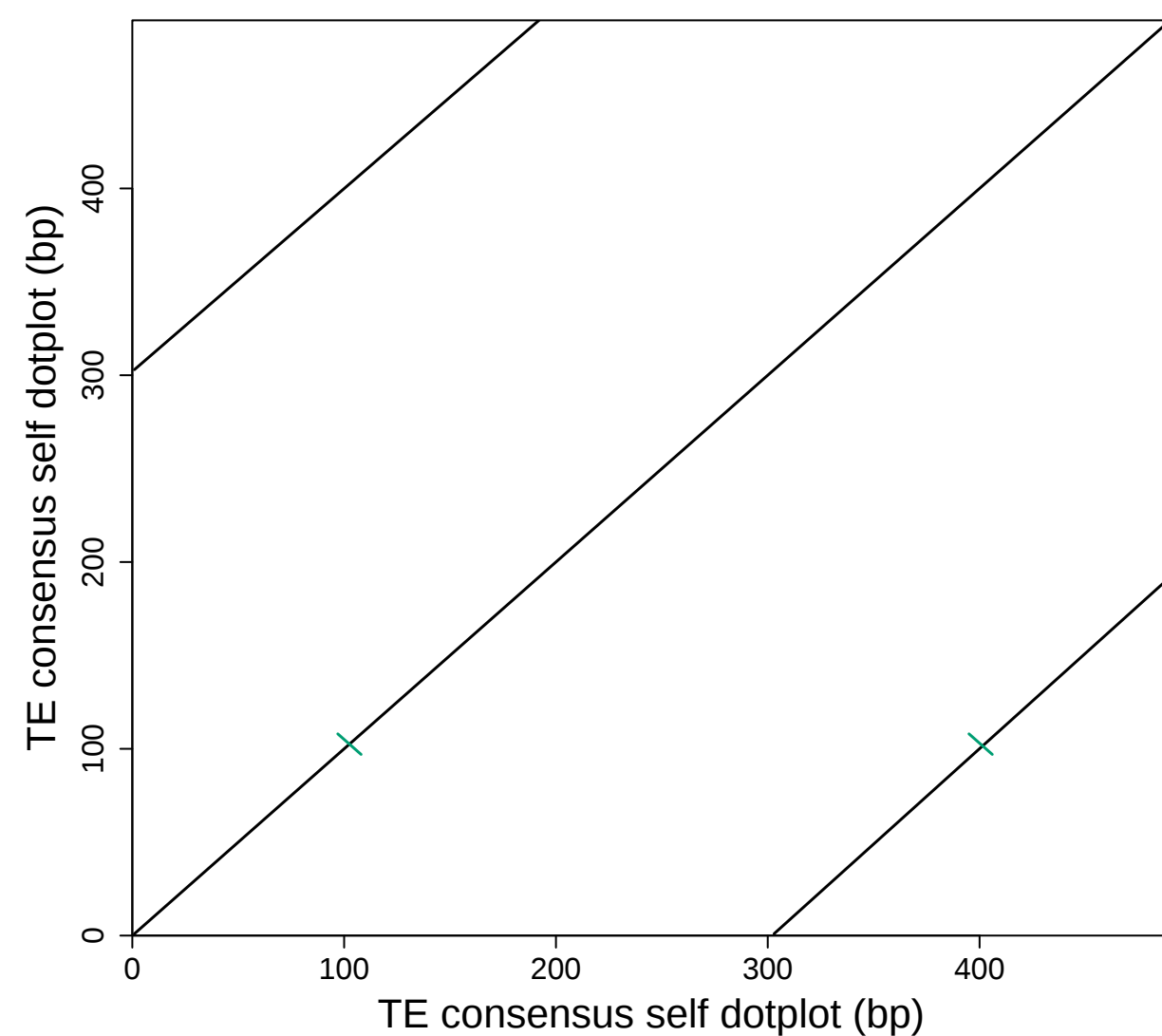
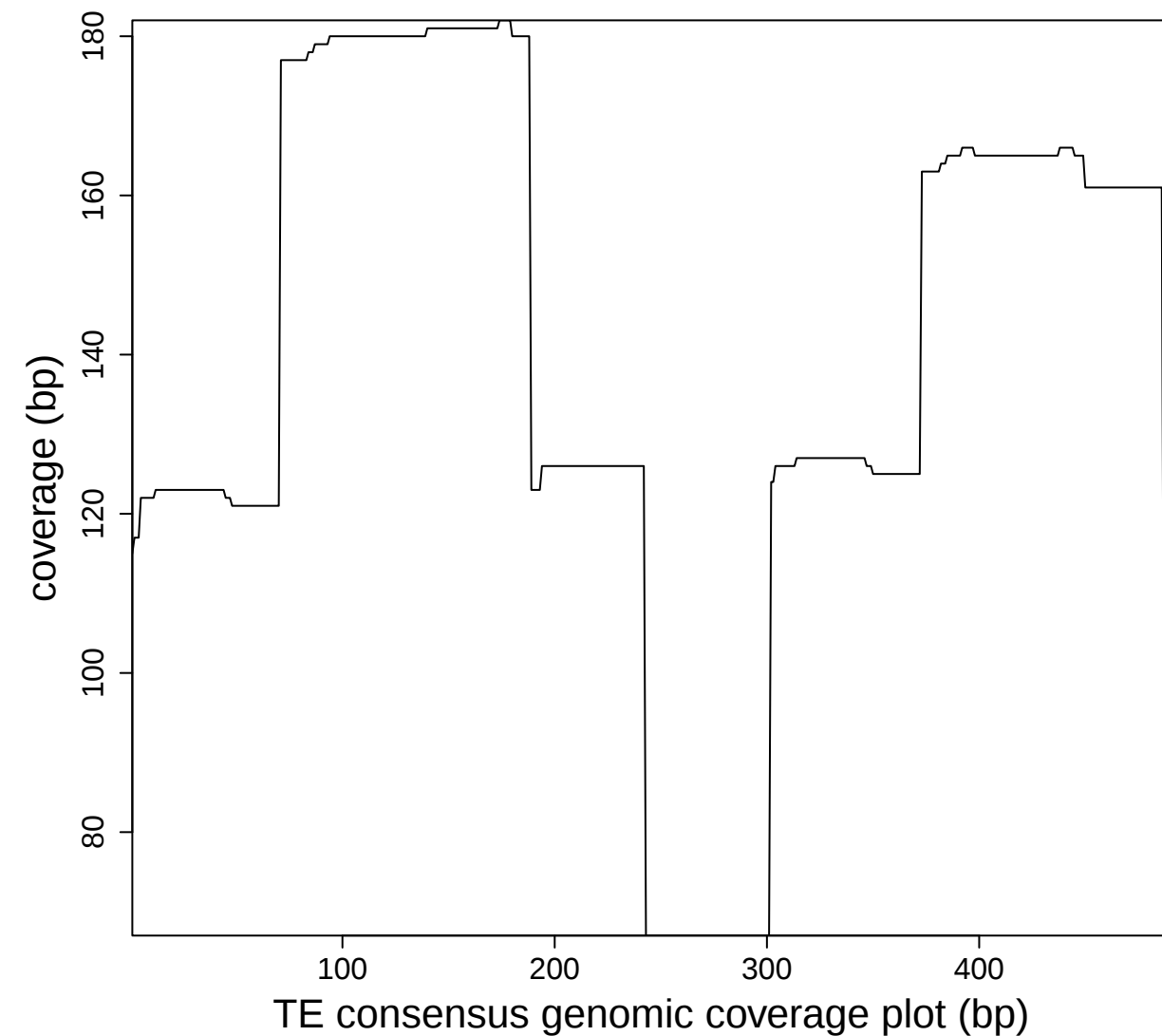
After TEtrimmer Extended plot Blue