

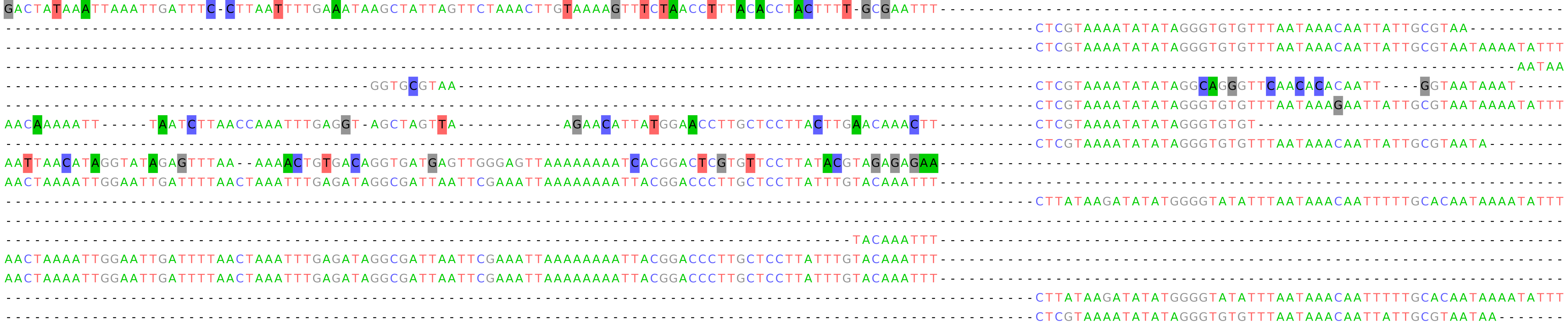
Start crop Point

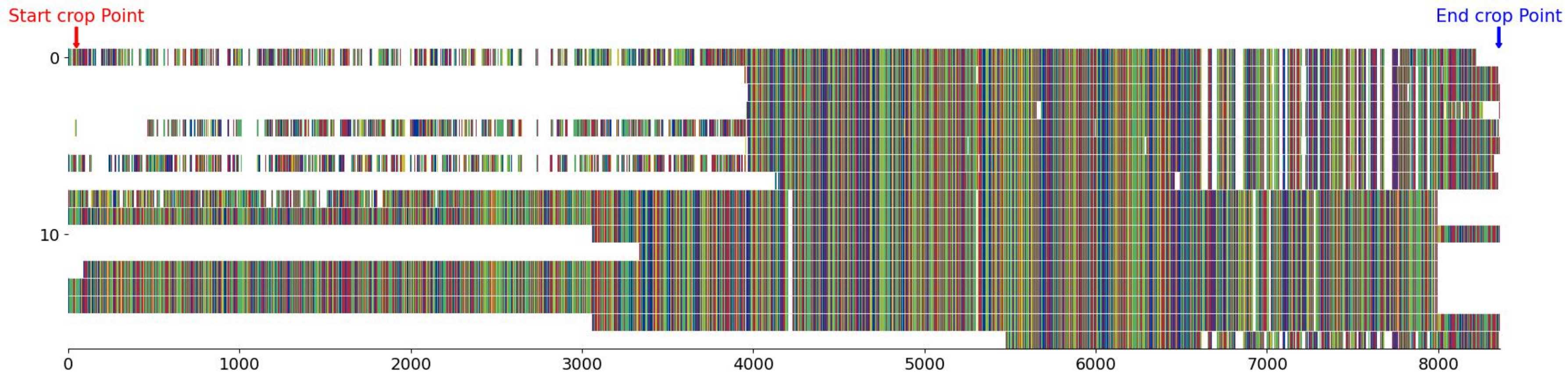


