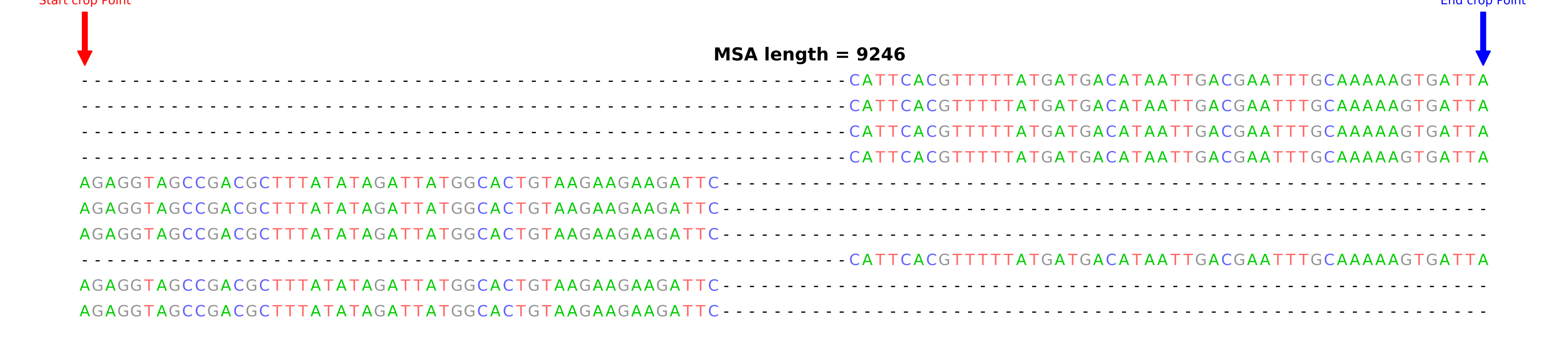
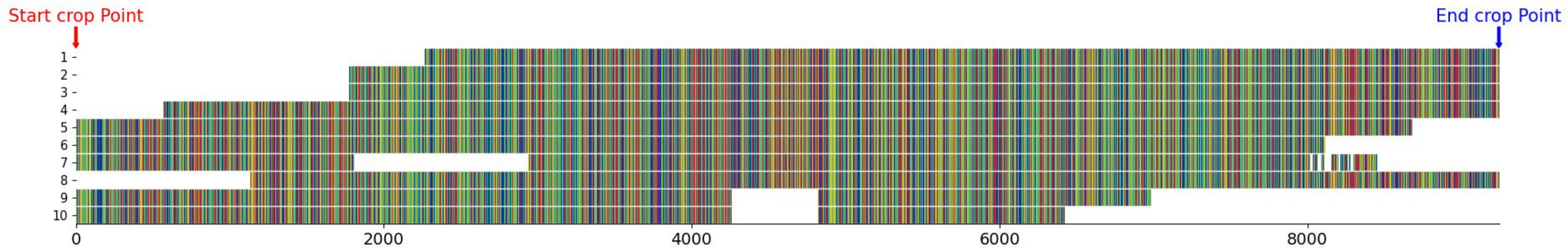
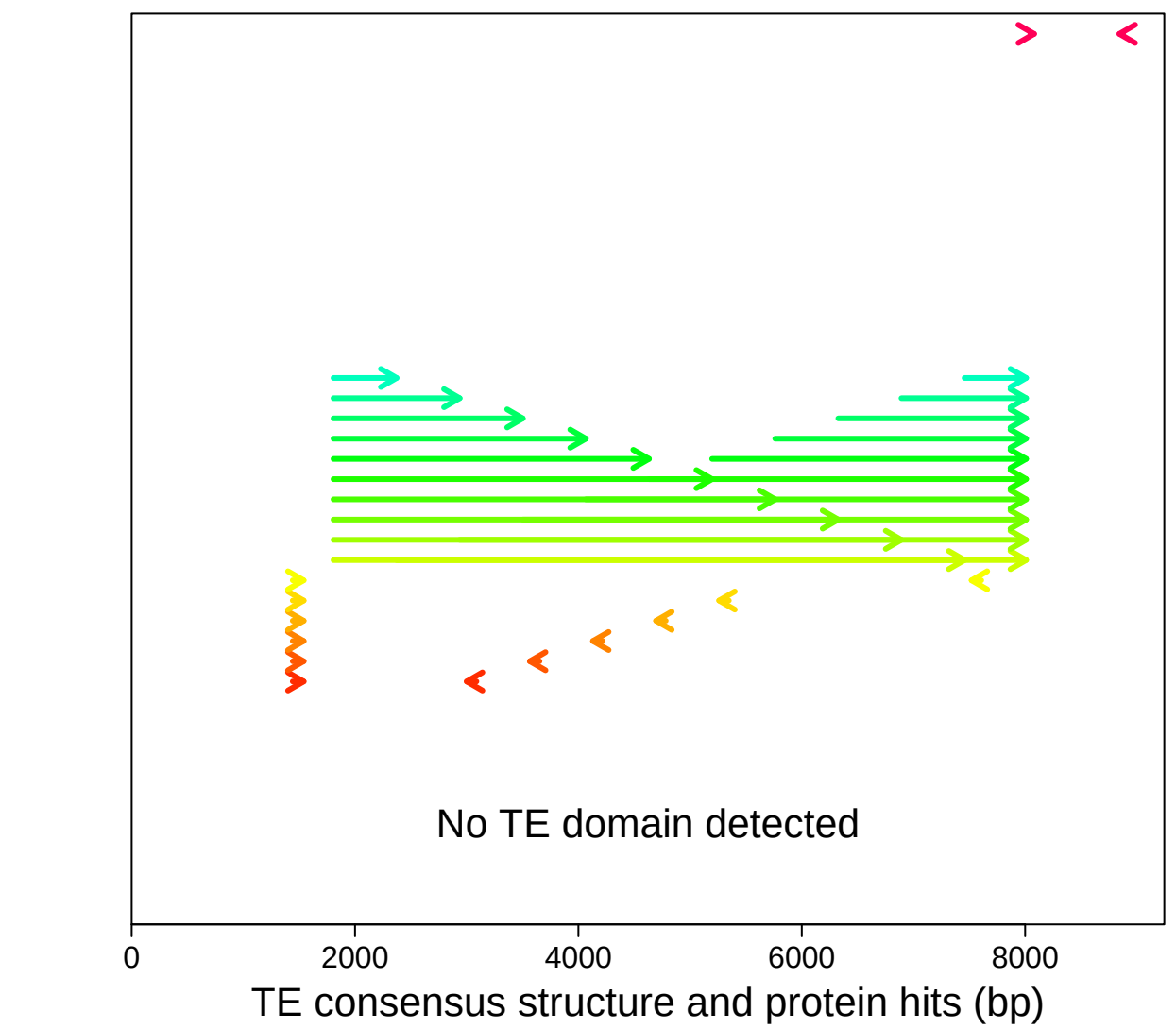
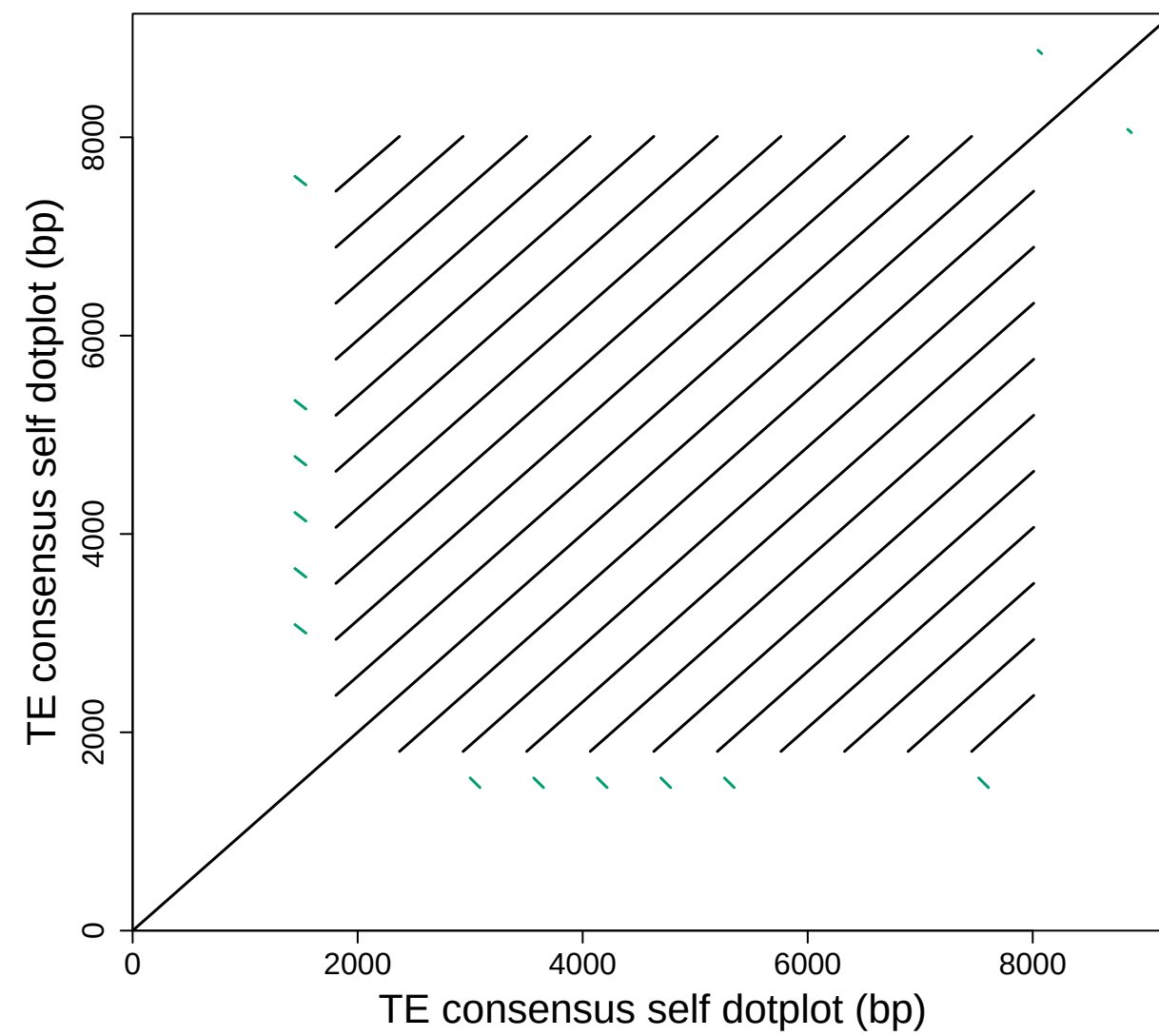
