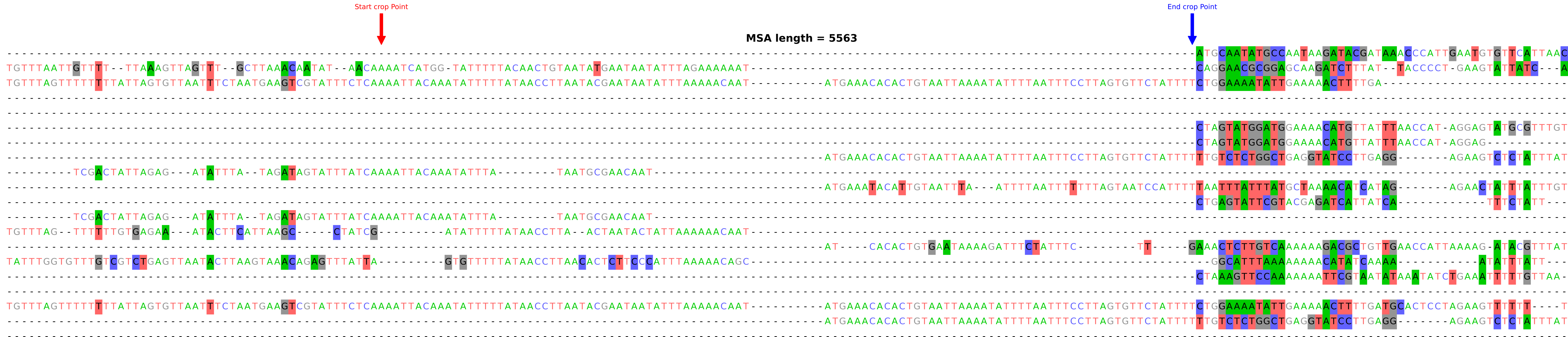
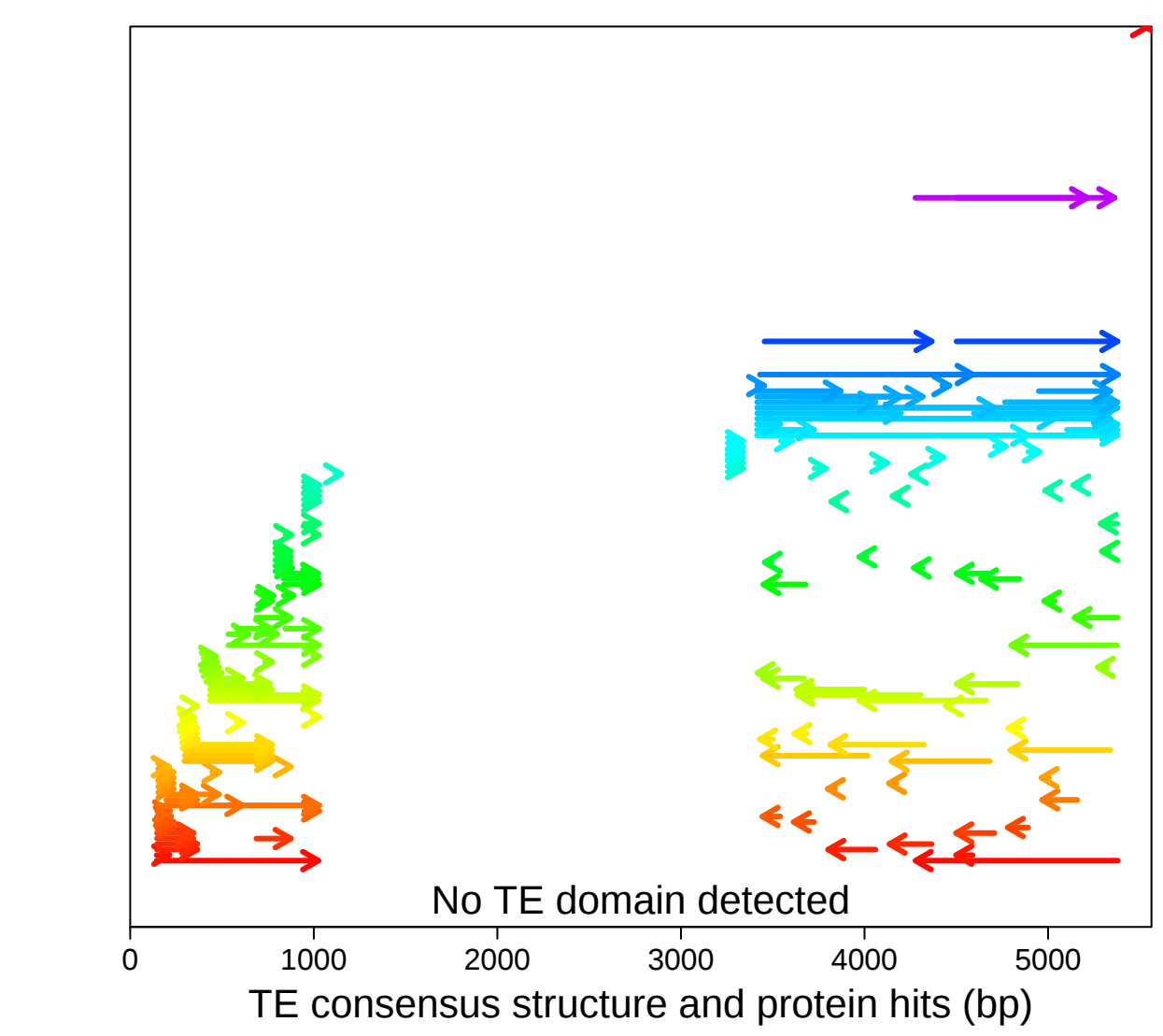
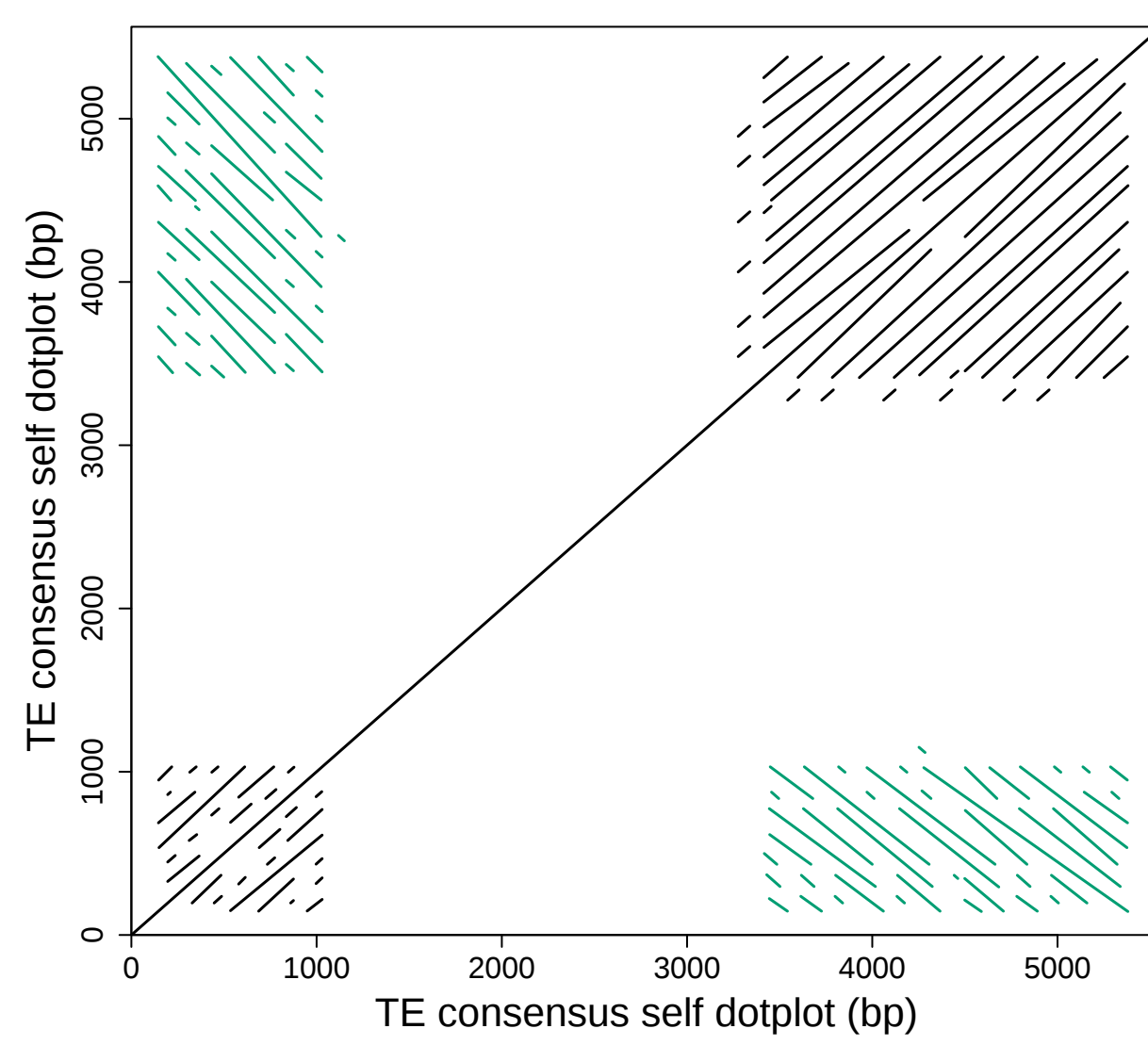
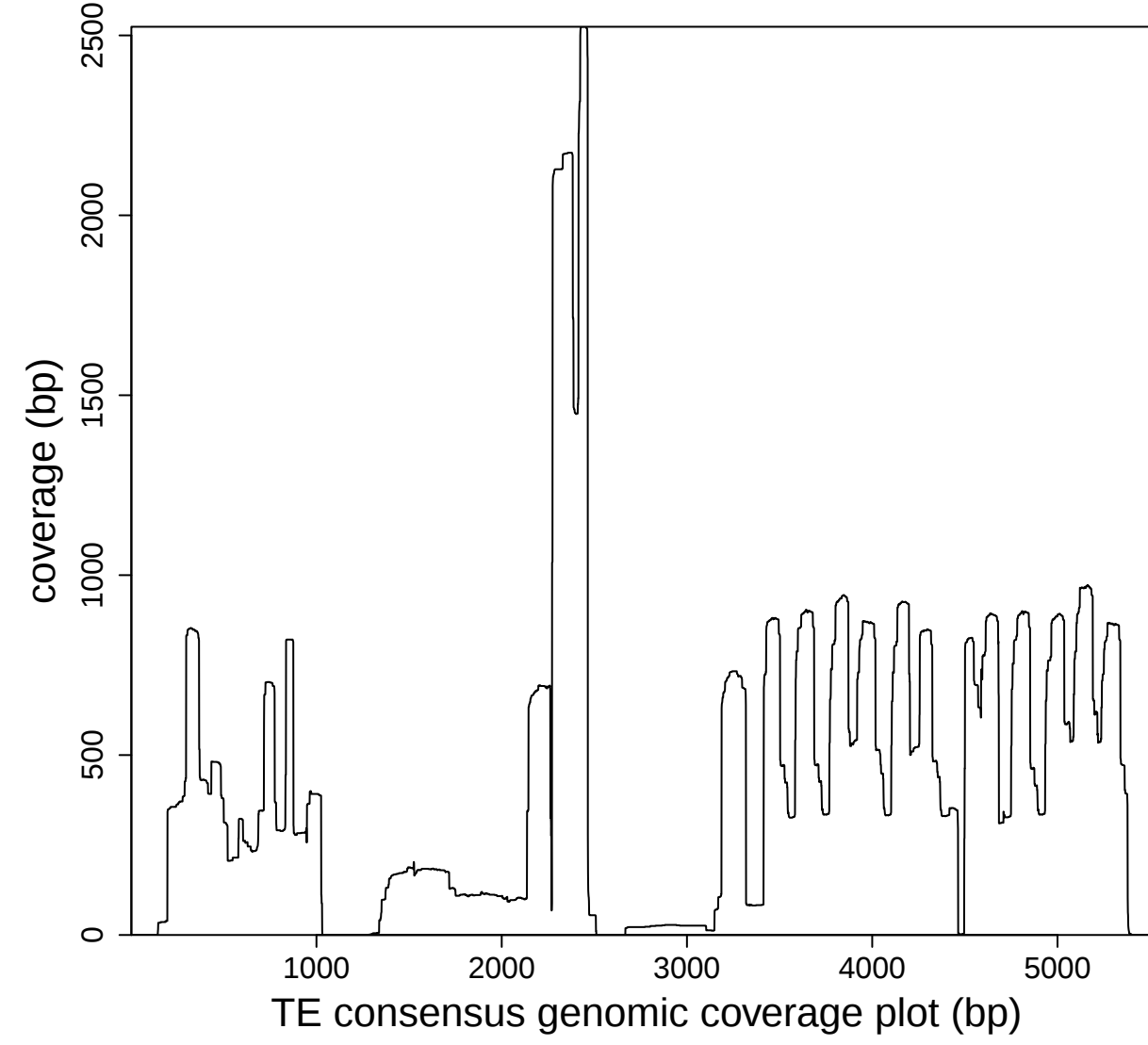