End crop Point



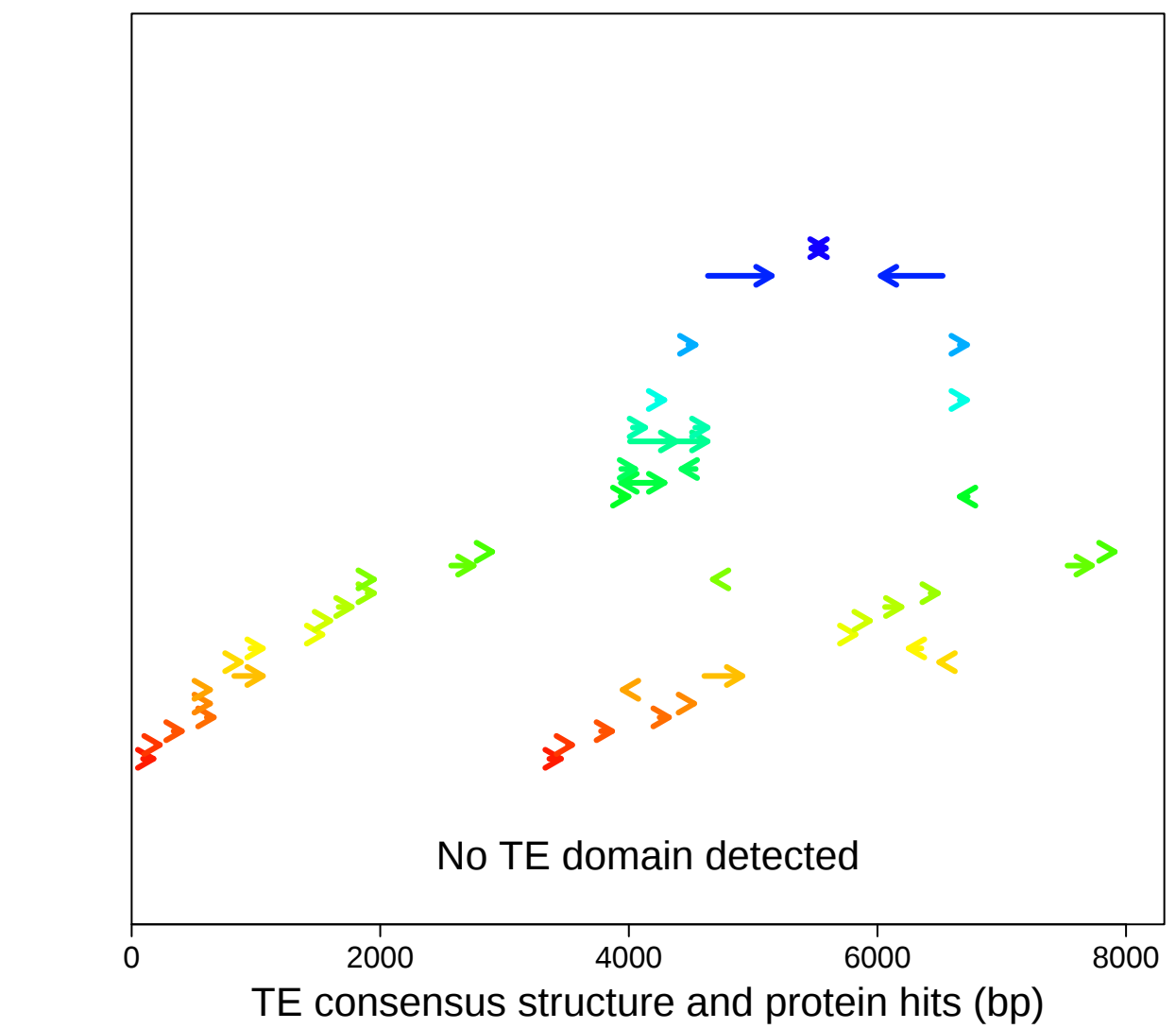
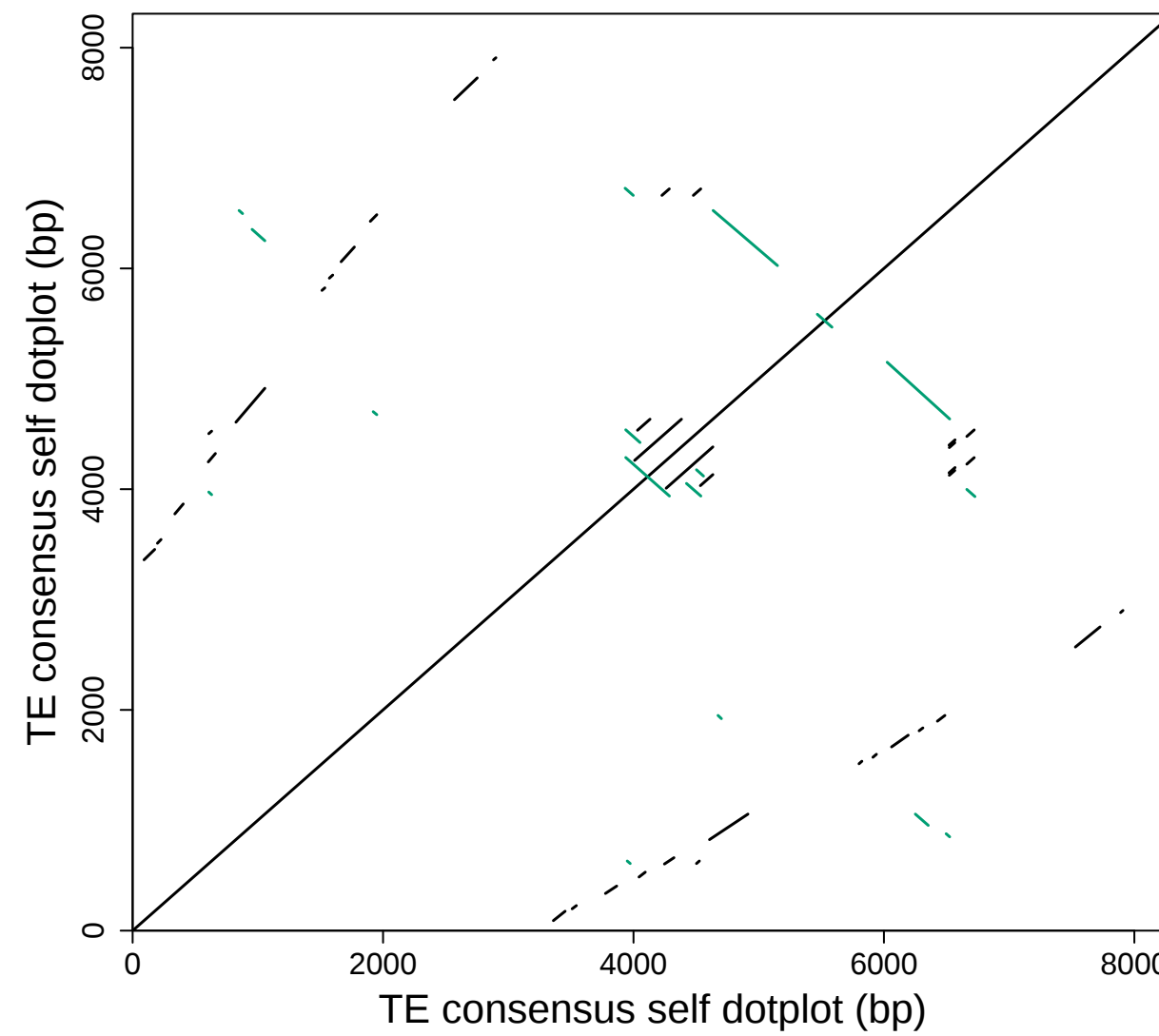