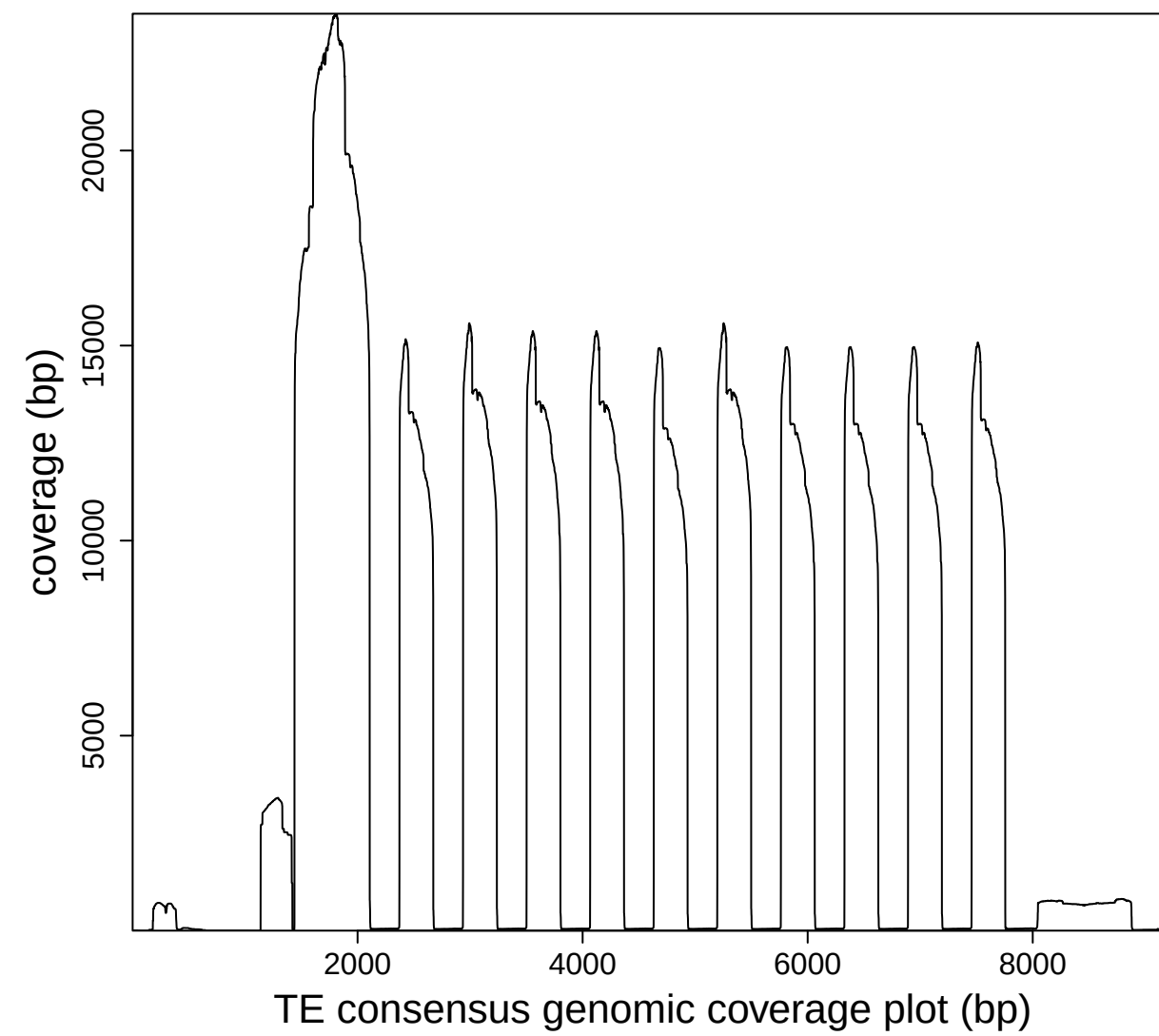
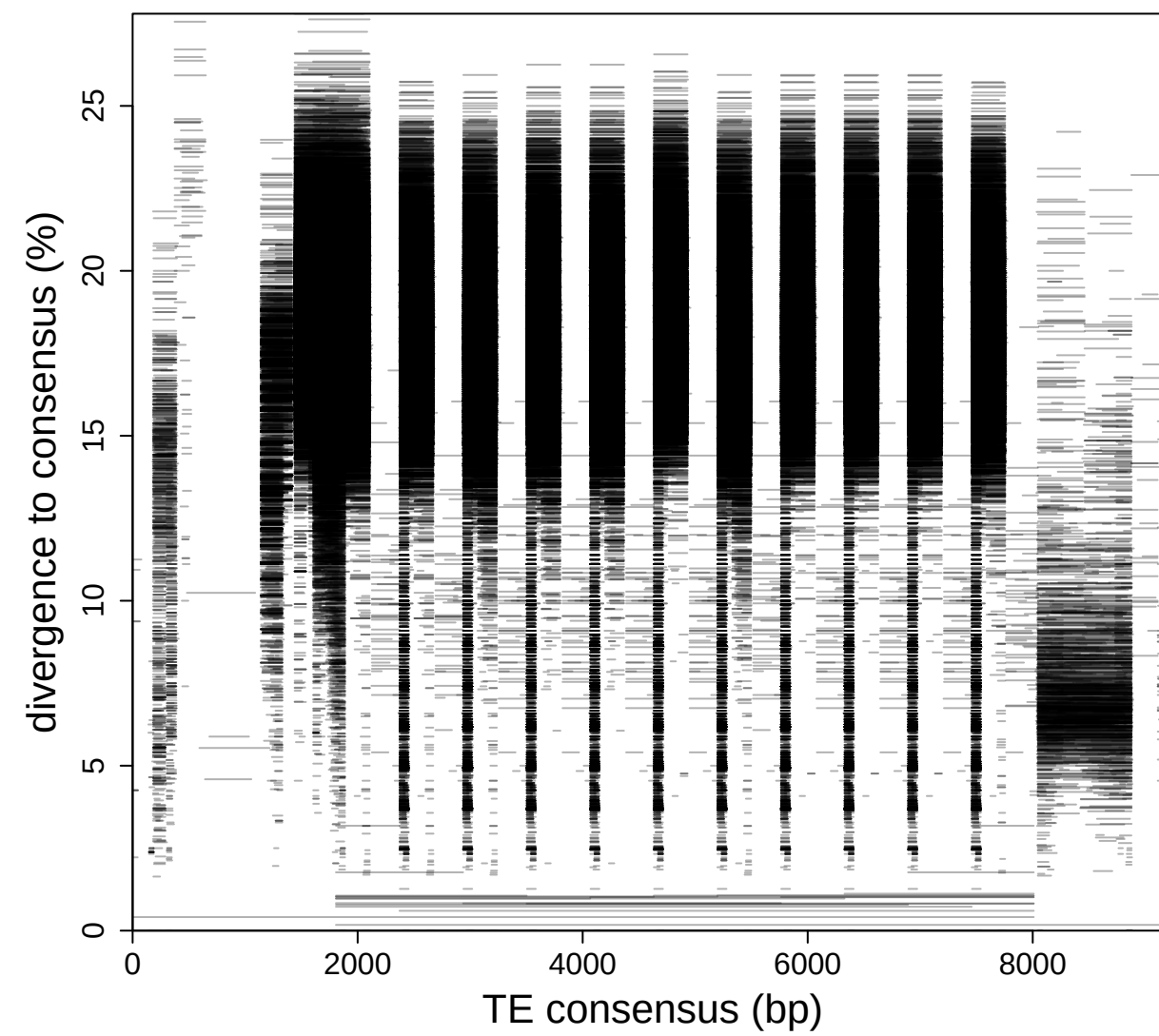






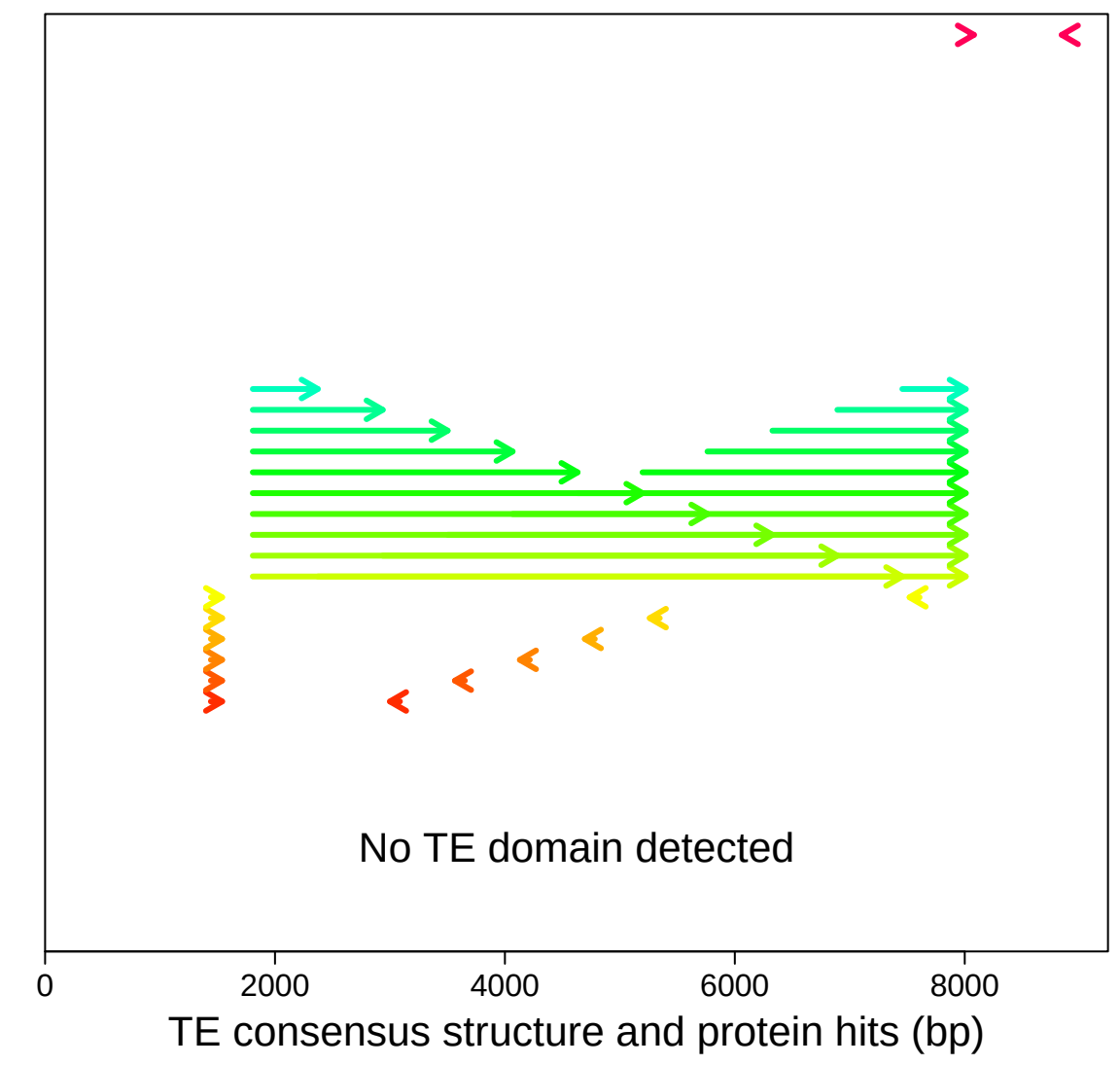
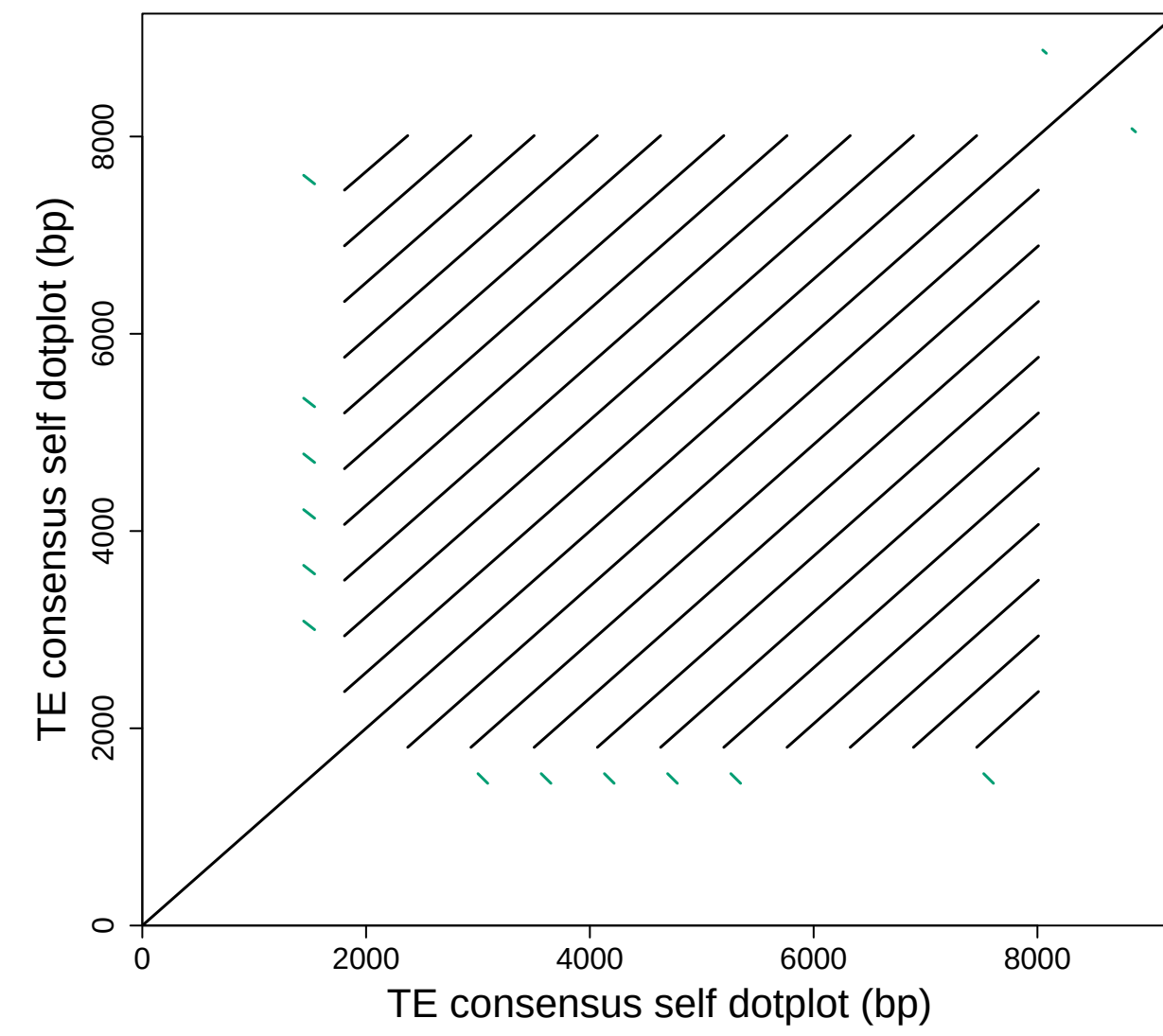