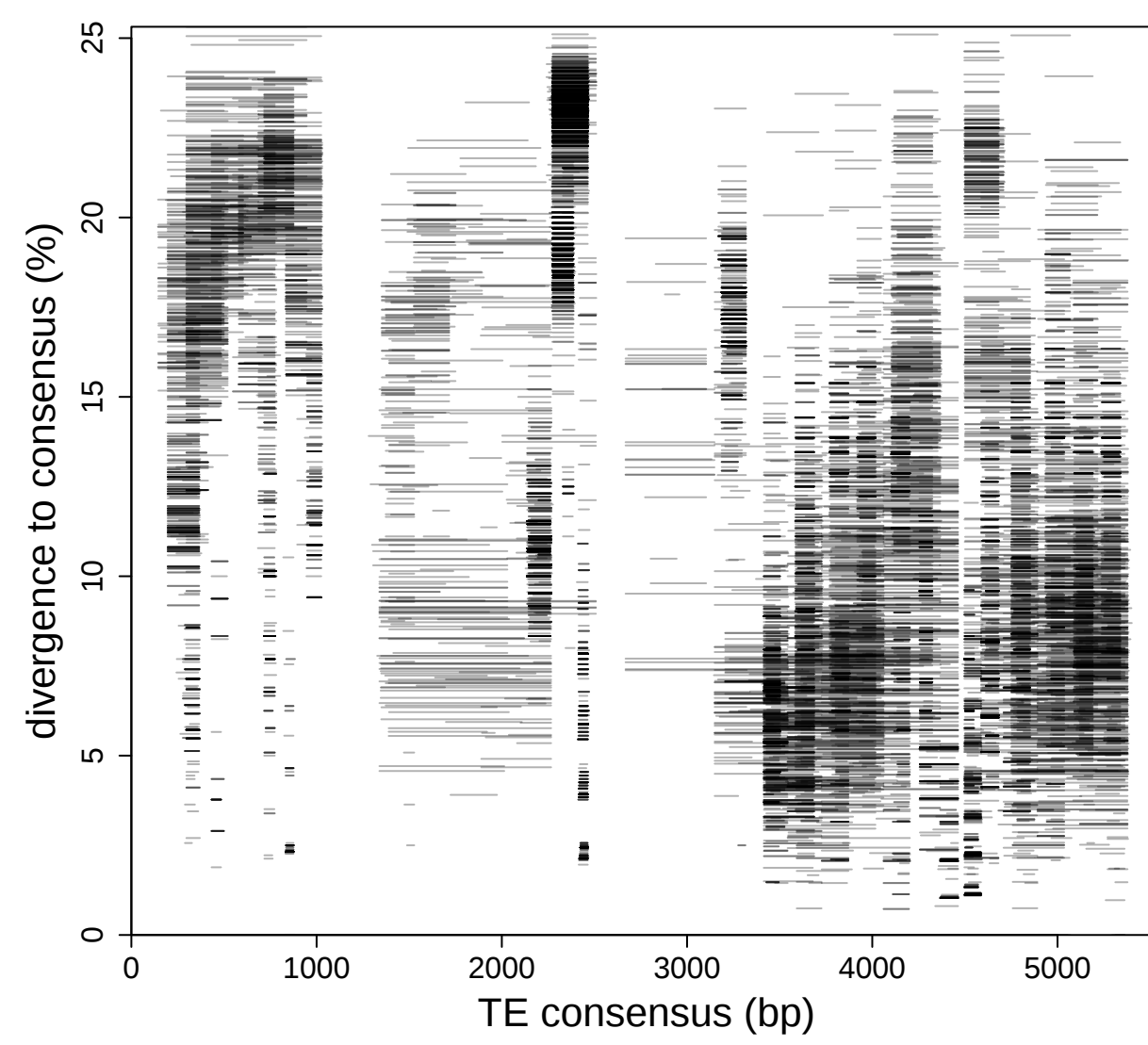






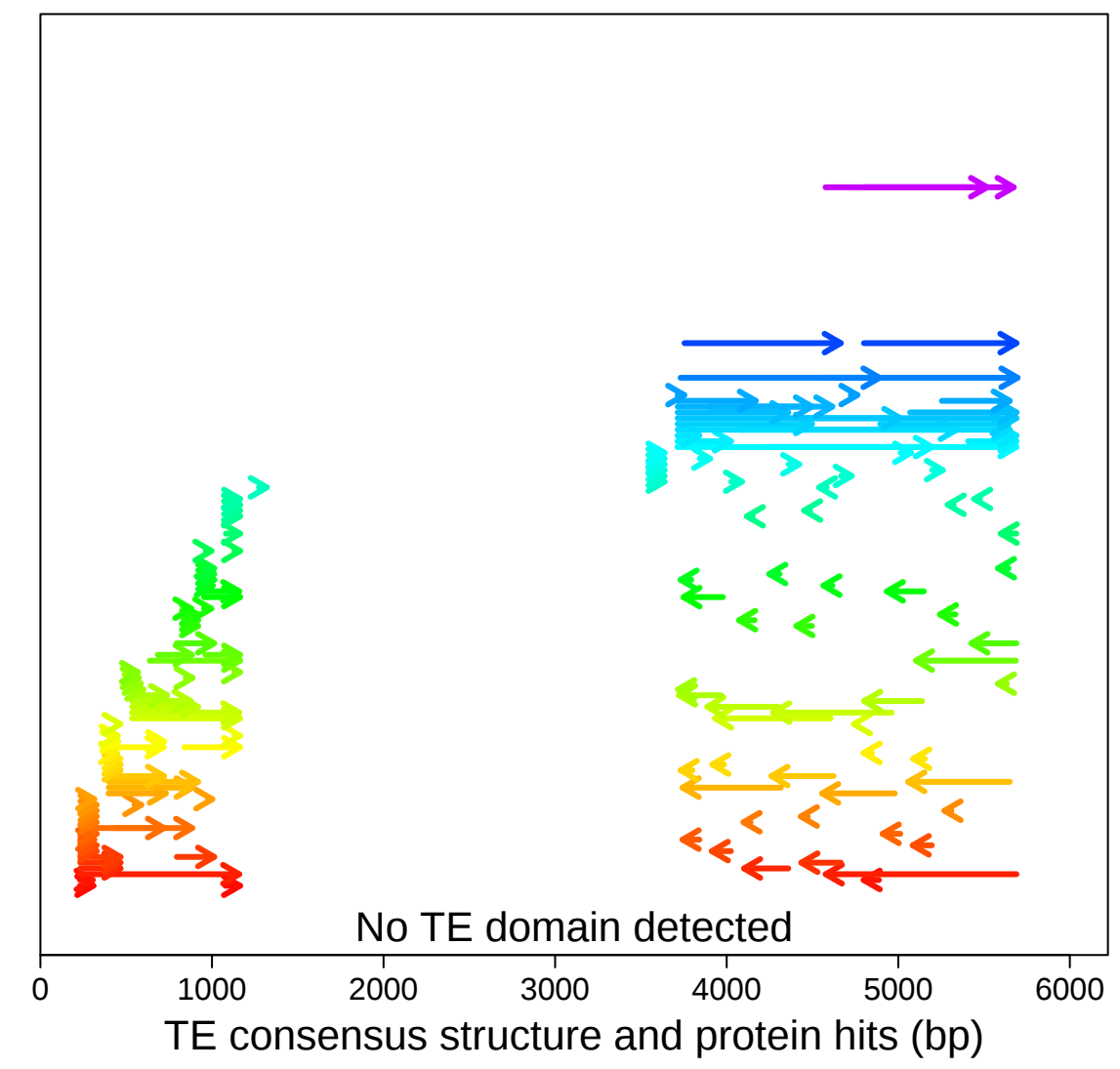
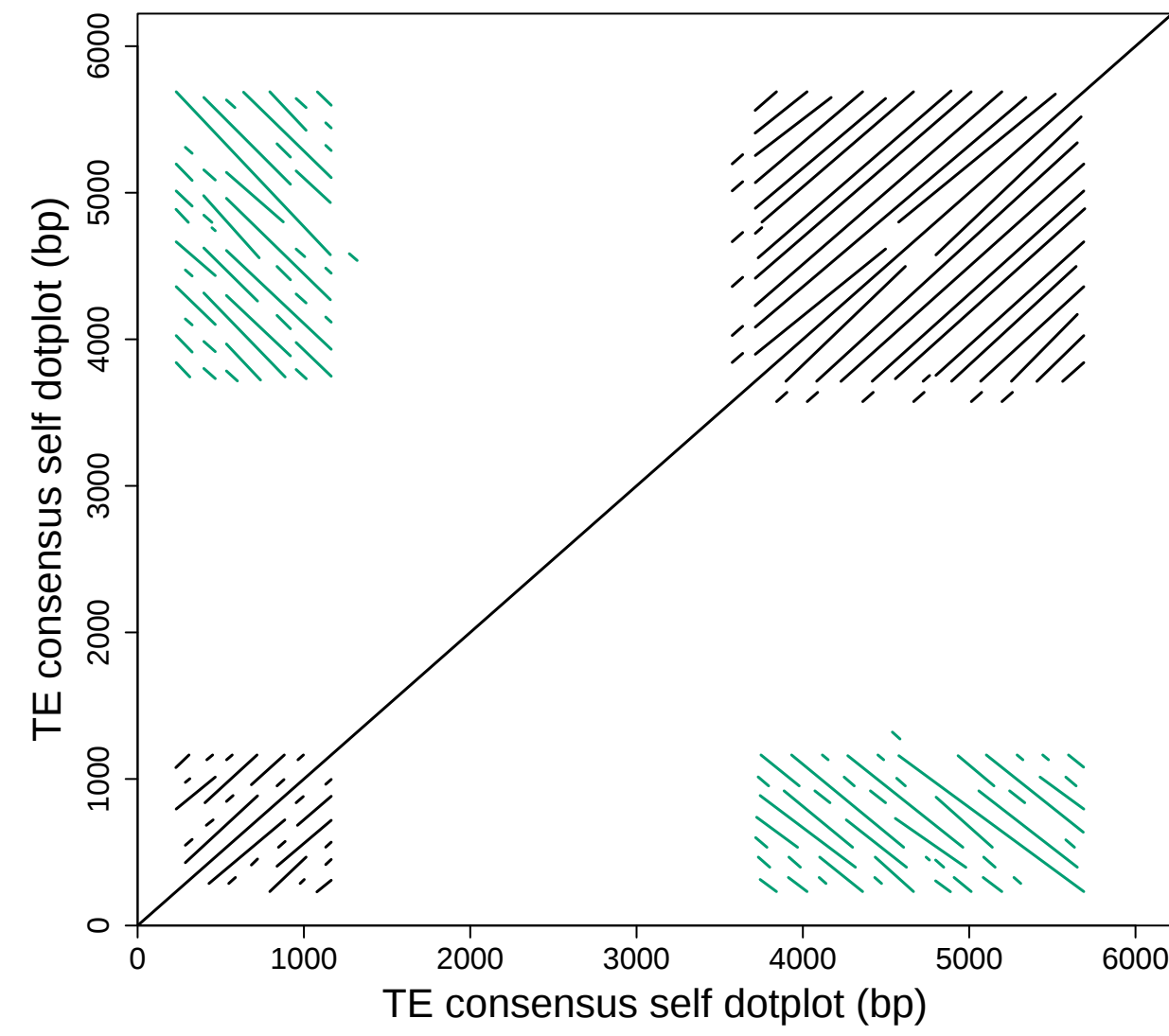
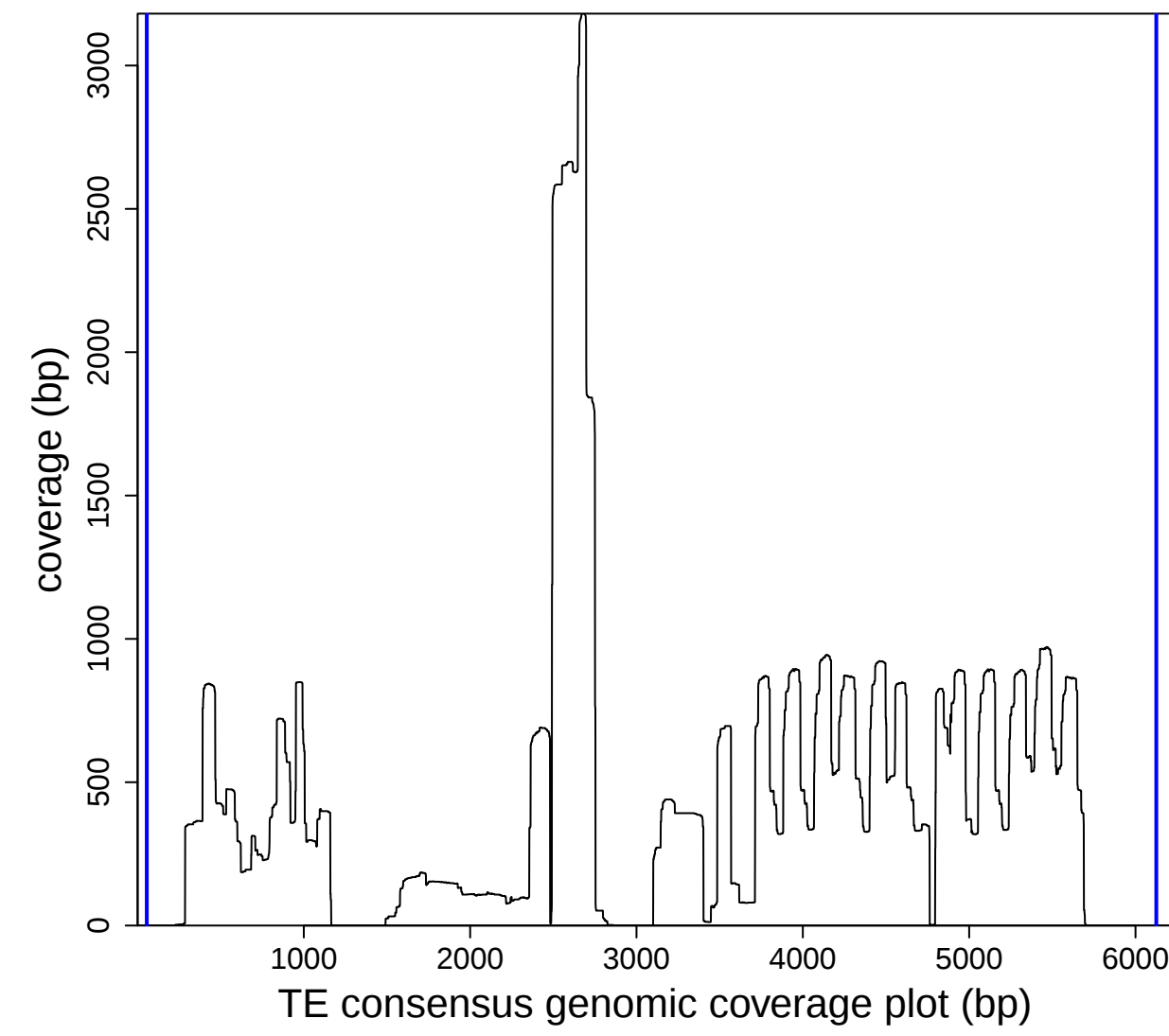