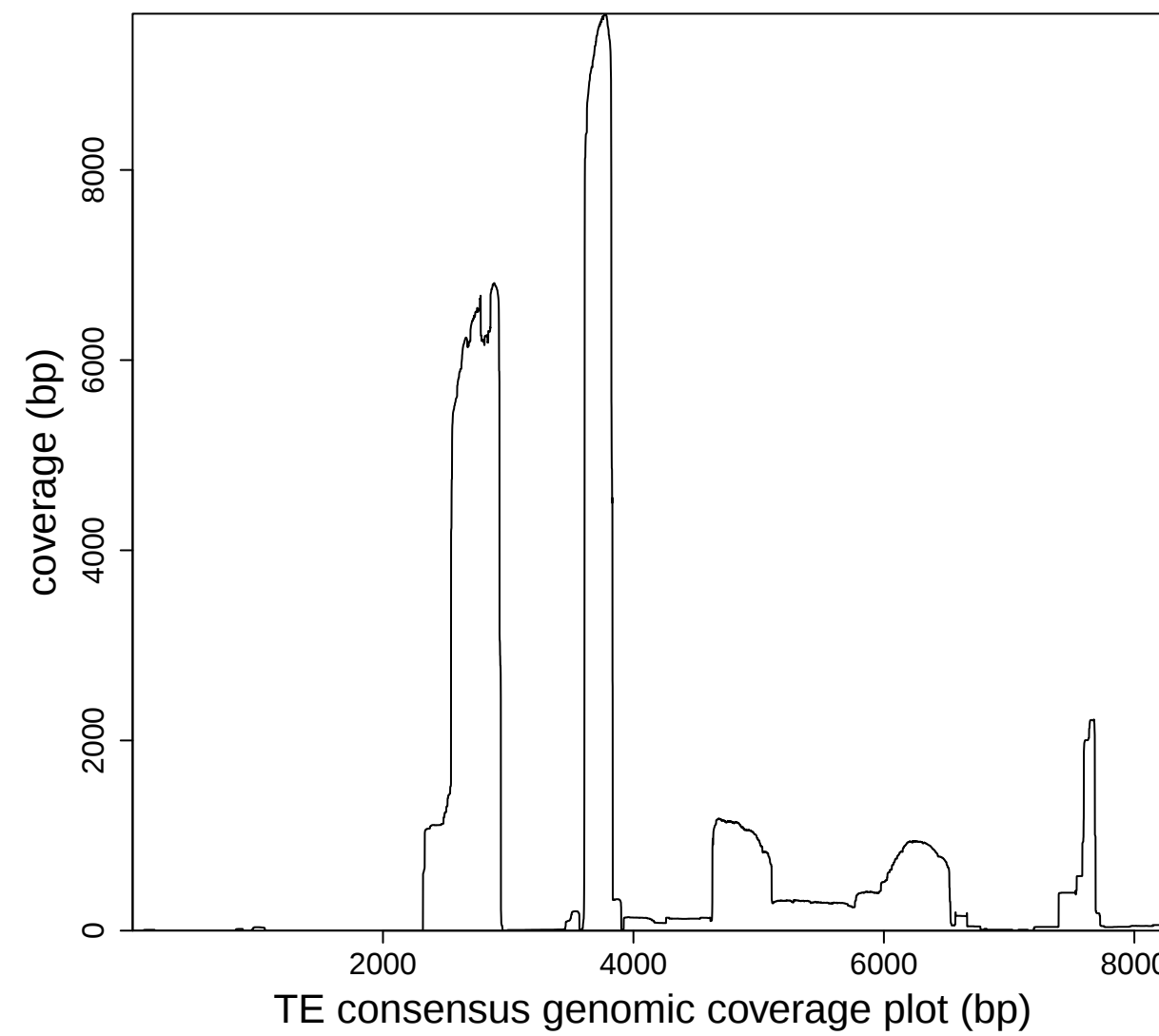
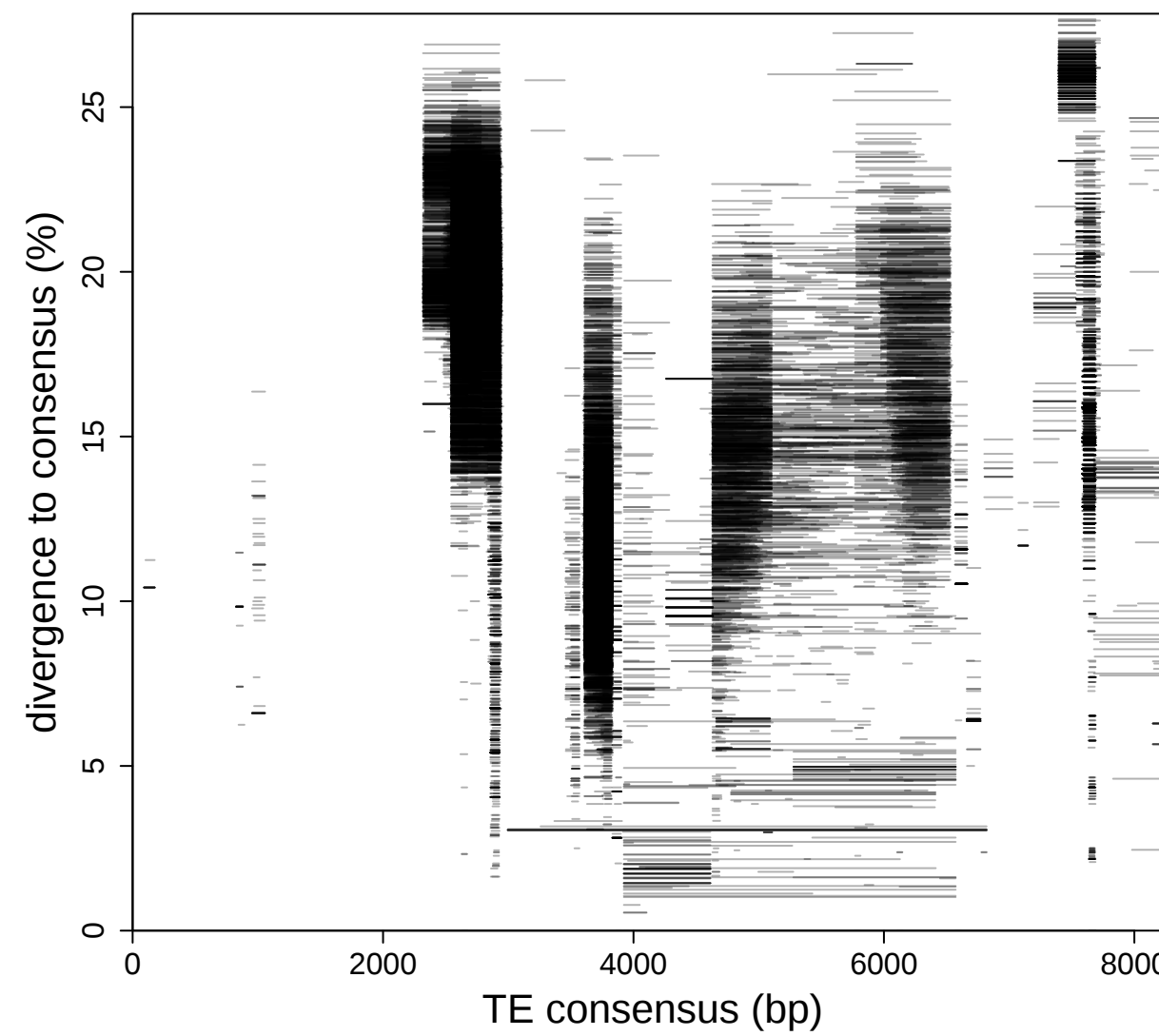