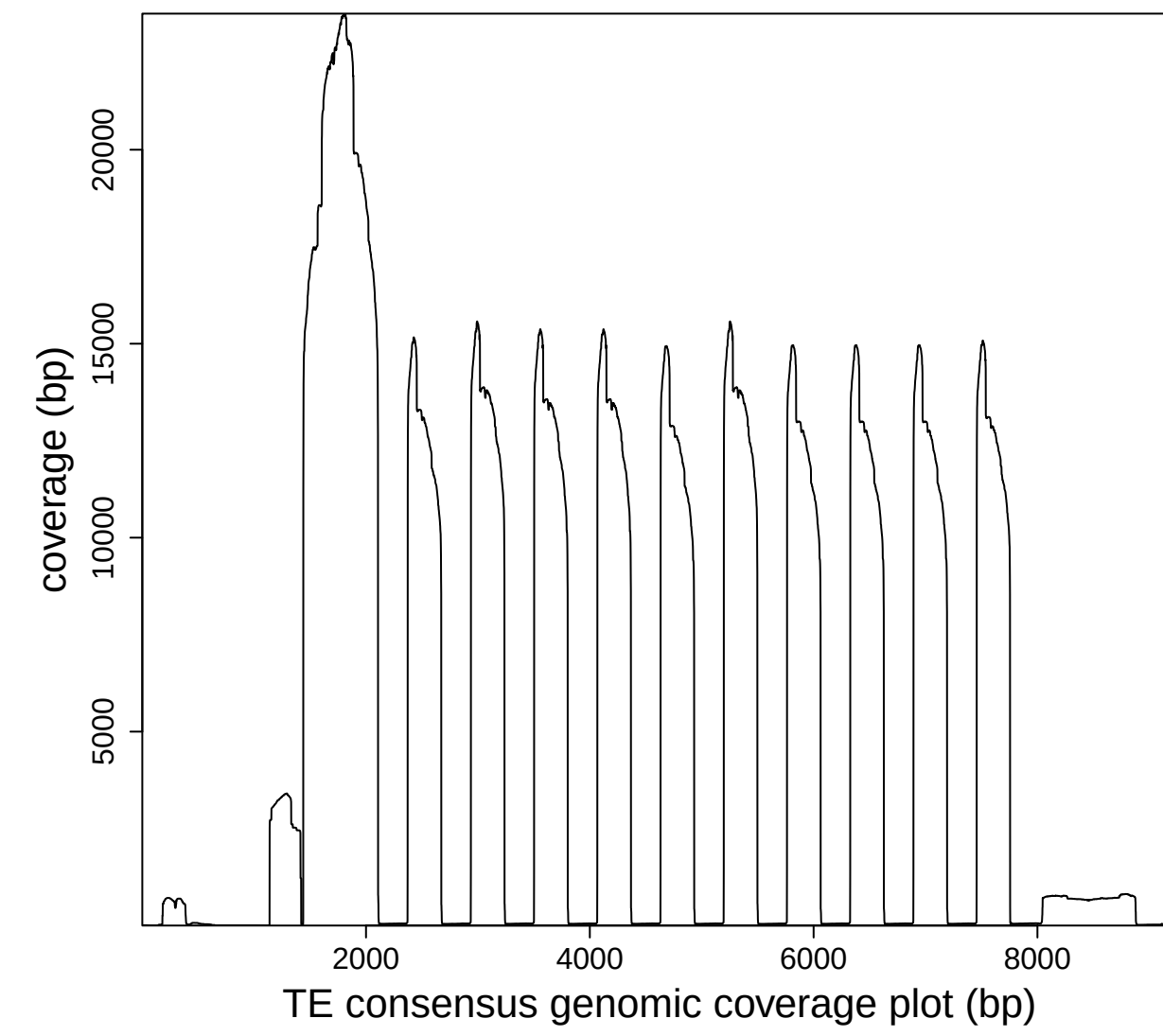
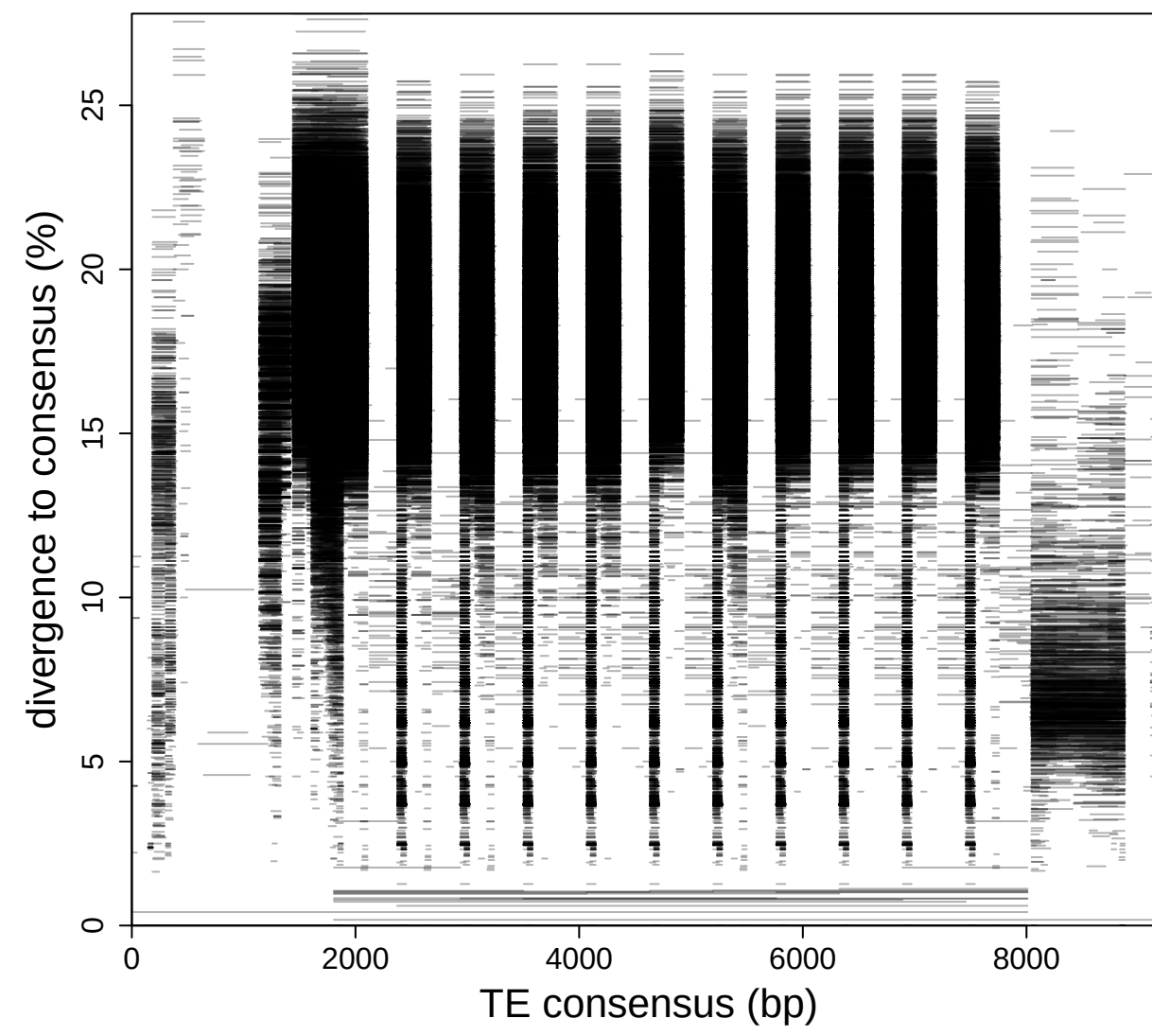
a.b.bed_uf.bed_g_3.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.f
