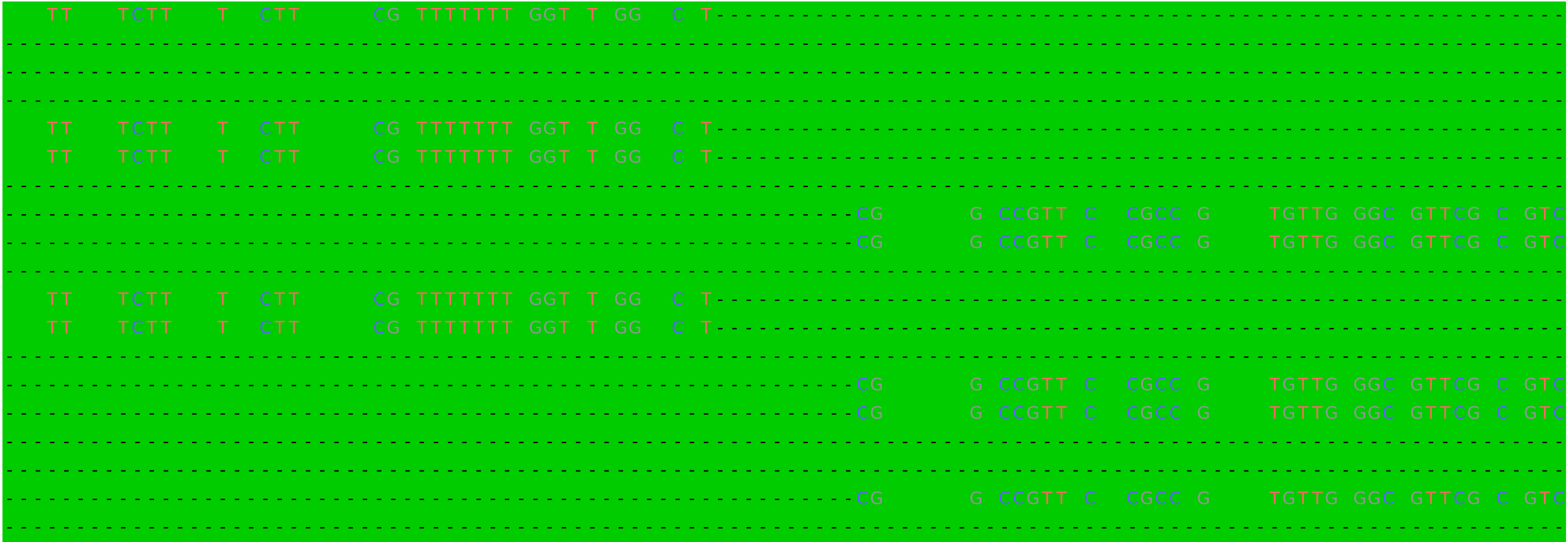
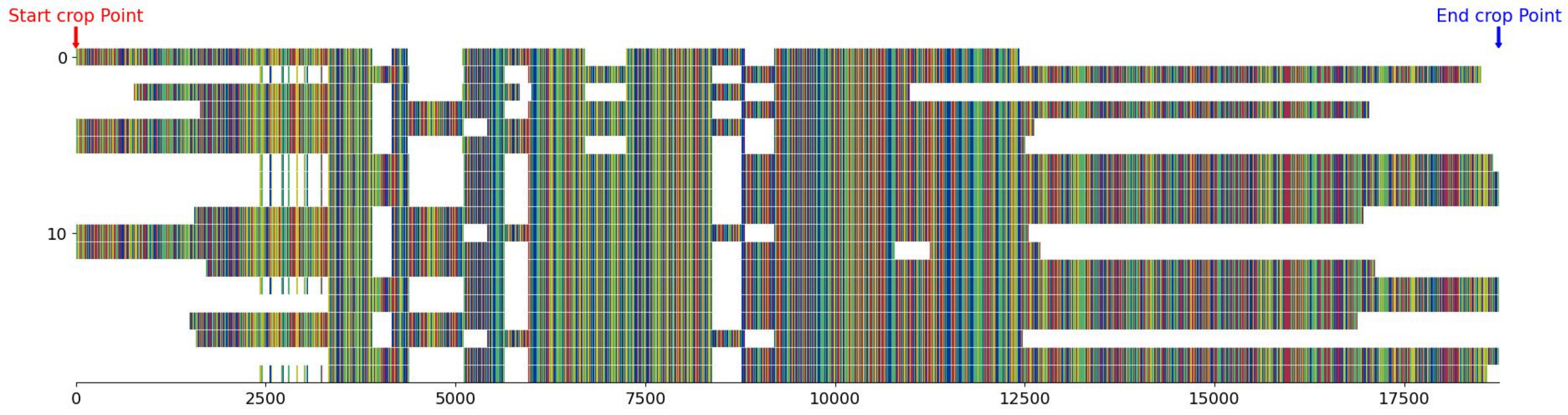


Start crop Point

End crop Point

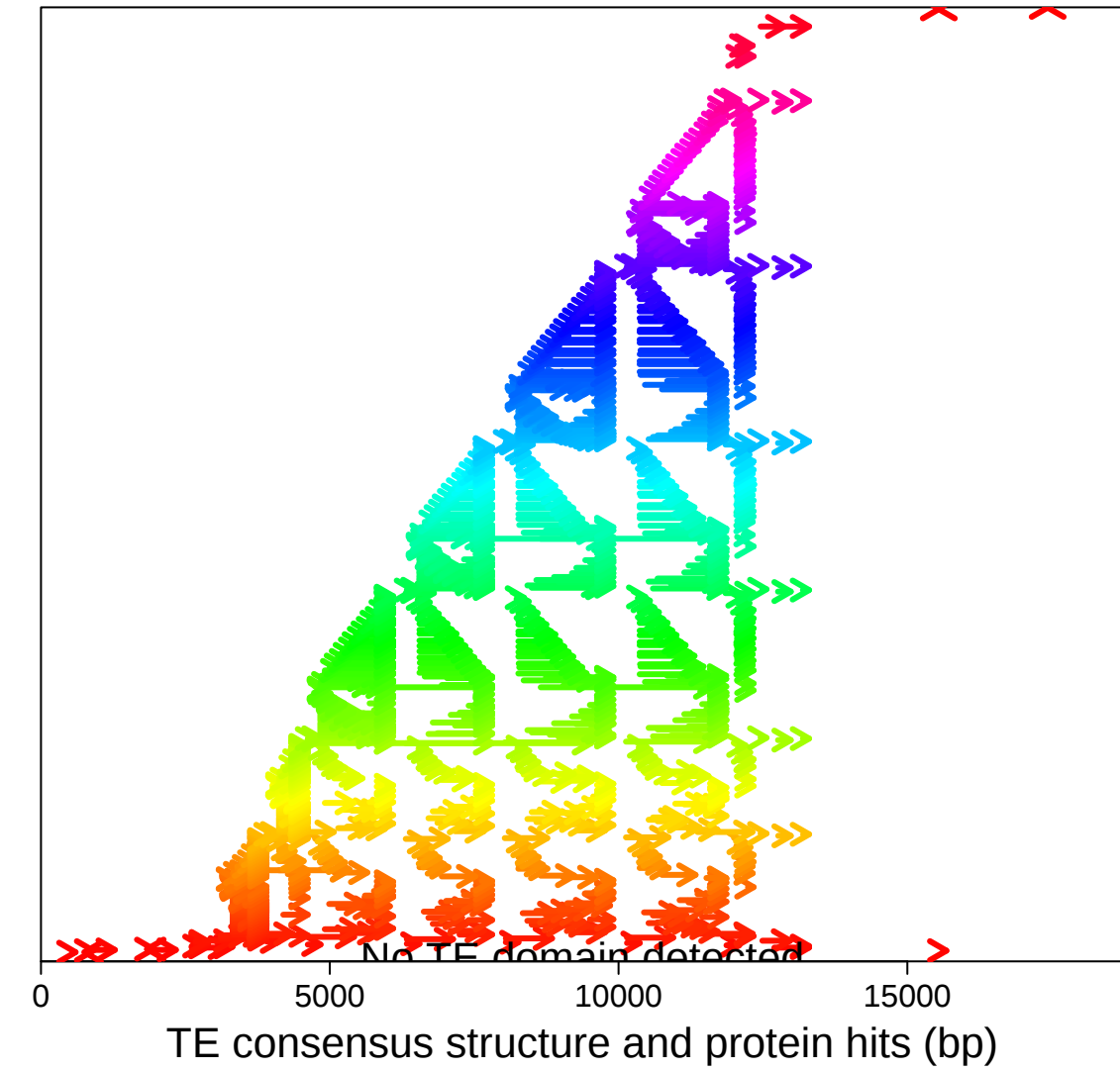