lines are boundaries



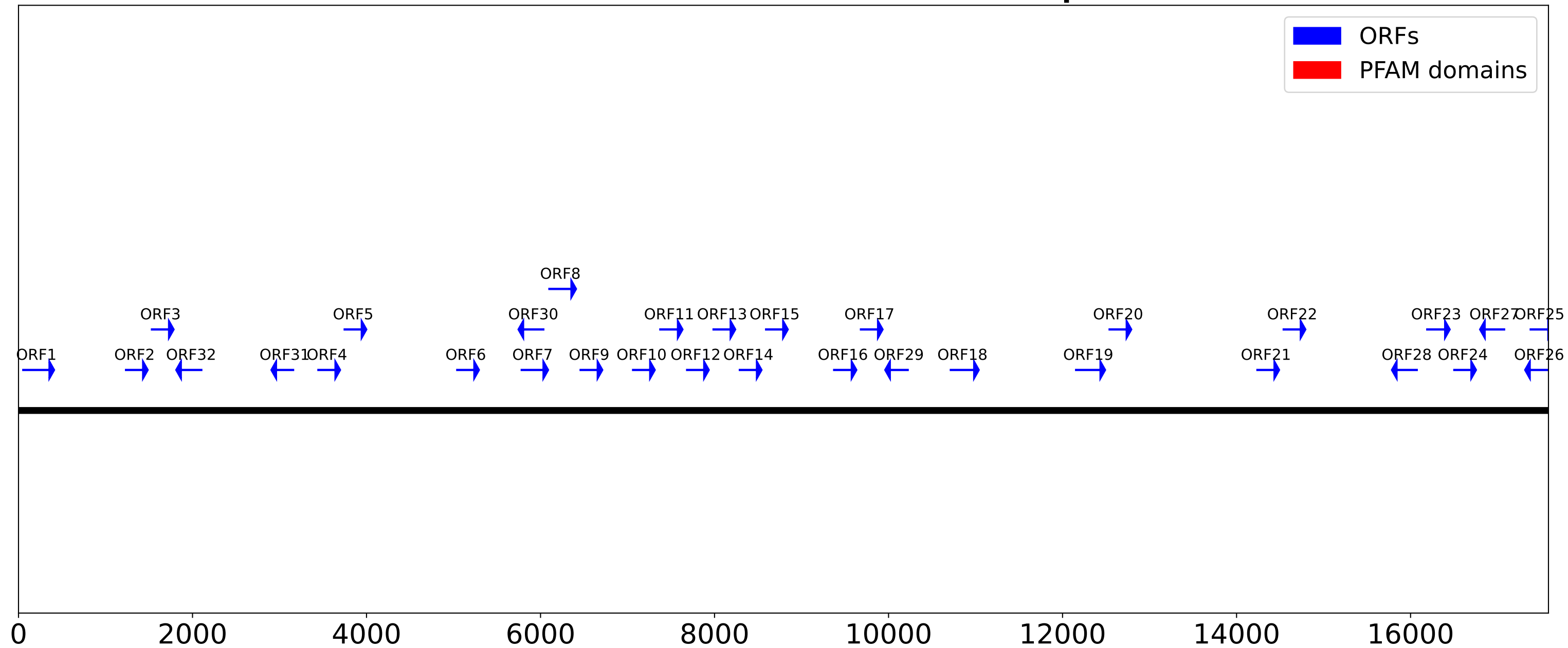
TE: ltr_1_family_68
size: 490bp; fragments: 289; full