size: 9246bp; fragments: 190349; full length: 0 (>=8321.4bp)

After TEtrimmer 9246 bp

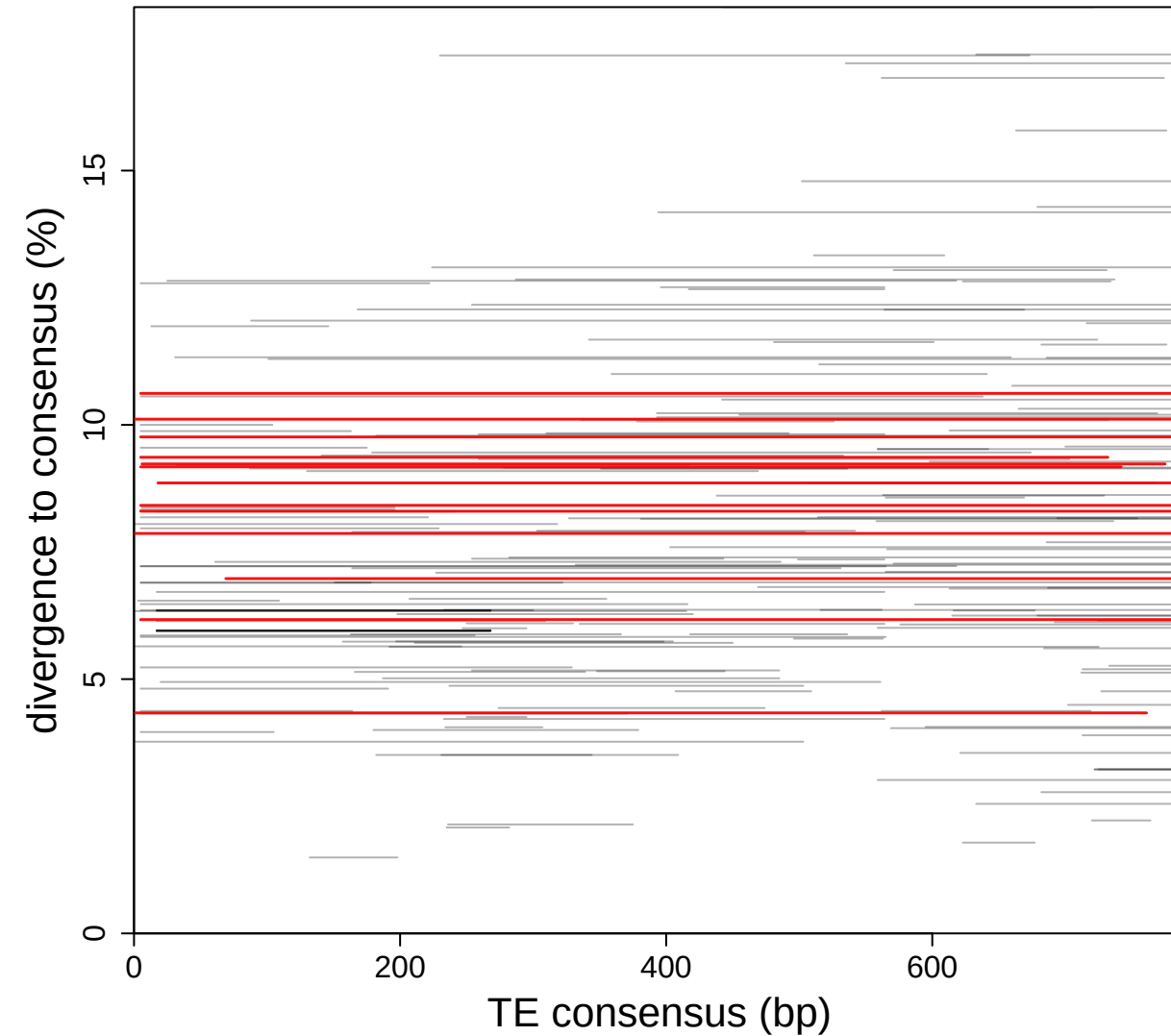


7.fasta.b.bed uf.bed g_3.bed fm_1.bed 