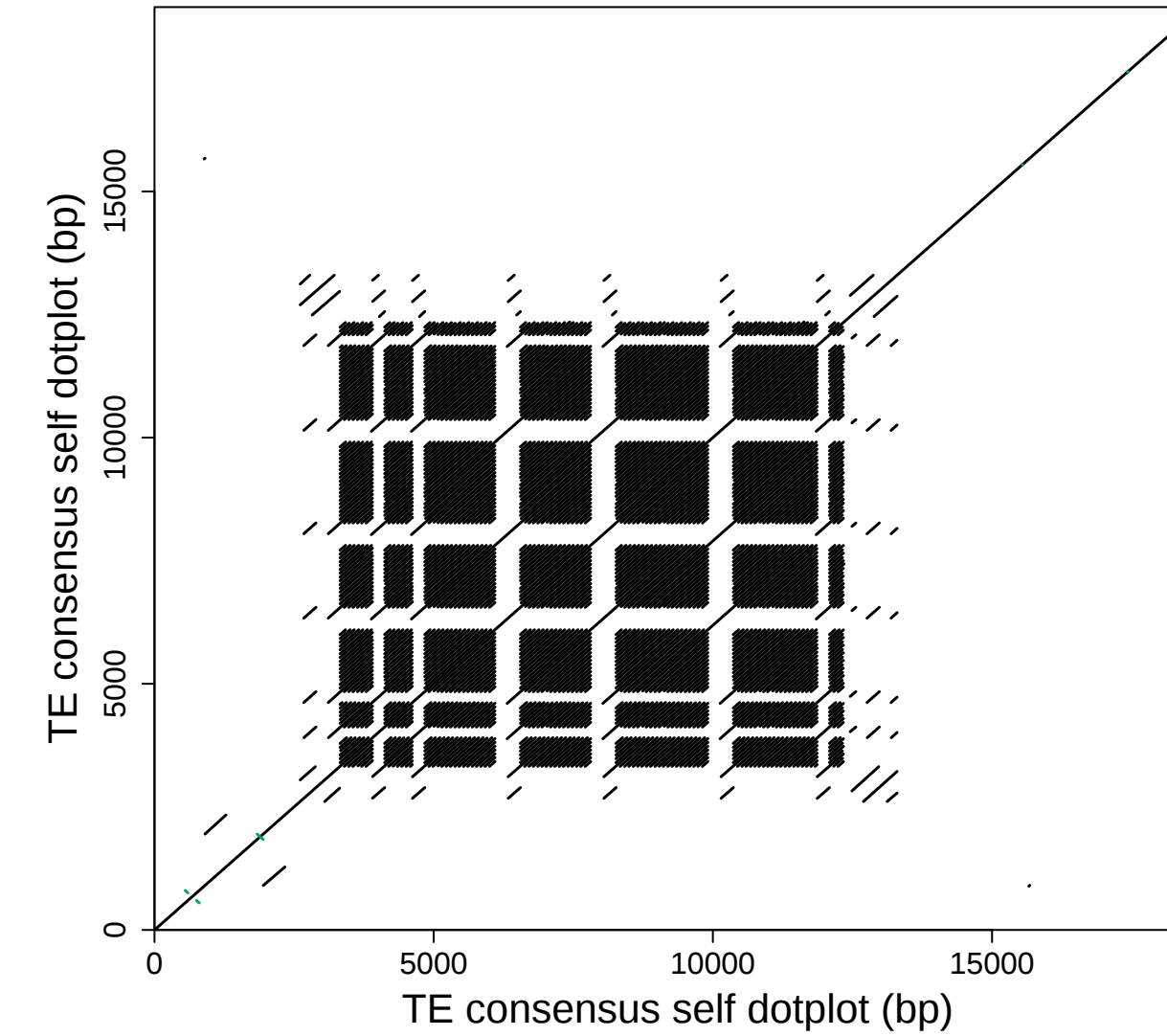
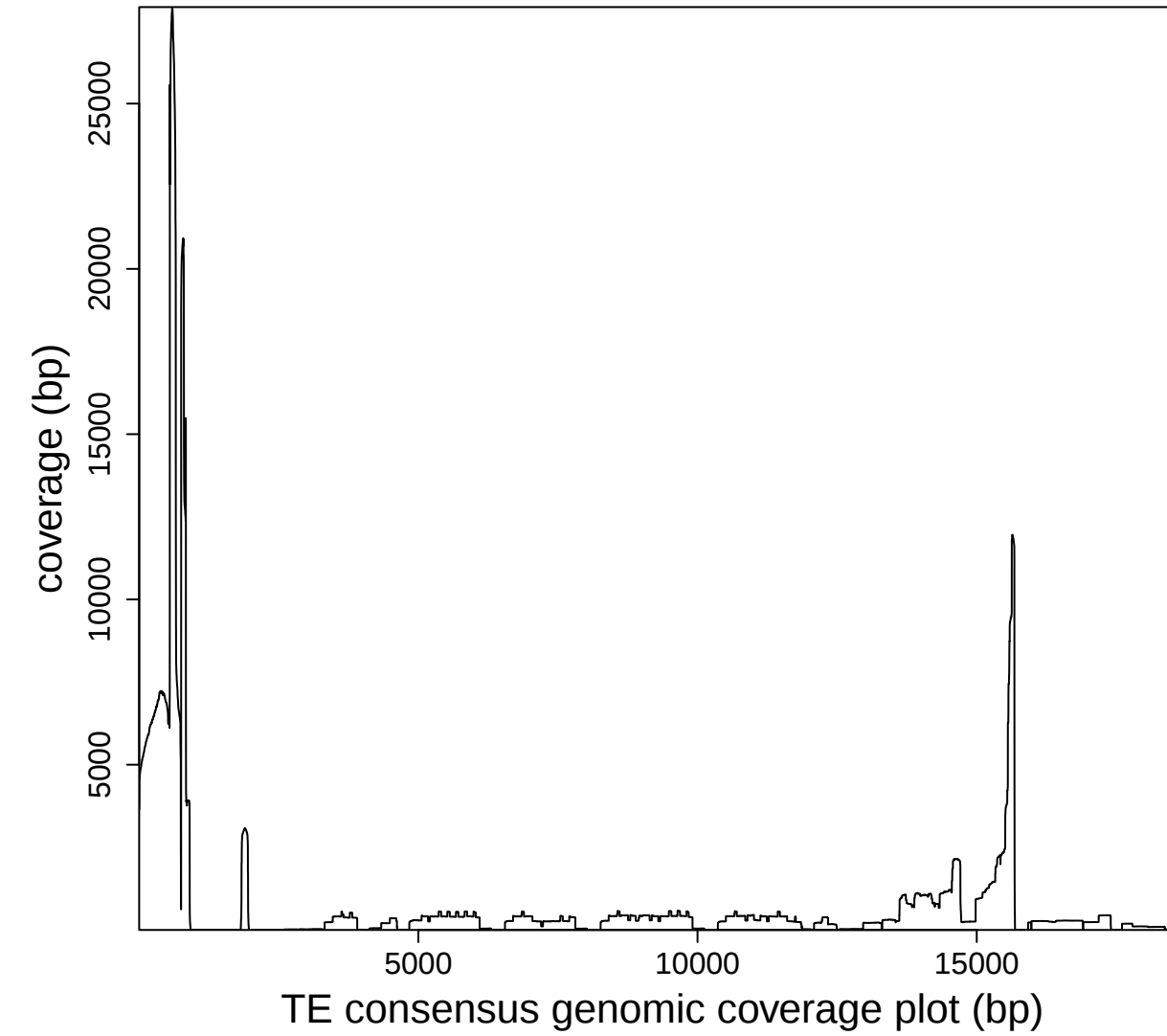
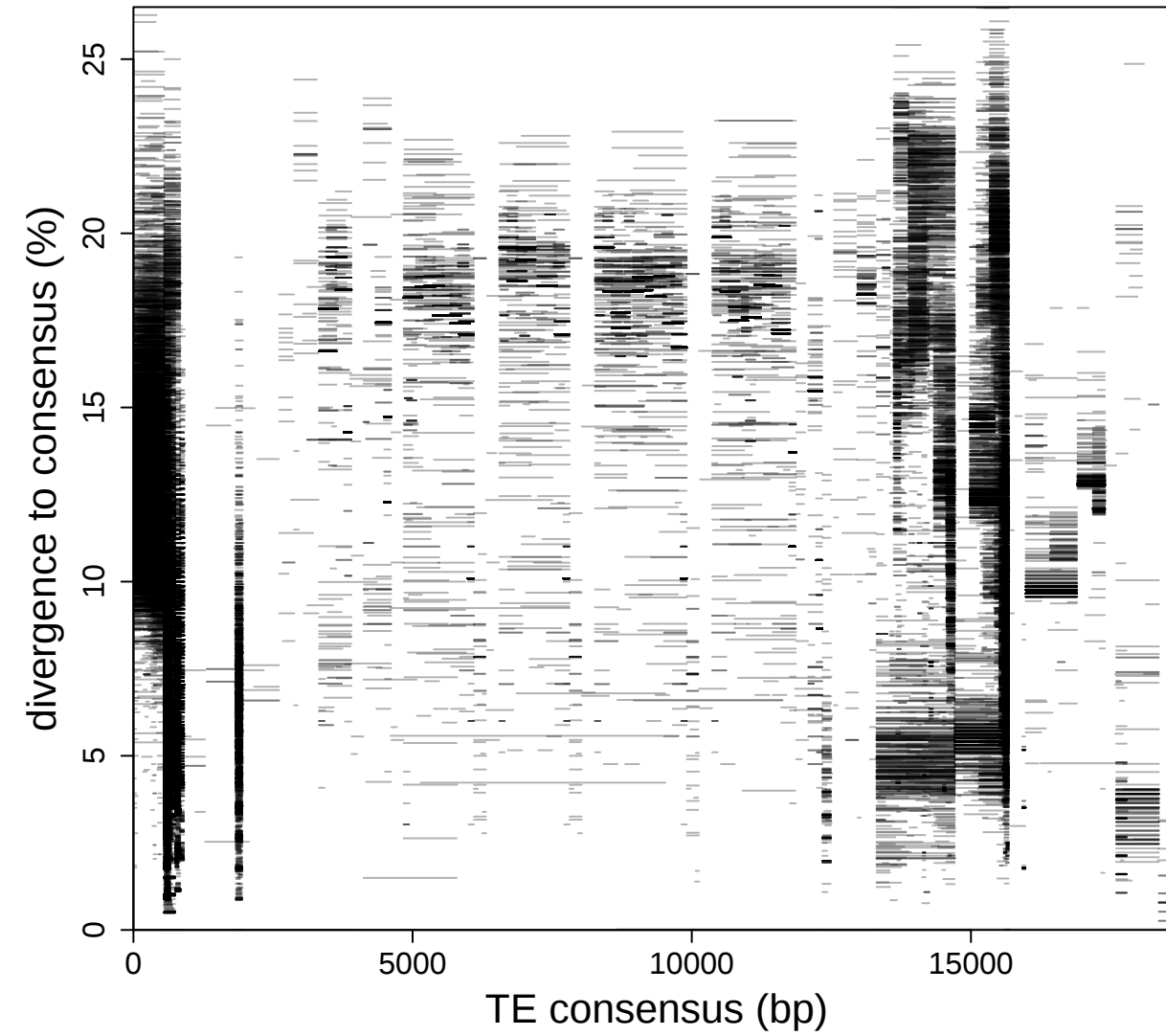
MSA length = 18744





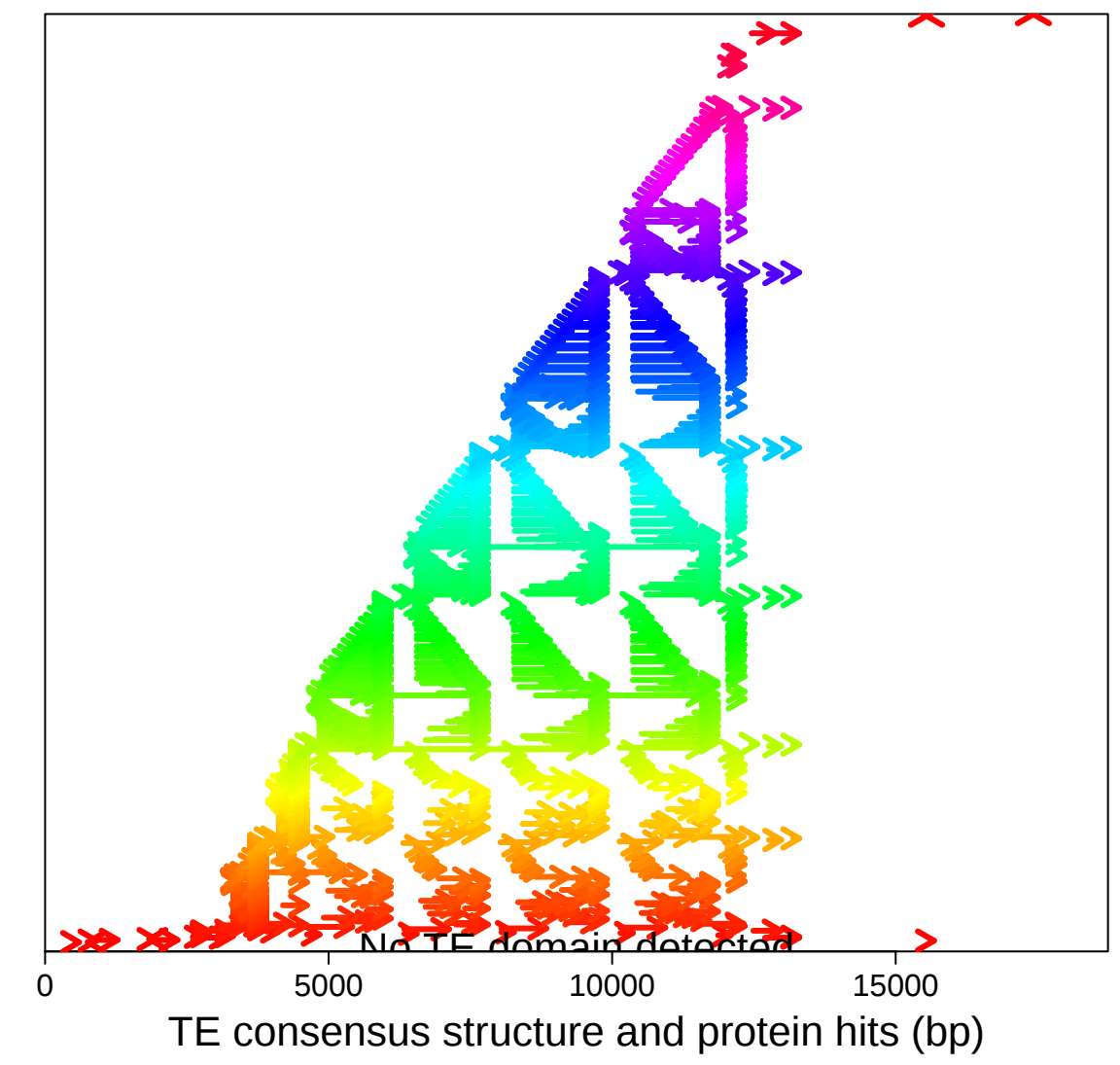