MSA length = 8308





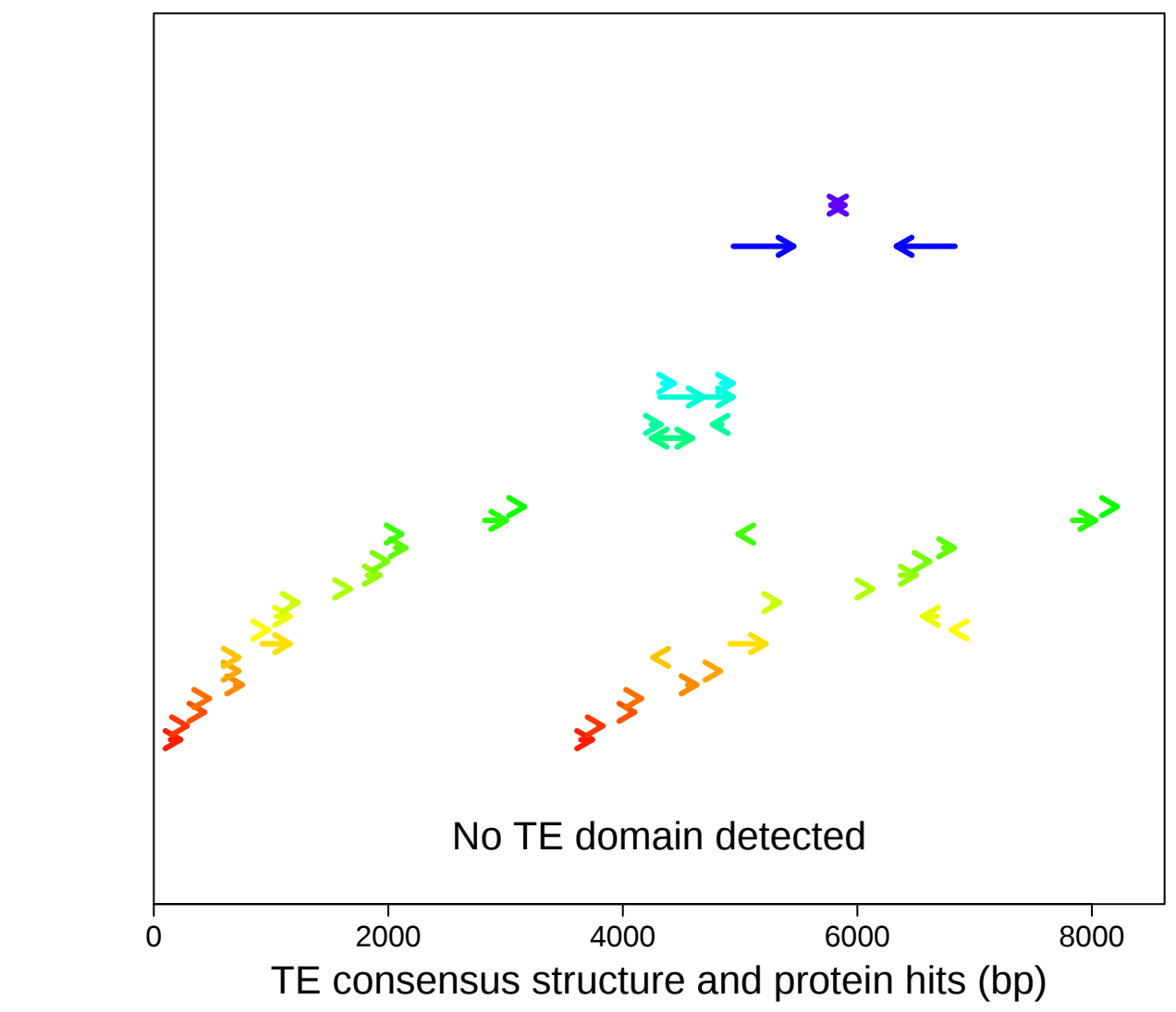
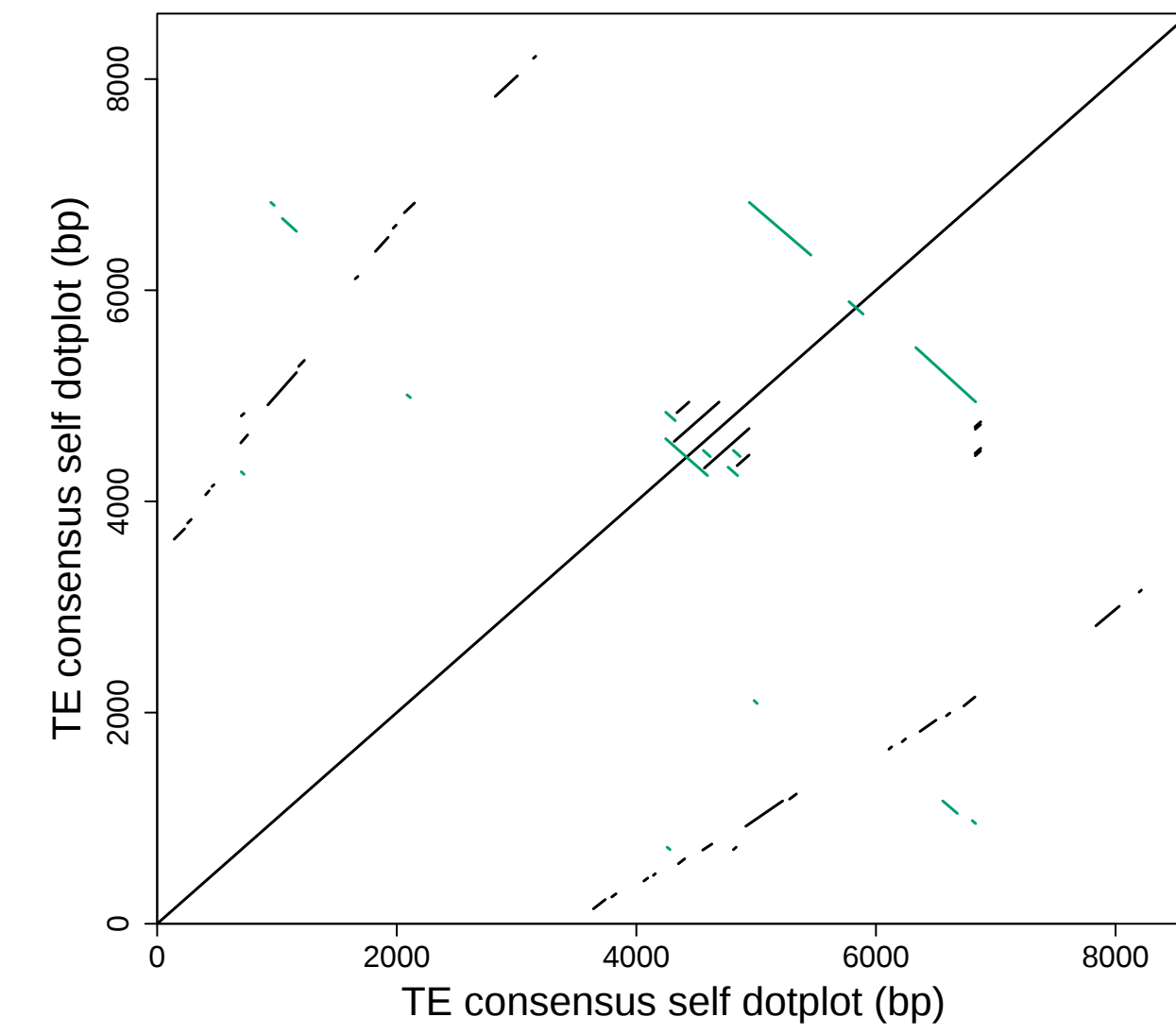