0 0 bcln.fa aln.fa cl.f
size: 9246bp; fragments: 190349; full length: 0 (>=8321.4bp)

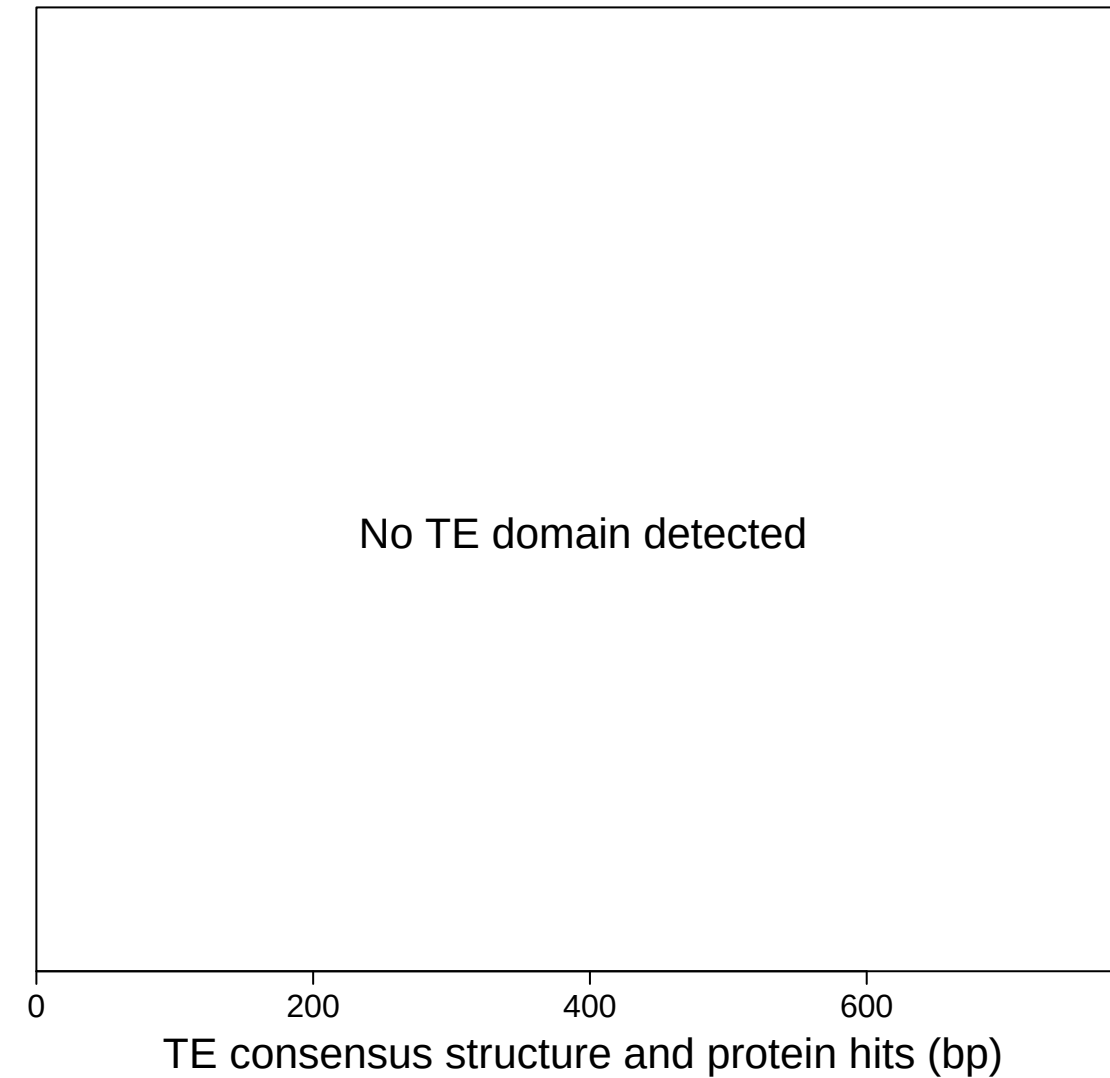
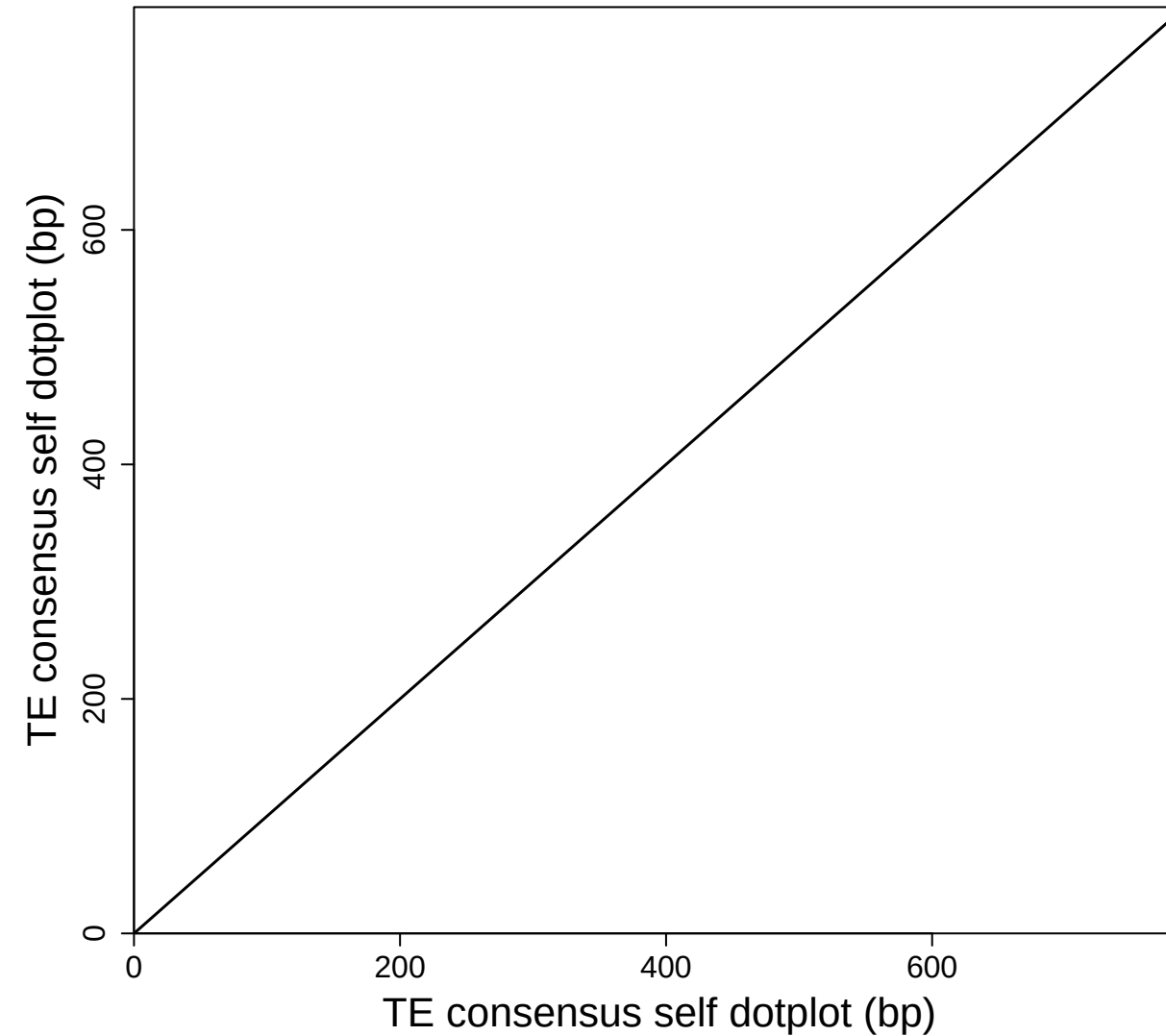