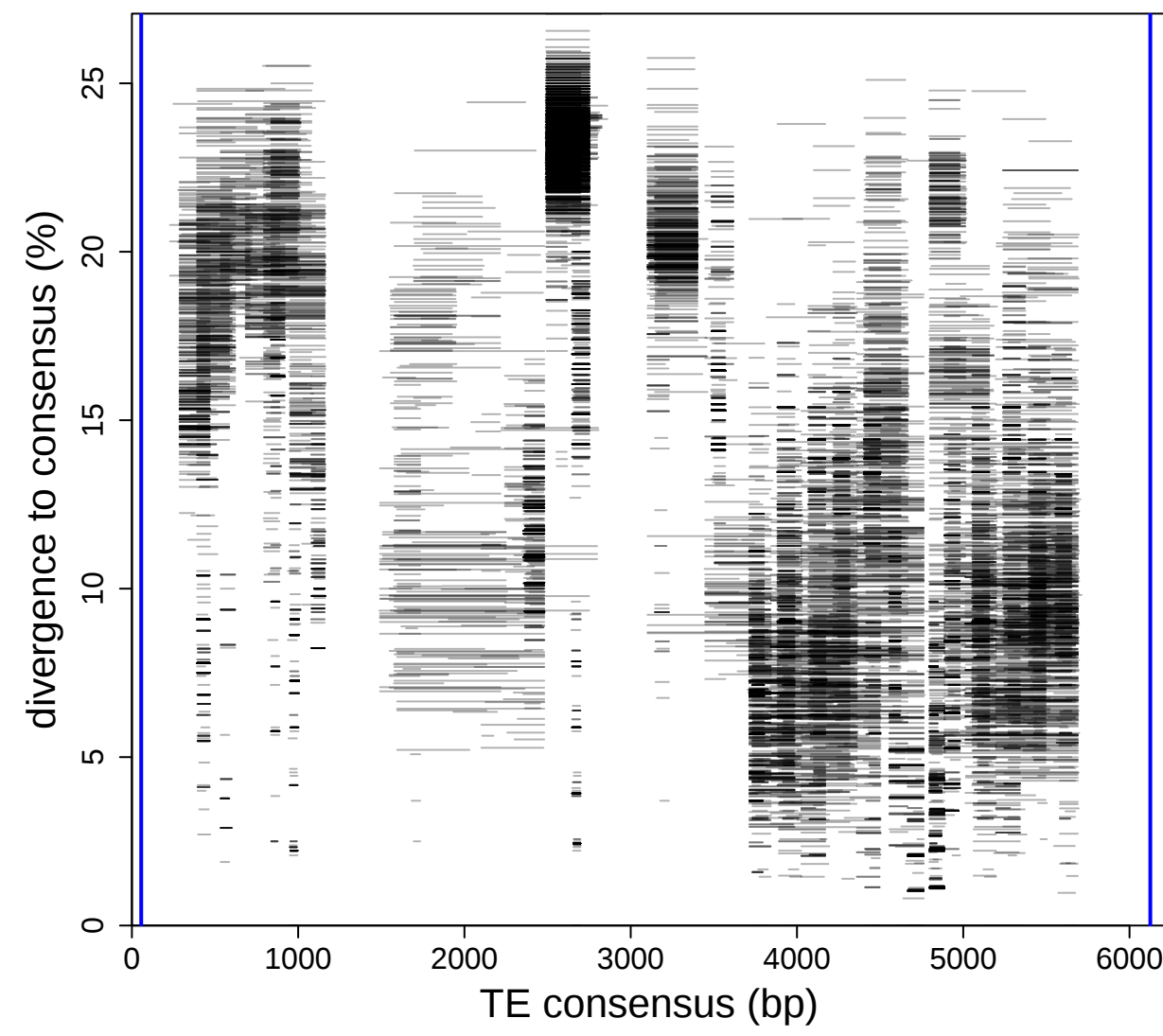
l.b.bed_uf.bed_g_2.bed_fm_1.bed_0_0_bcIn.fa_aln.fa_cl.fa_gs.fa
size: 5563bp; fragments: 14524; full length: 0 (>=5006.7bp)