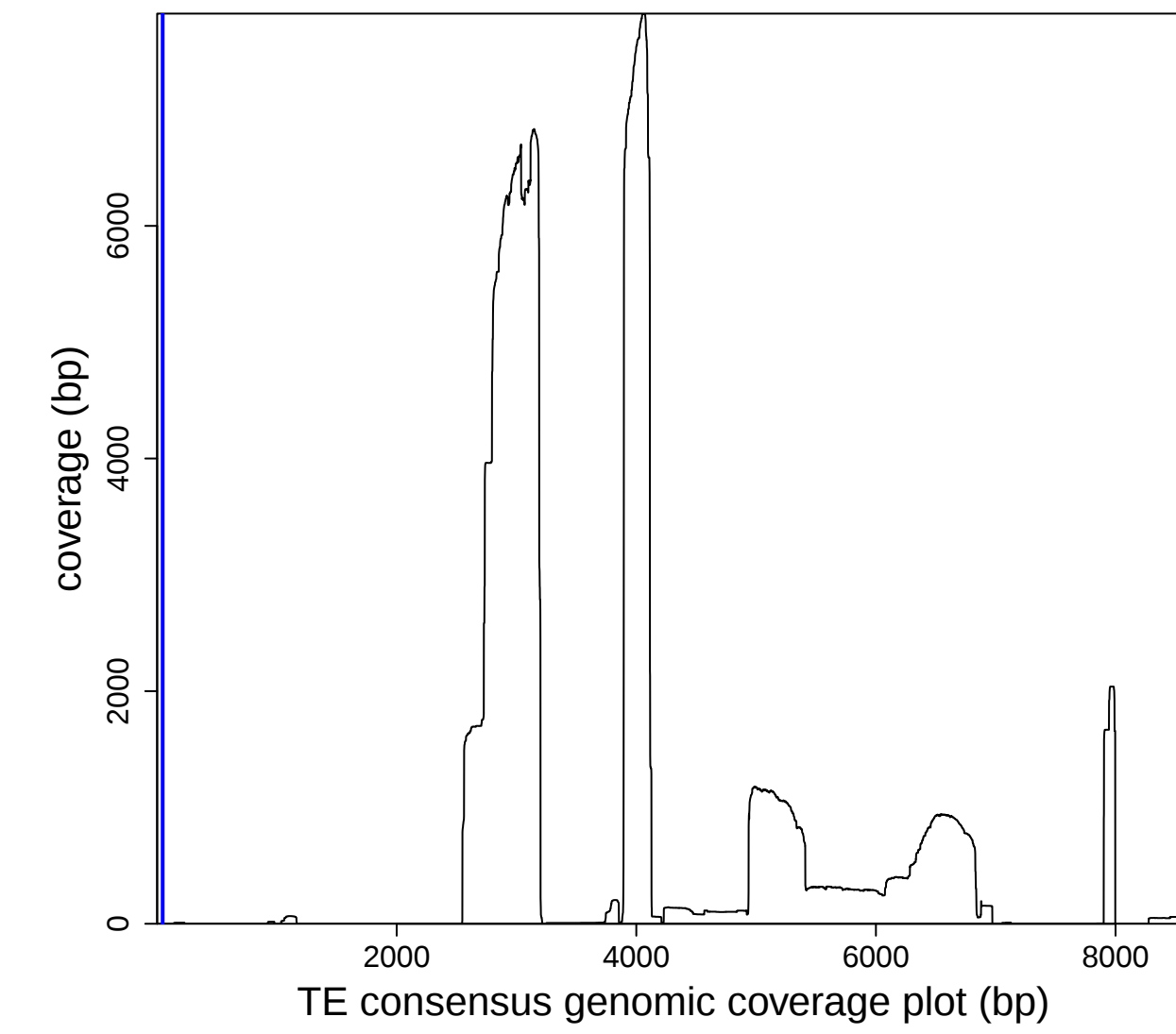
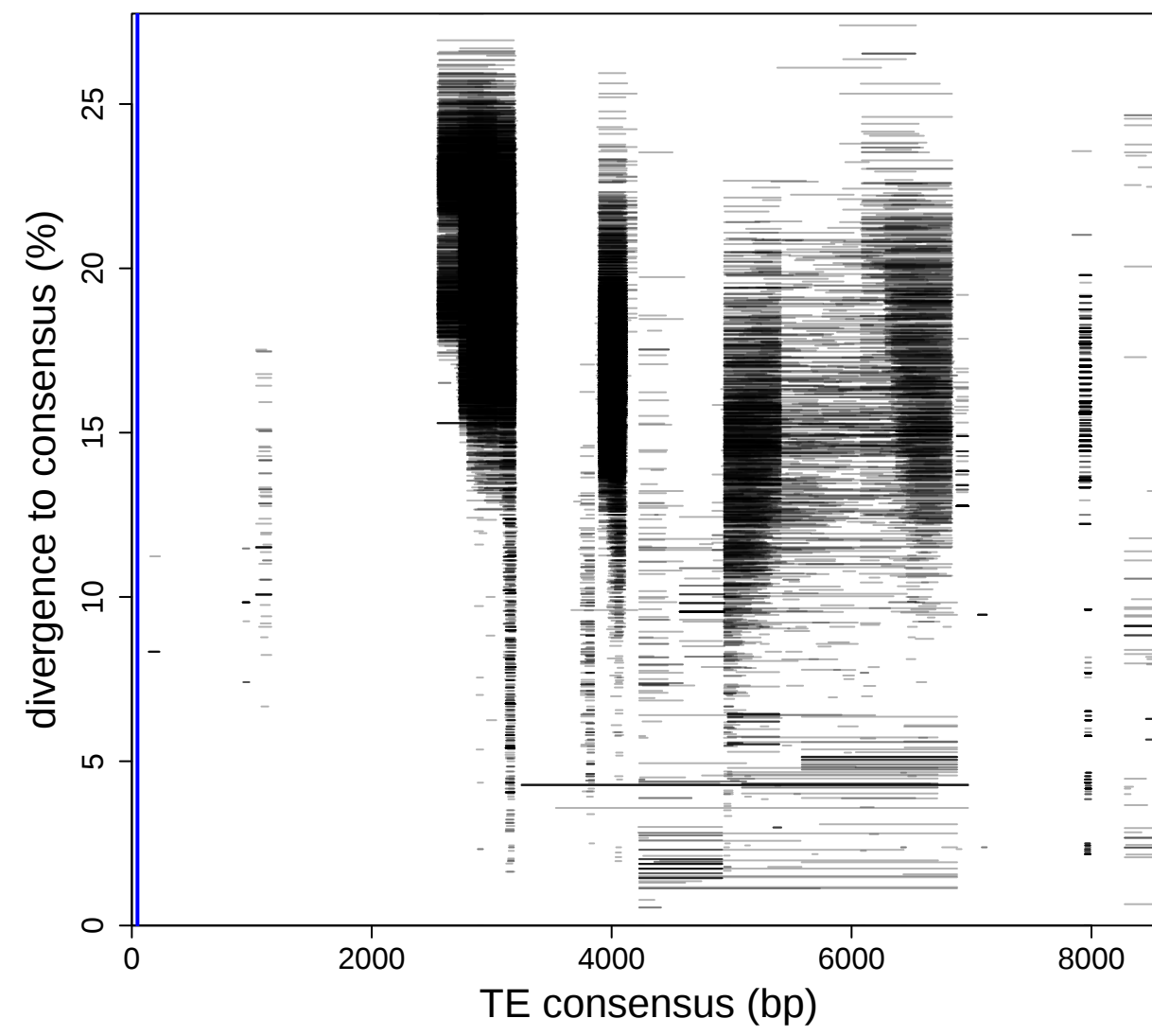