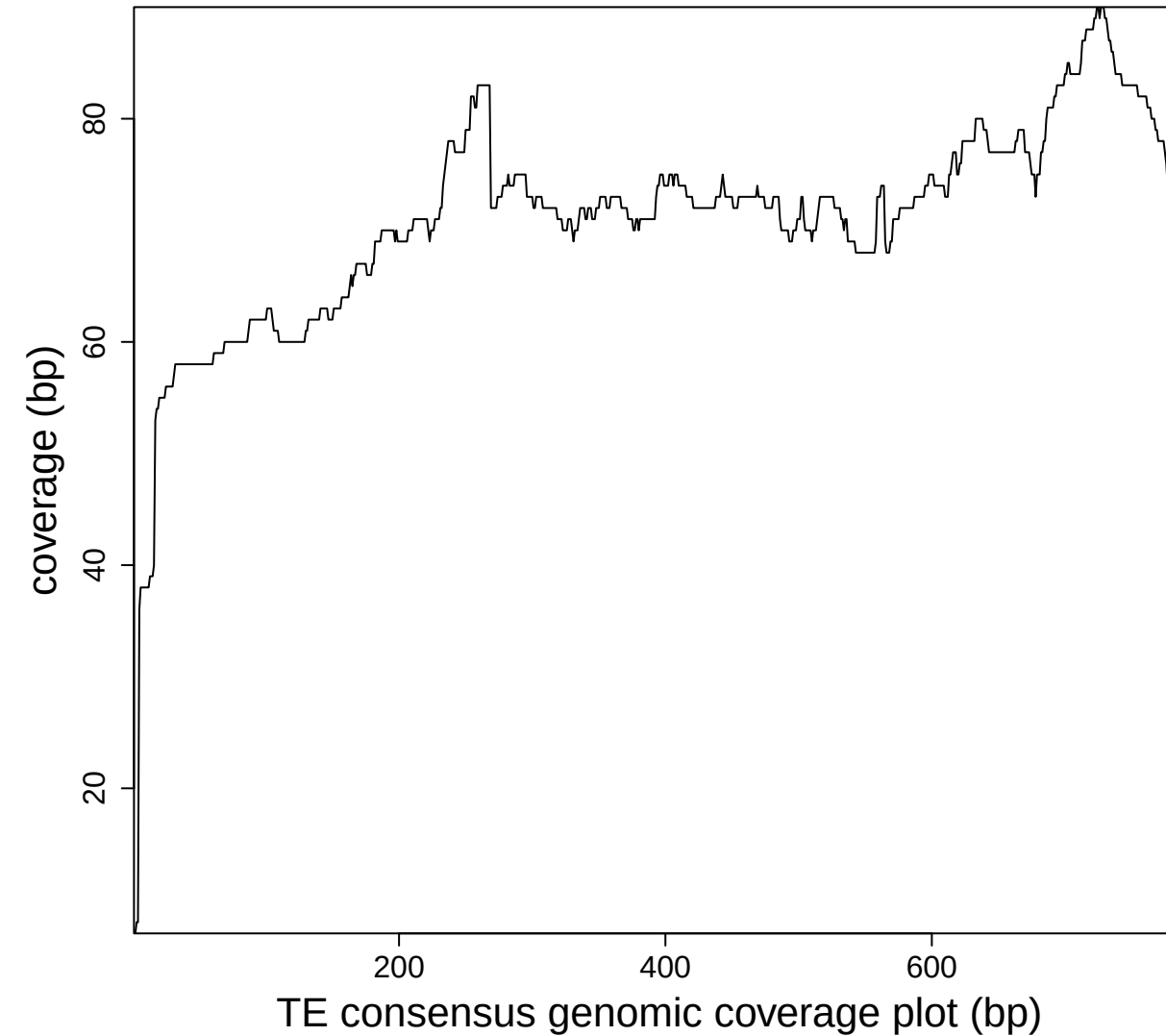
After TEtrimmer Extended plot Blue lines are boundaries



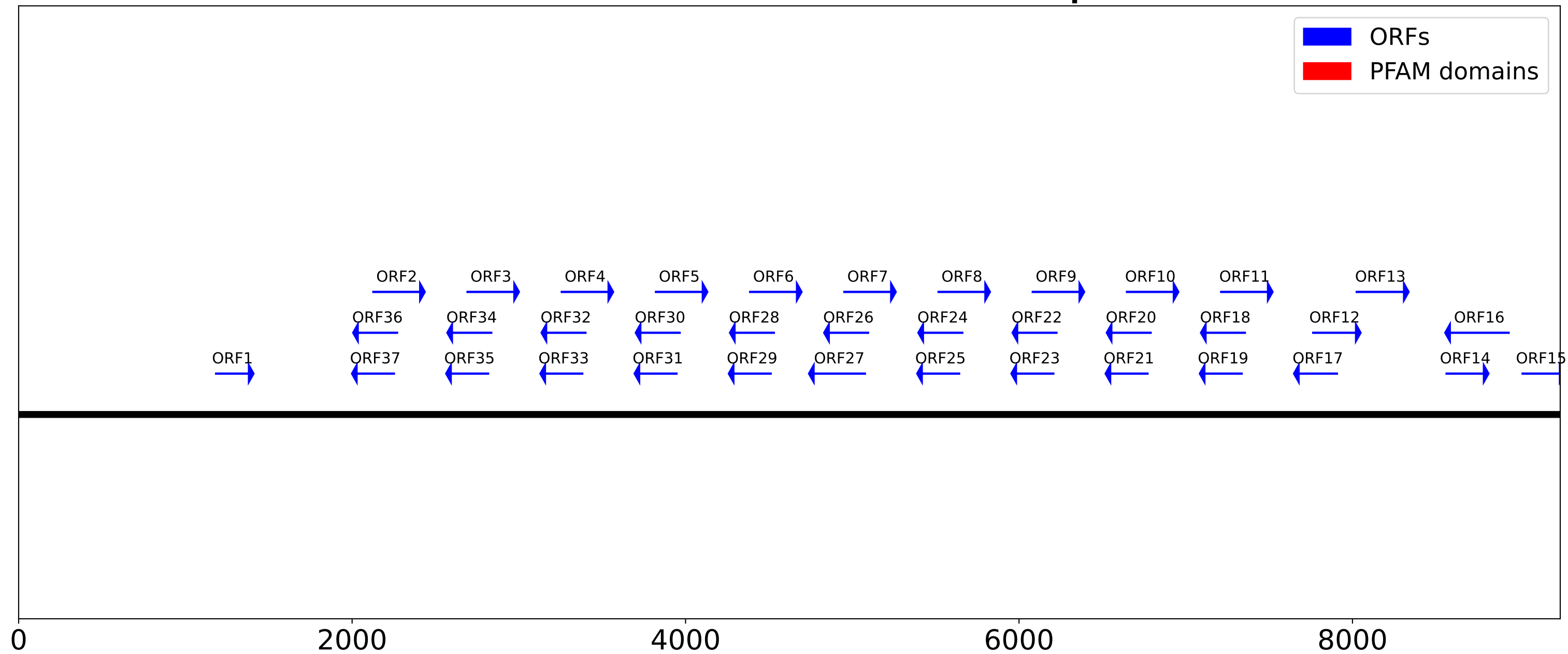
TE: rnd_5_family_8967