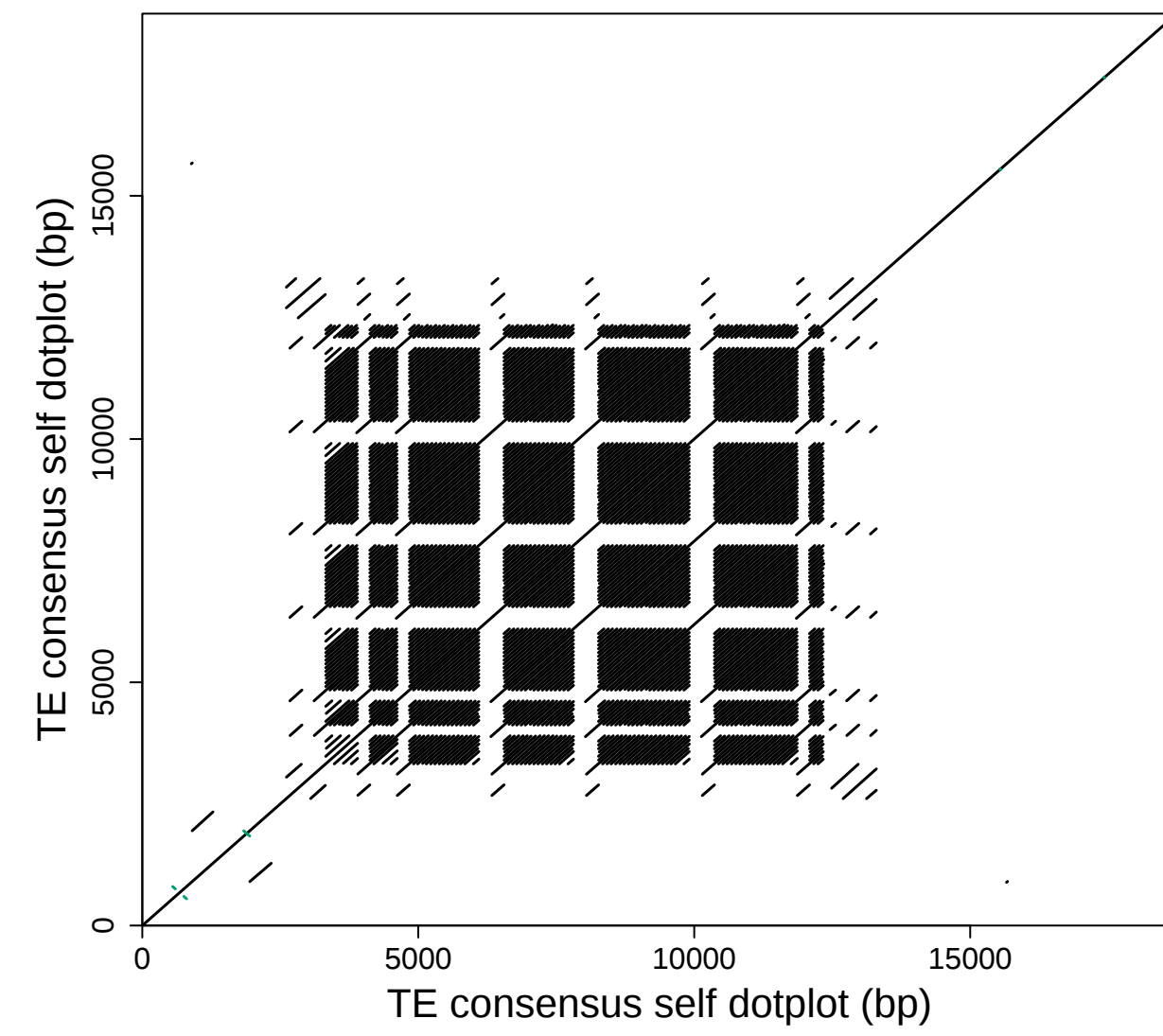
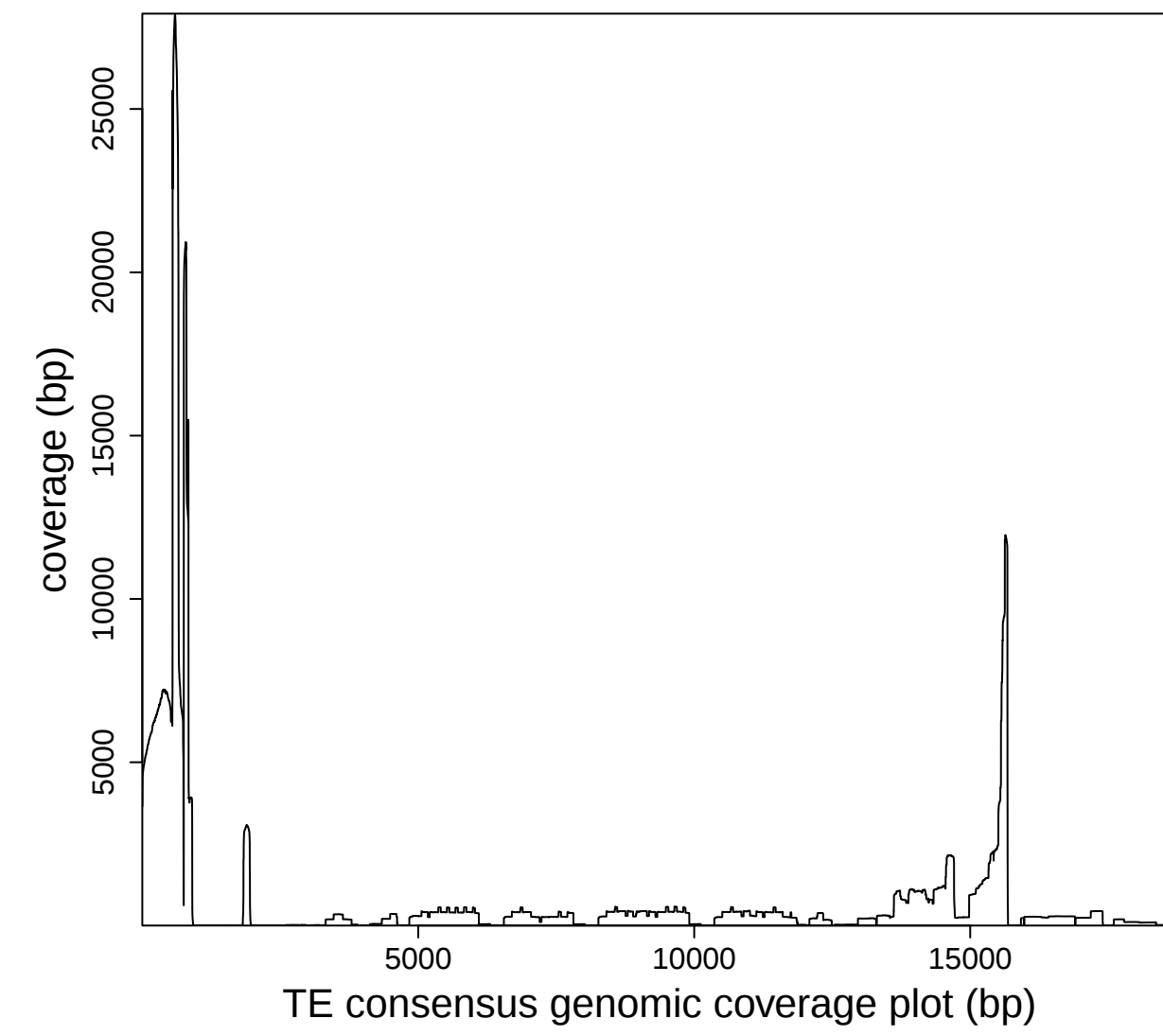
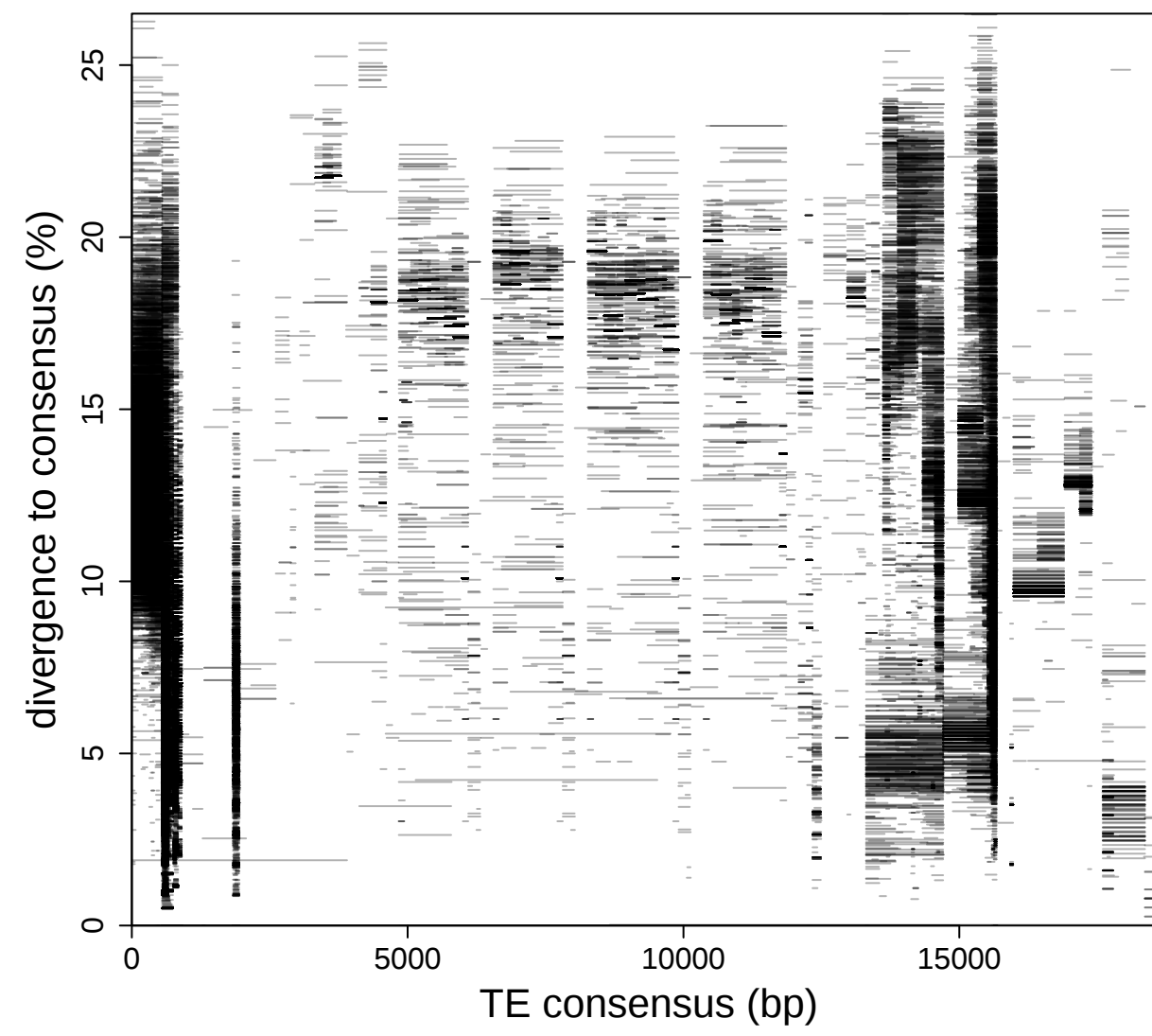
l.b.bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcIn.fa_aln.fa_cl.fa_gs.fa
size: 18744bp; fragments: 90741; full length: 0 (>=16869.6bp)

After TEtrimmer 18744 bp



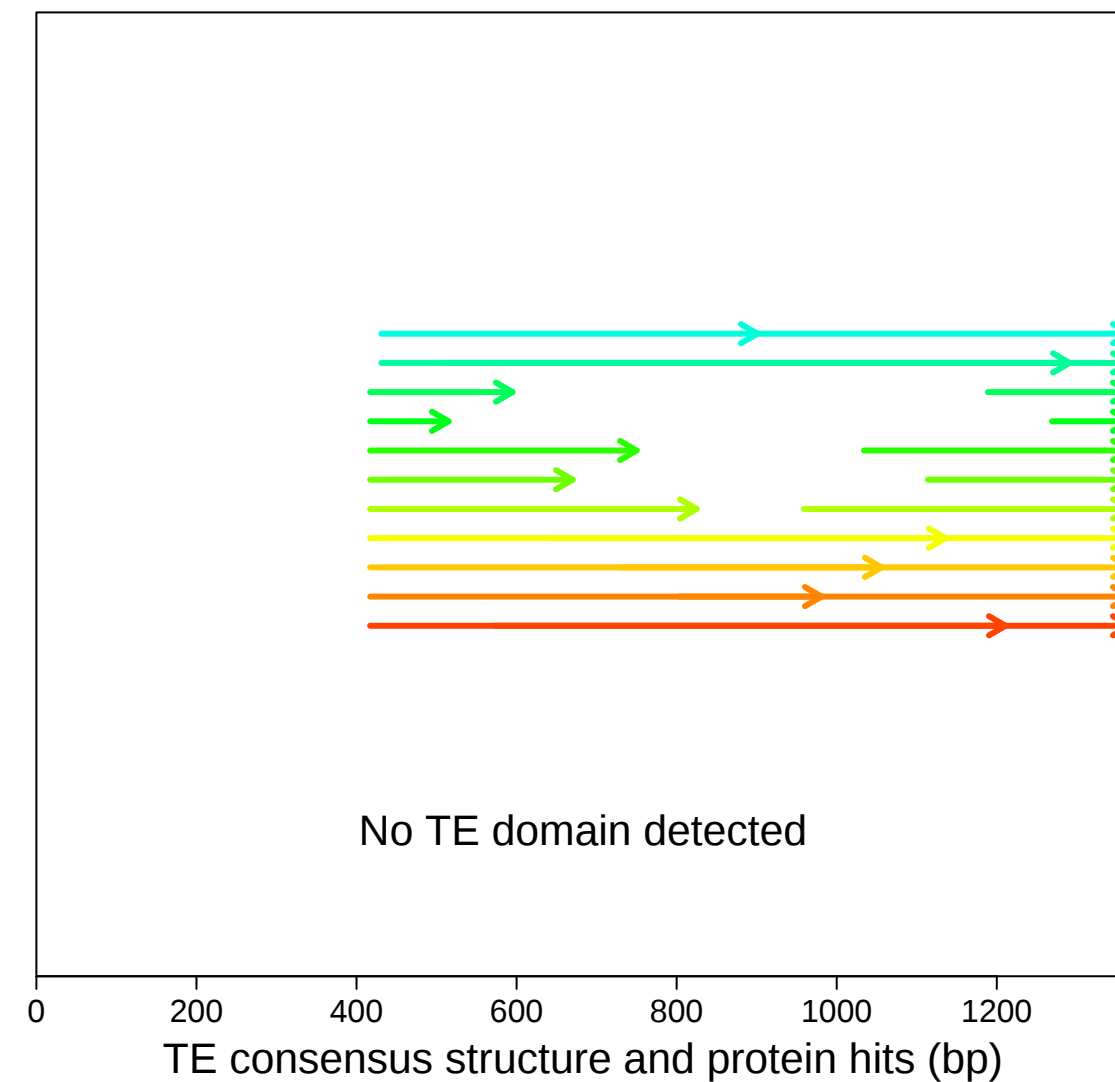
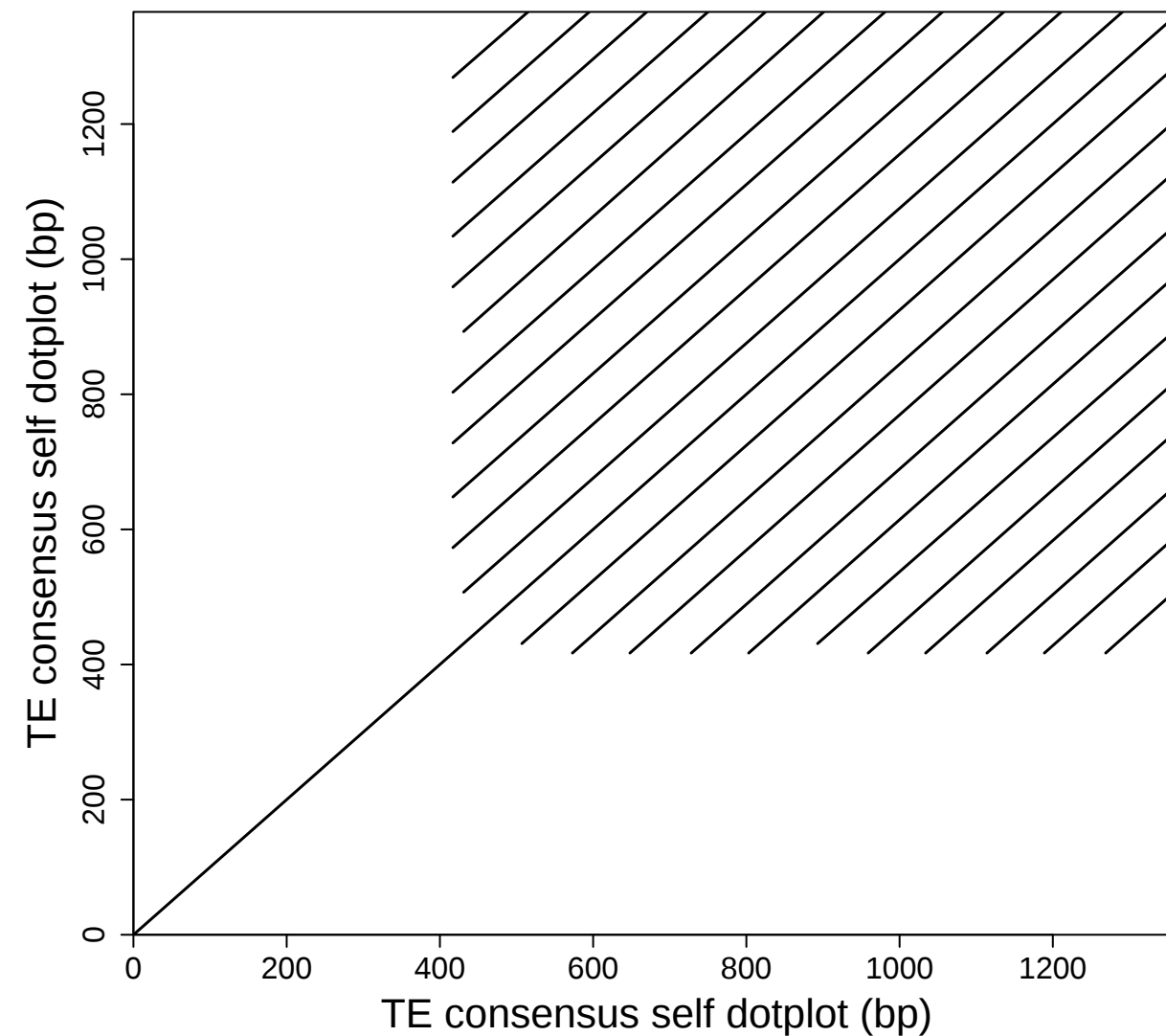
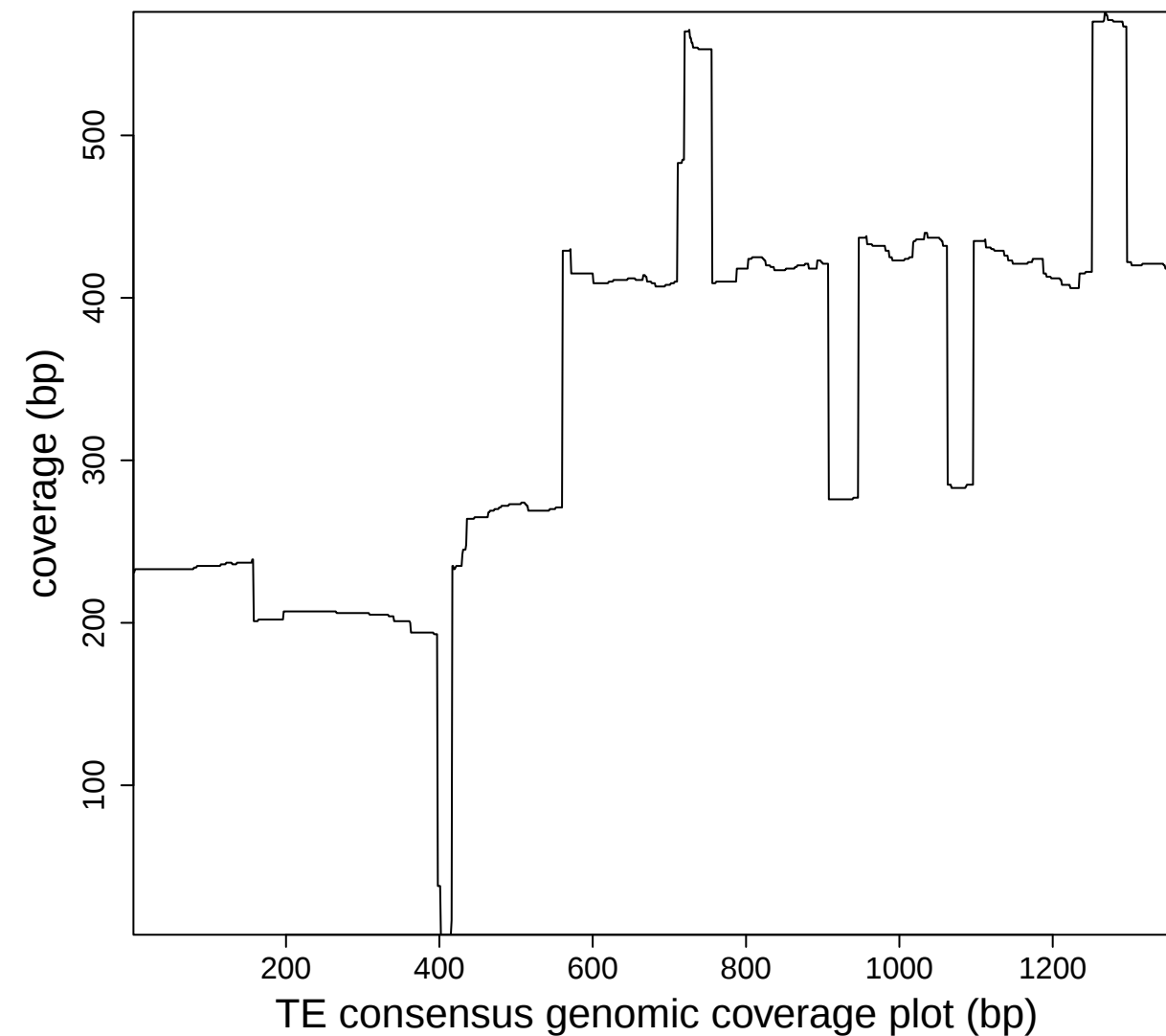
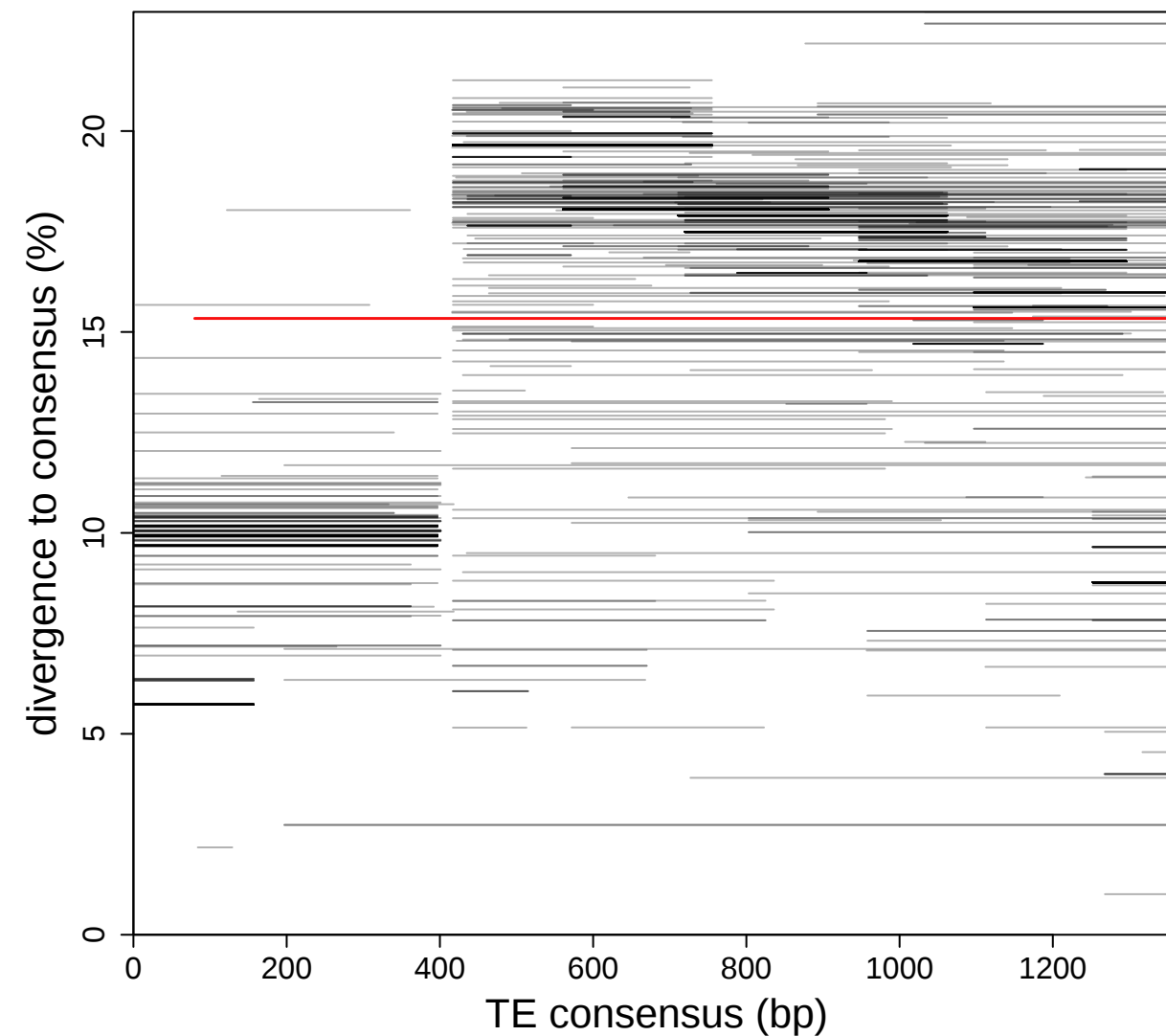