a.b.bed_uf.bed_g_2.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.f
size: 8308bp; fragments: 23645; full length: 0 (>=7477.2bp)

After TEtrimmer 8308 bp

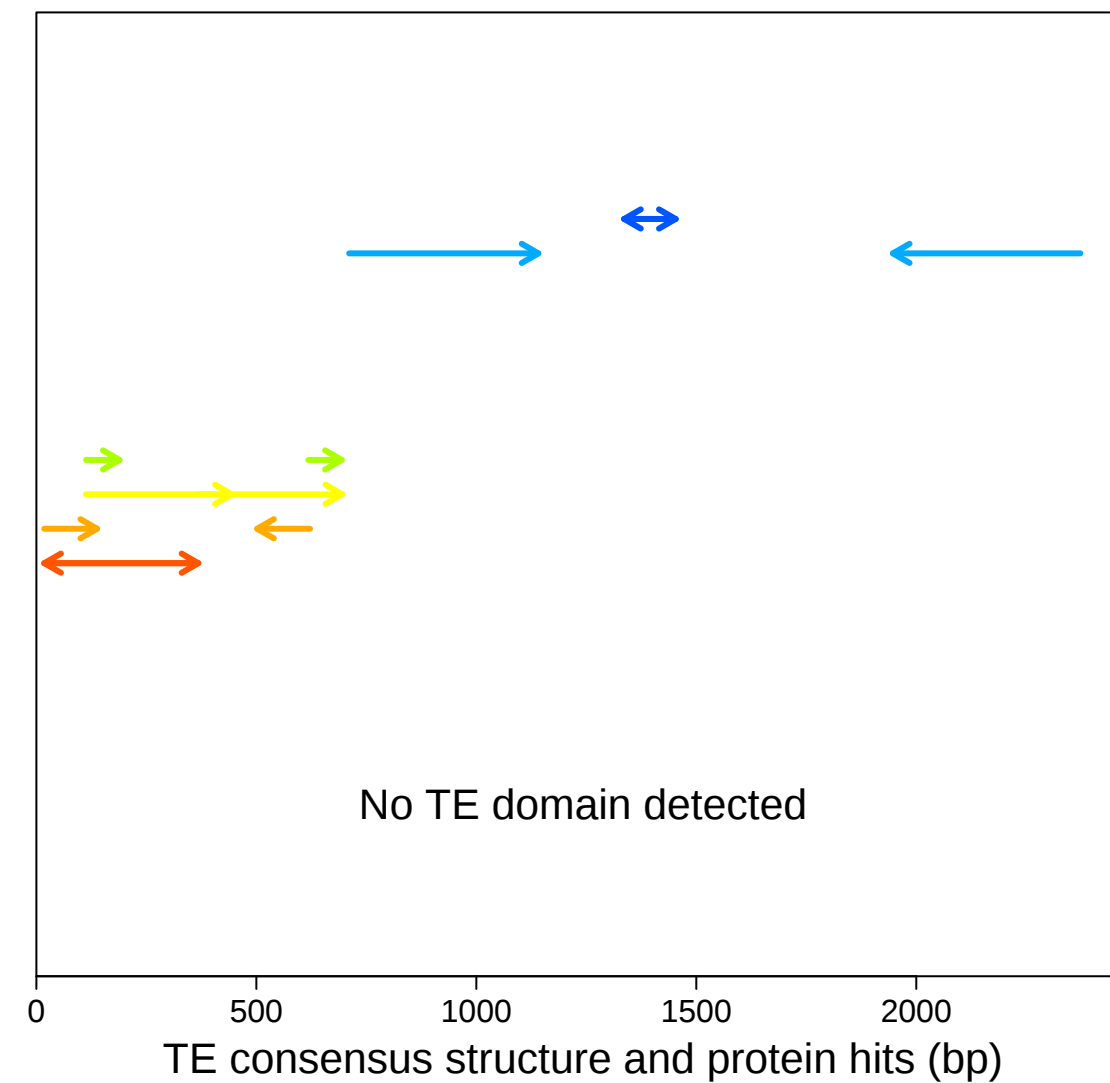