size: 790bp; fragments: 194; full length: 13 (≥ 711 bp)



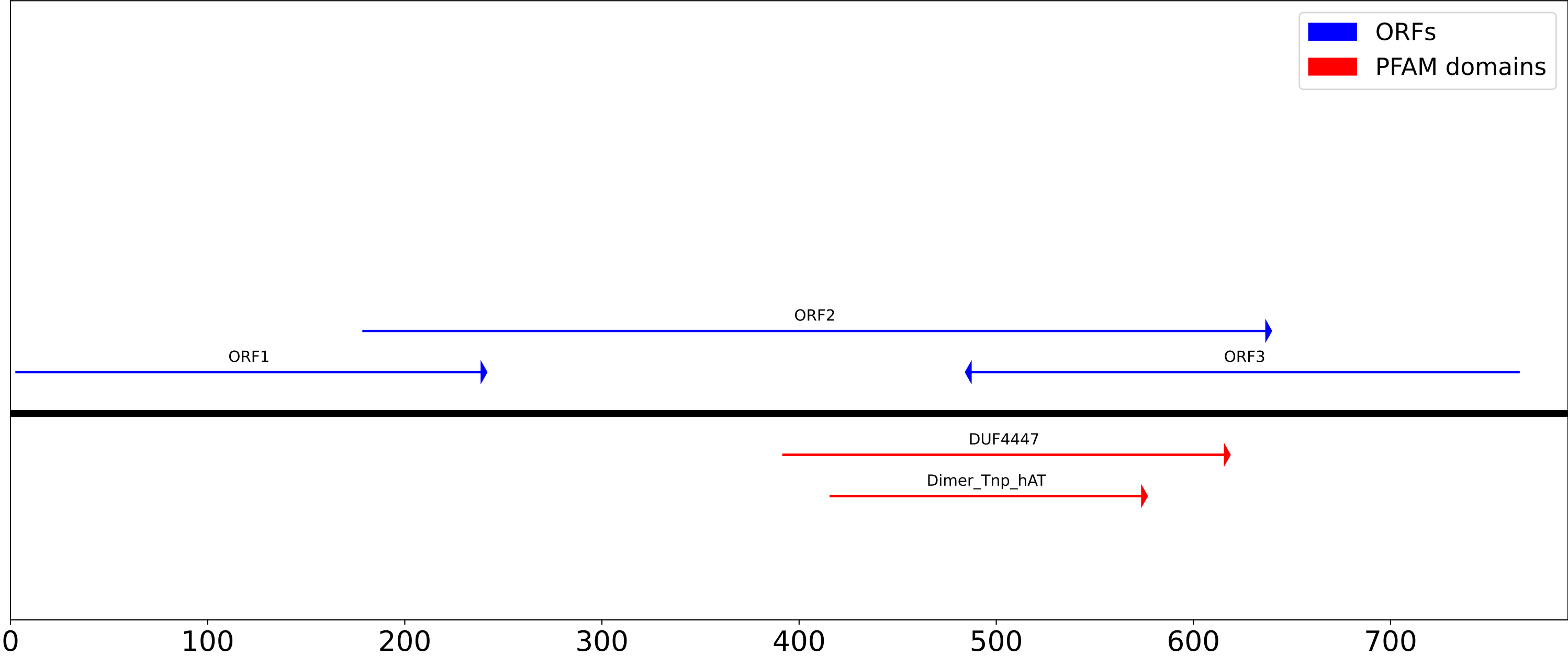
Before TEtrimmer 790 bp



After TEtrimmer ORF and PFAM domain plot



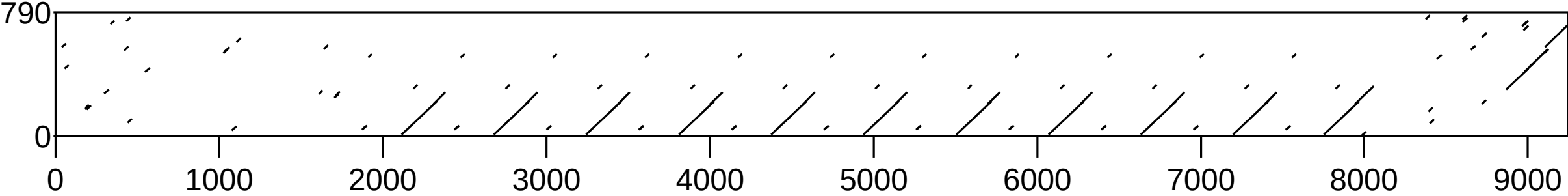
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

ensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)