After TEtrimmer 5563 bp



7.fasta.b.bed_uf.bed_g_2.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa
size: 6222bp; fragments: 14802; full length: 0 (>=5599.8bp)

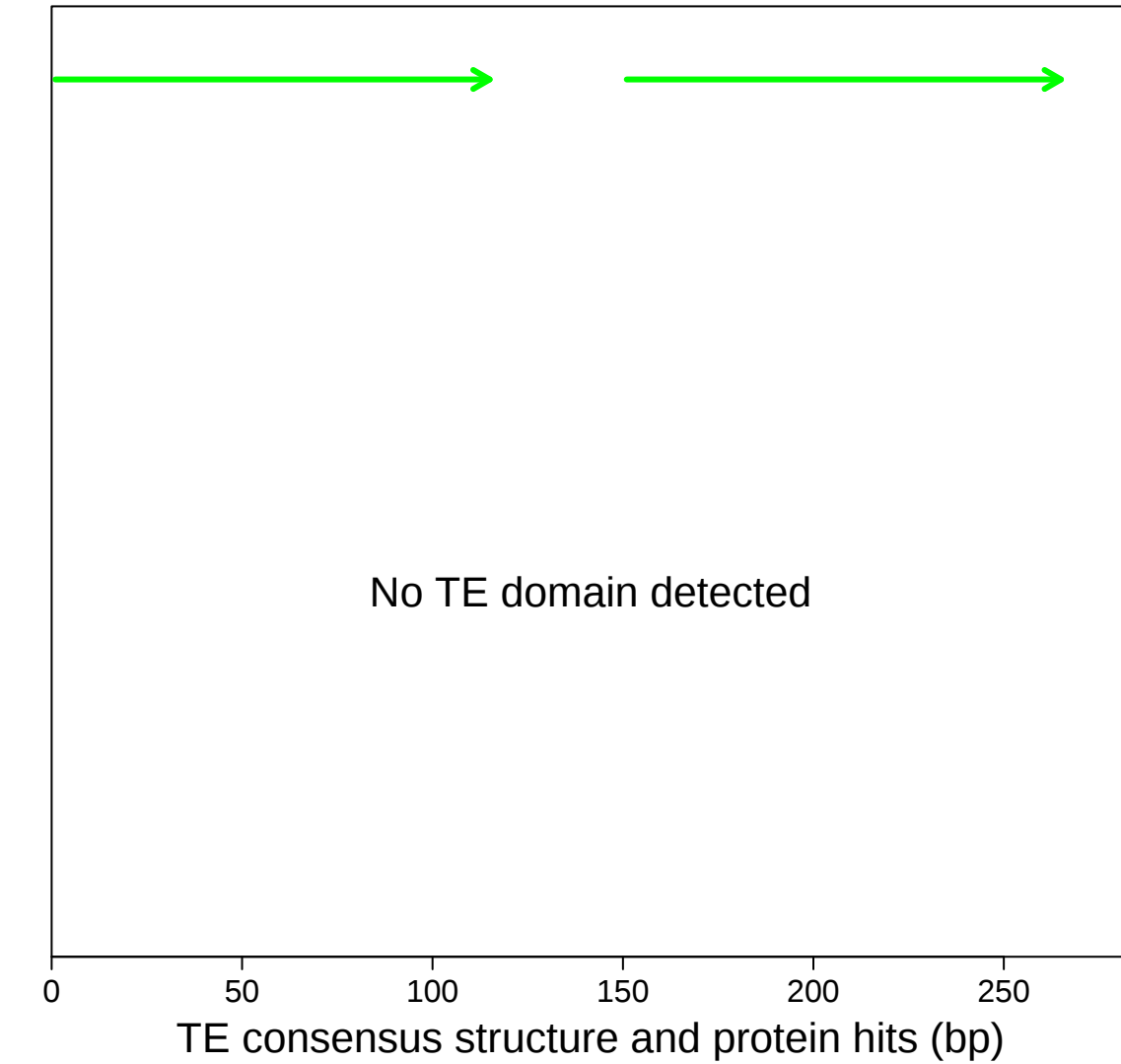
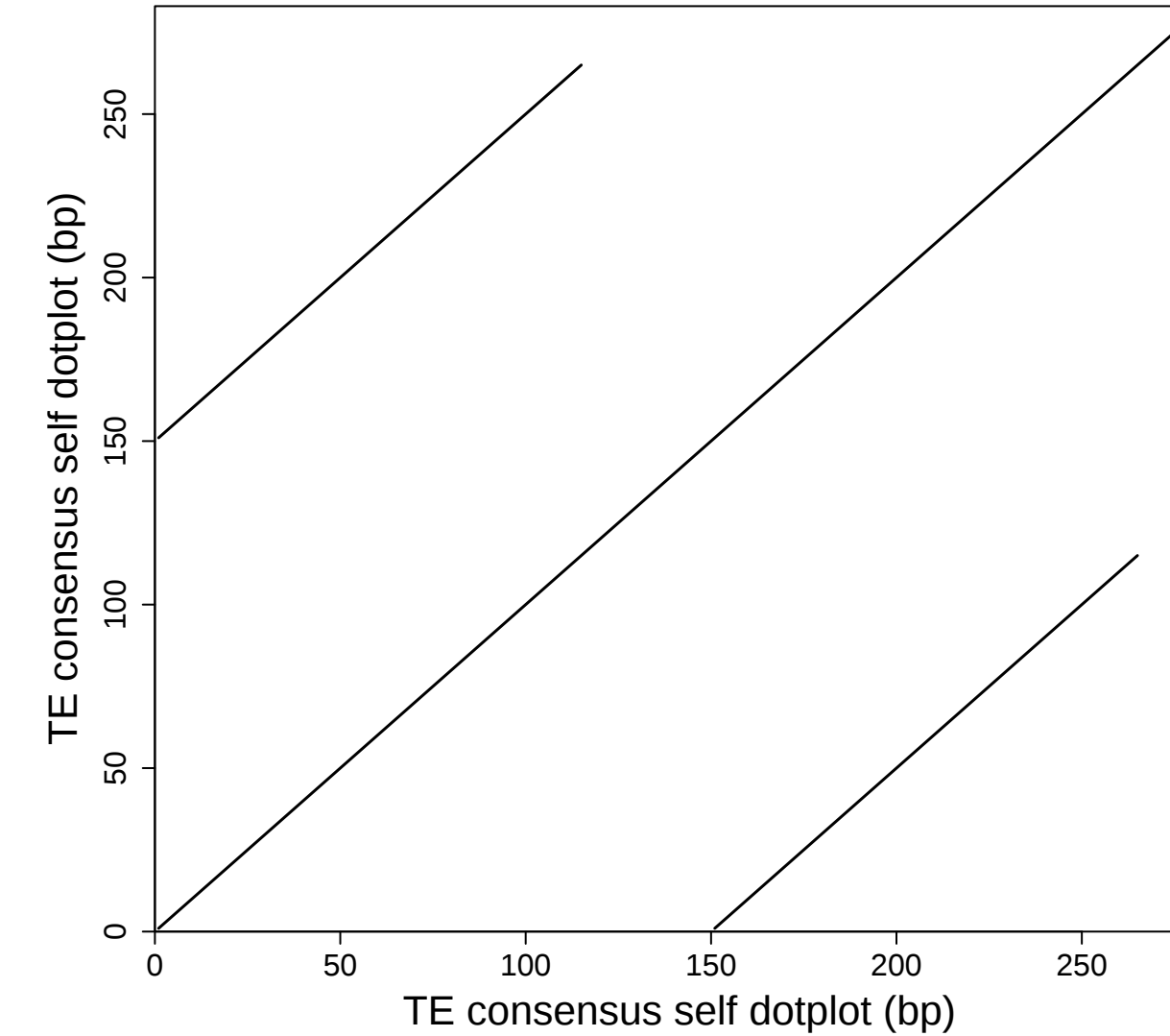
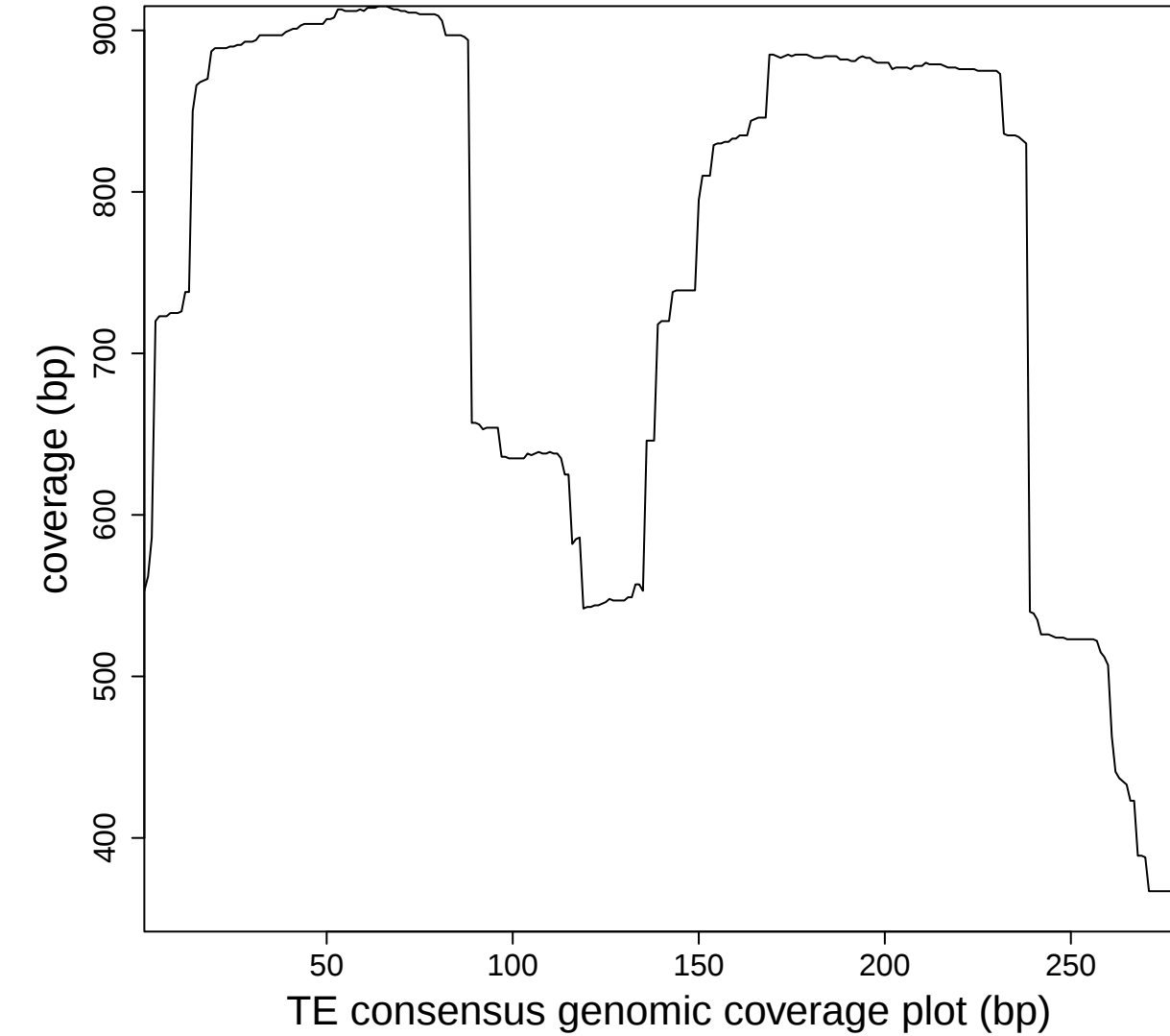