length: 61 (≥ 441 bp)



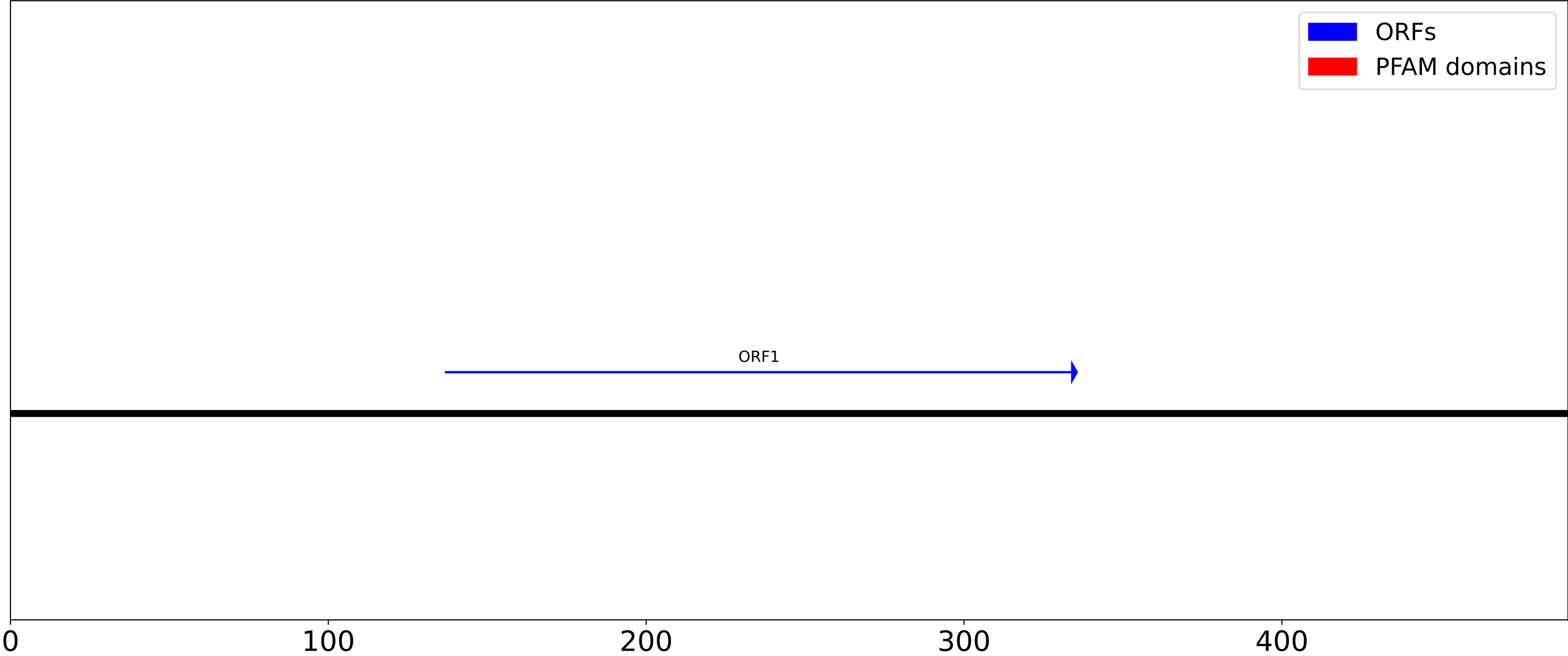
Before TEtrimmer 490 bp



After TEtrimmer ORF and PFAM domain plot



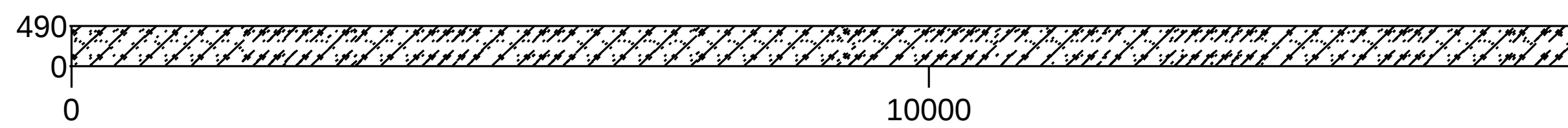
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)