5.fasta.b.bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa
size: 18744bp; fragments: 90350; full length: 0 (>=16869.6bp)

After TEtrimmer Extended plot Blue lines are boundaries

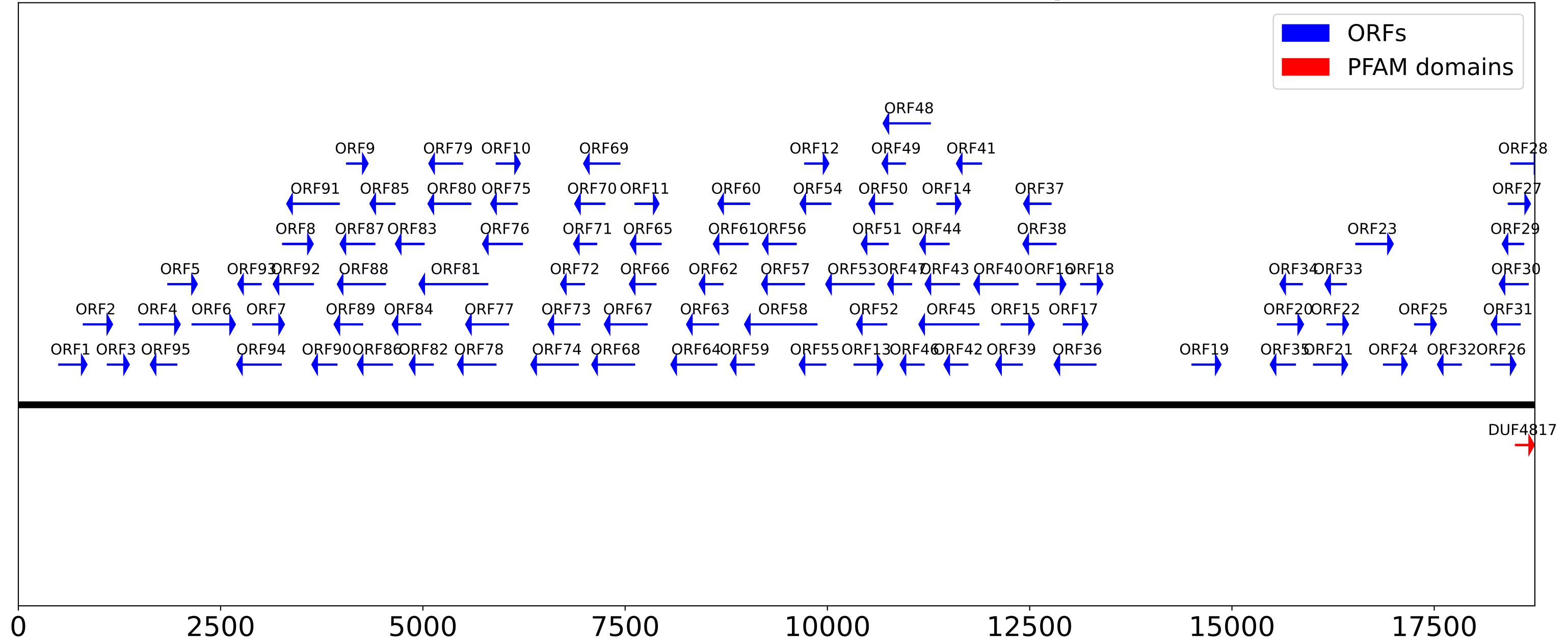


TE: rnd_1_family_825
size: 1366bp; fragments: 1391; full length: 1 (≥ 1229.4 bp)

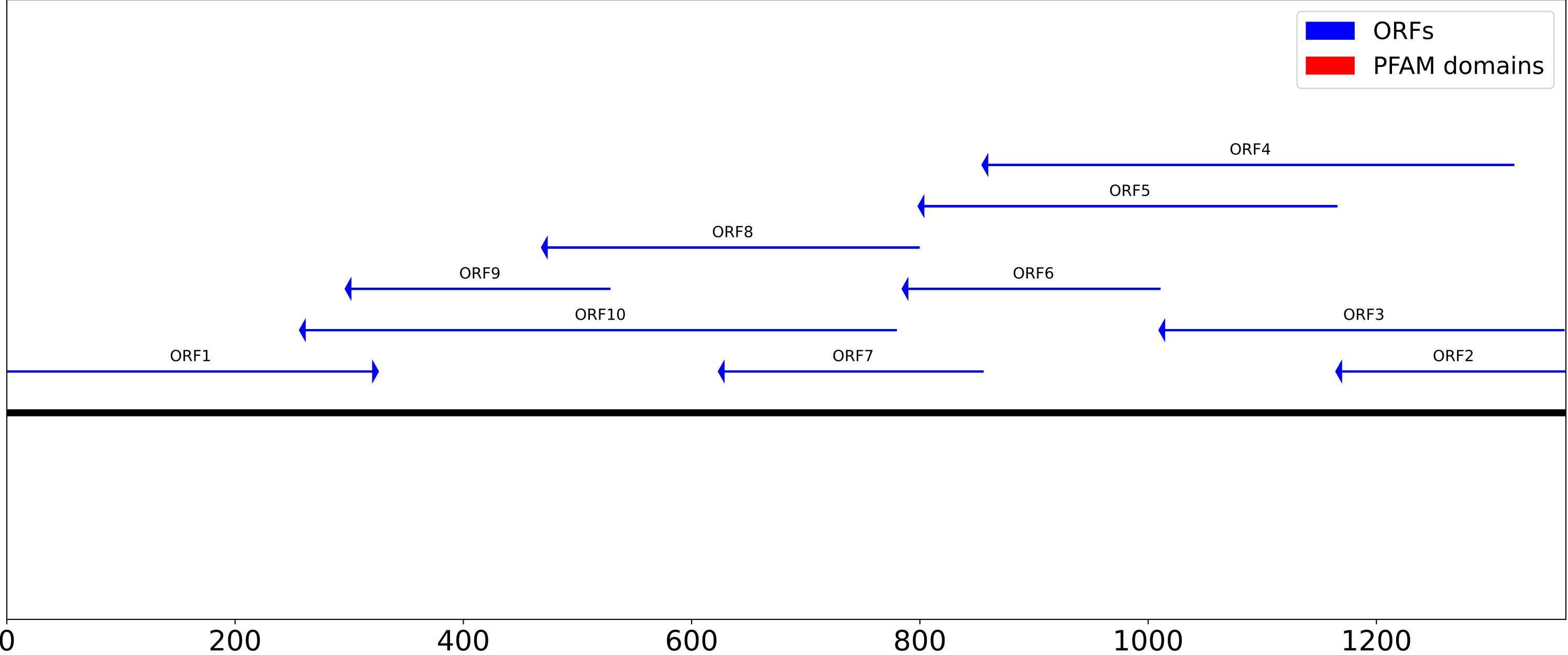
Before TEtrimmer 1366 bp



After TETrimmer ORF and PFAM domain plot



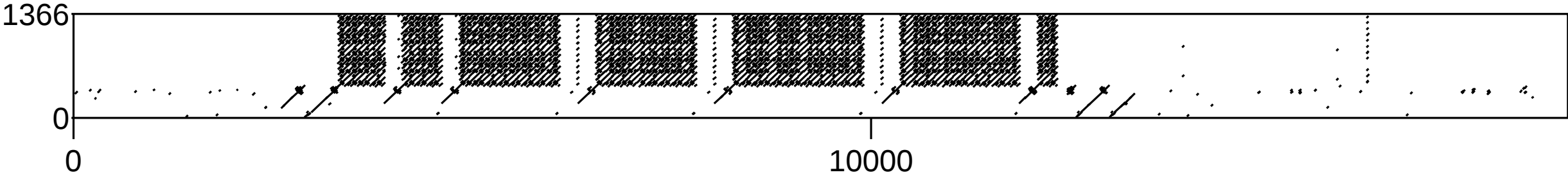
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)