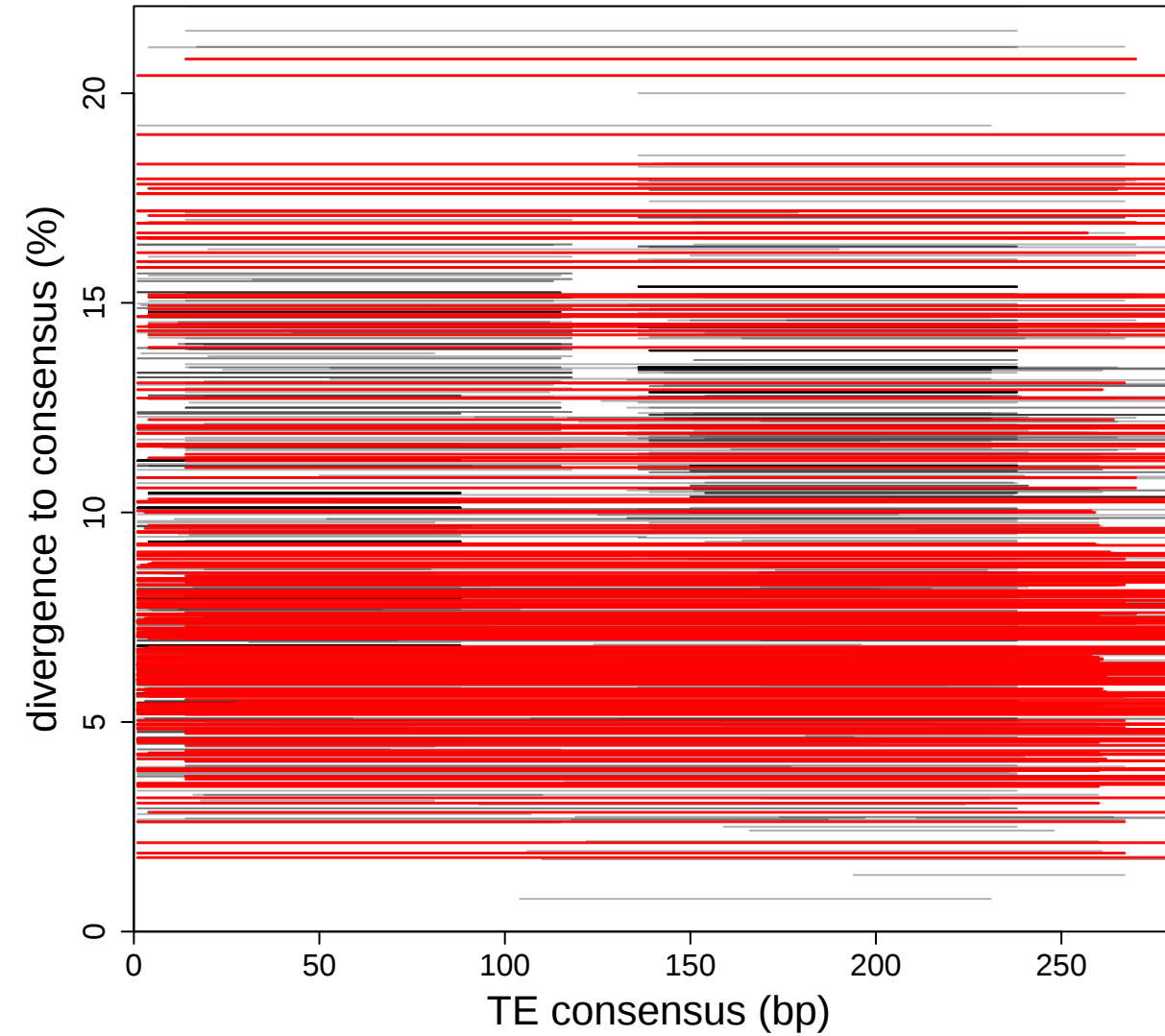
After TEtrimmer Extended plot Blue lines are boundaries



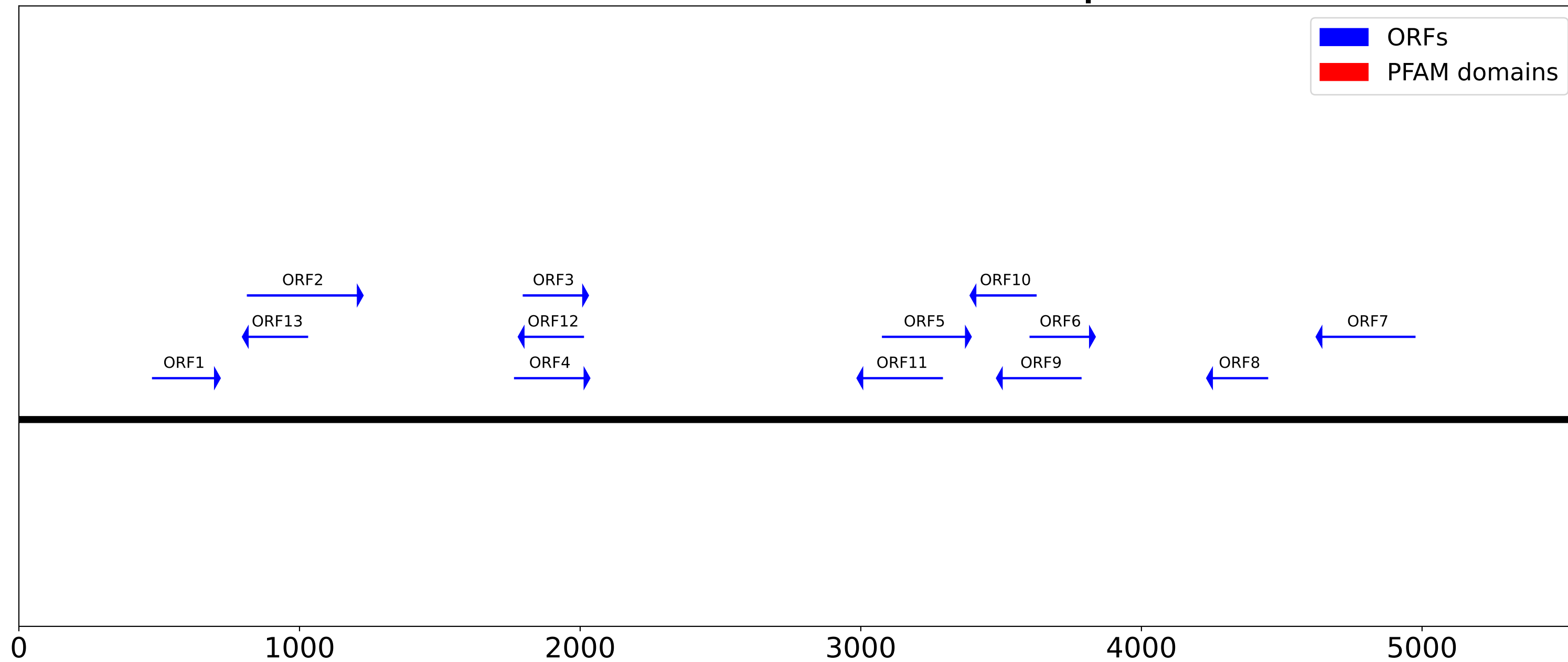
Before TEtrimmer 283 bp

TE: rnd_1_family_917

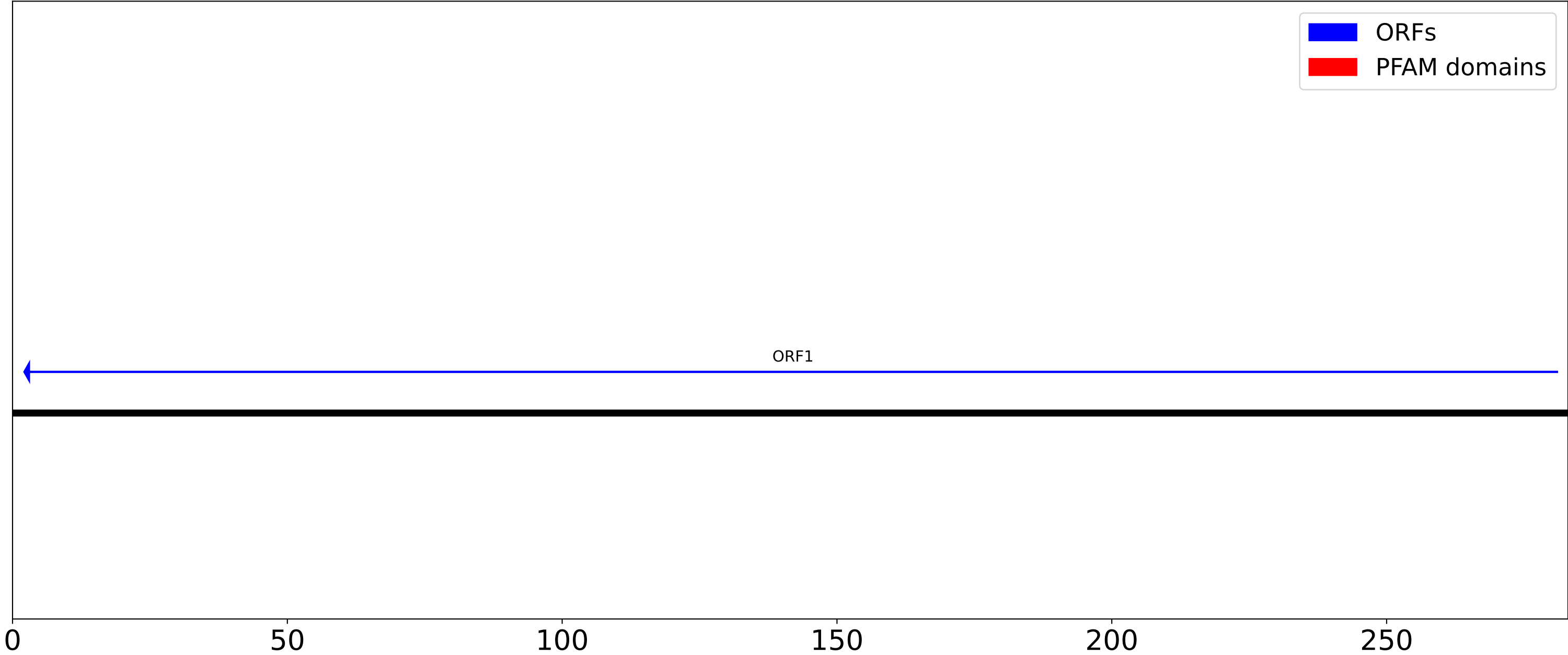
size: 283bp; fragments: 1302; full length: 339 (≥ 254.7 bp)



After TEtrimmer ORF and PFAM domain plot



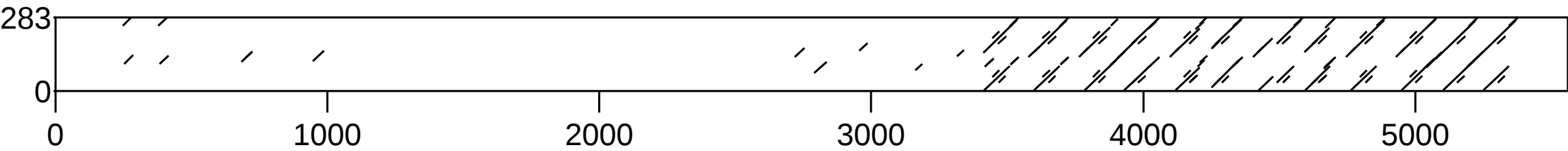
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

Consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)