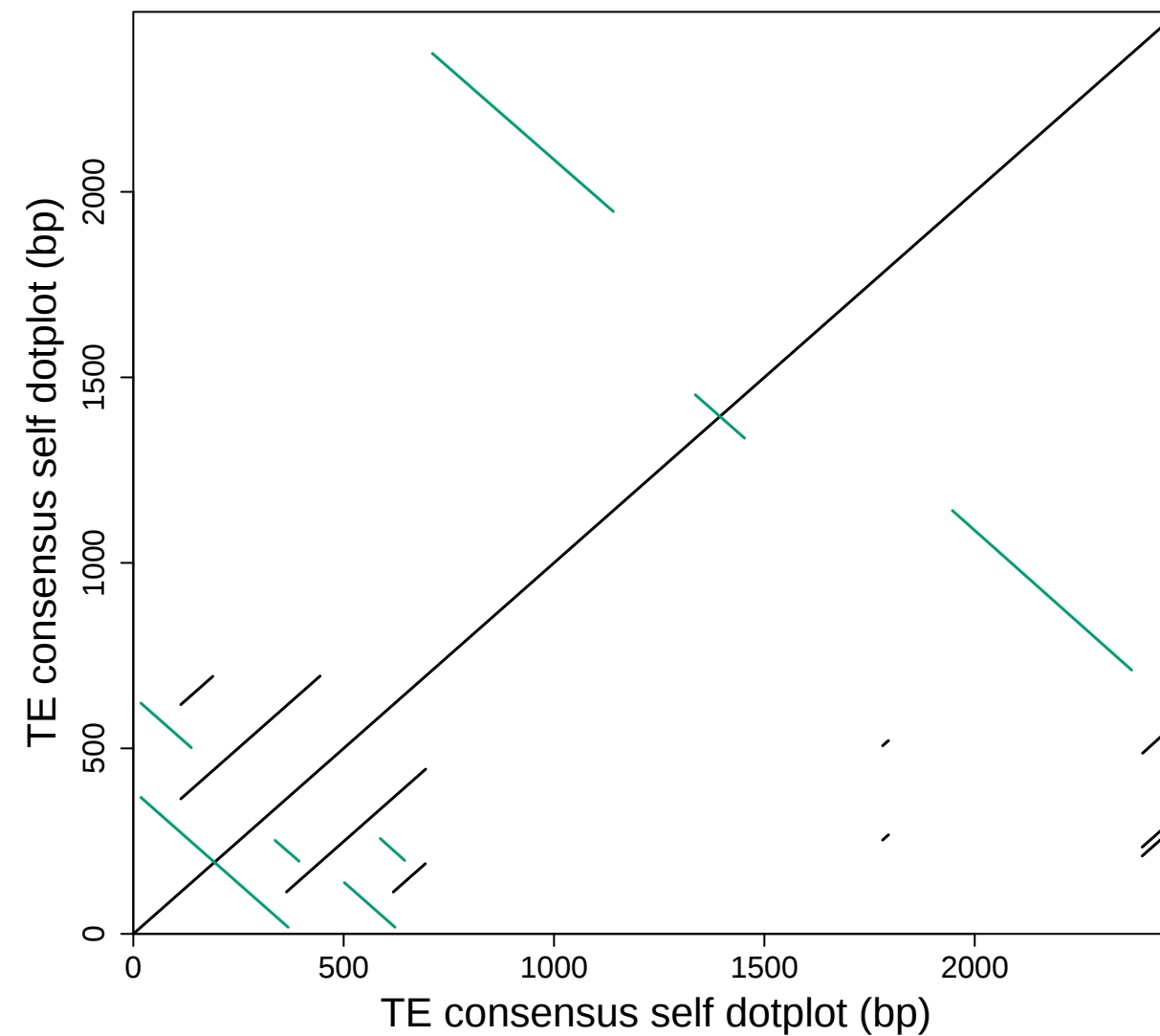
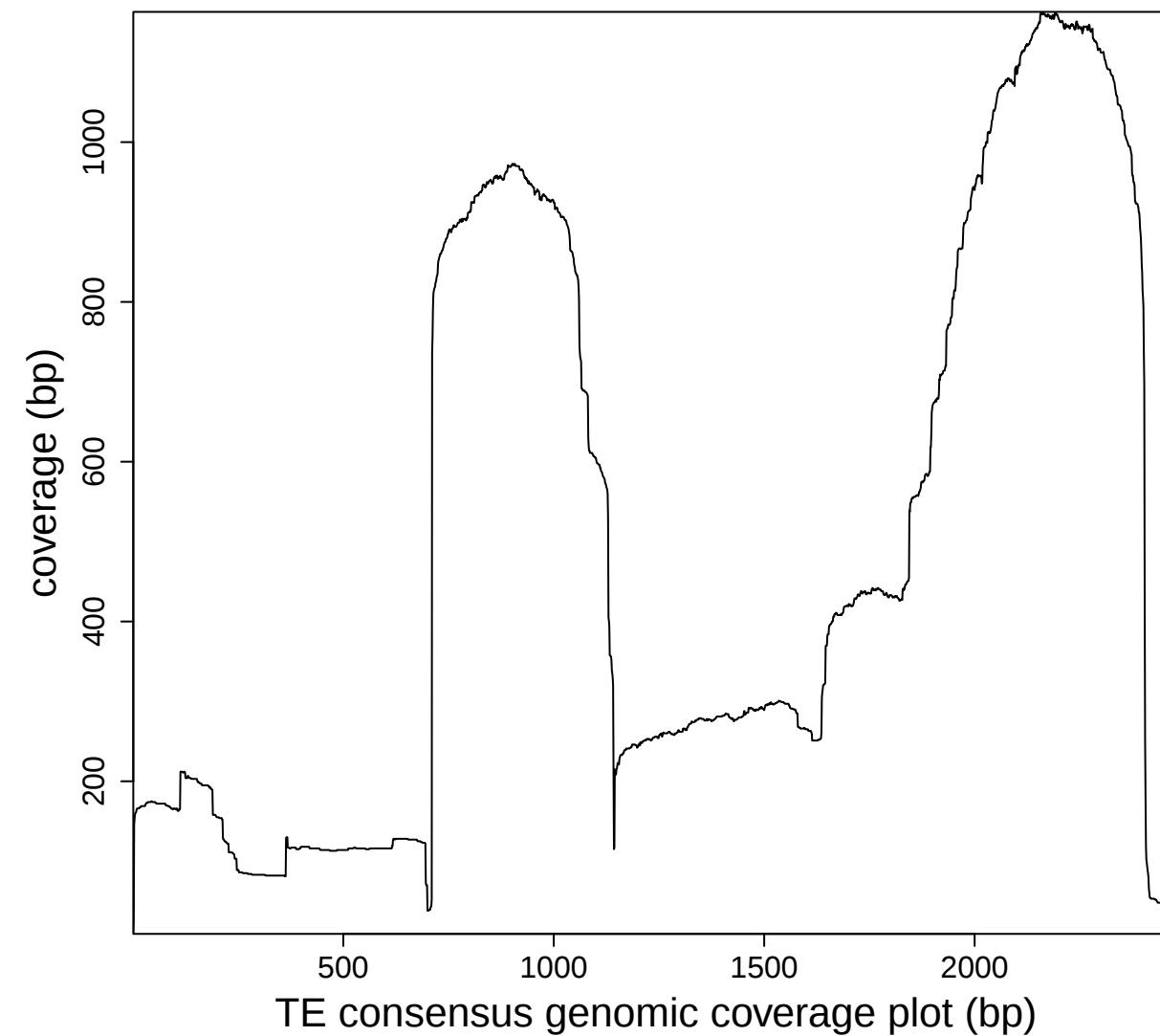
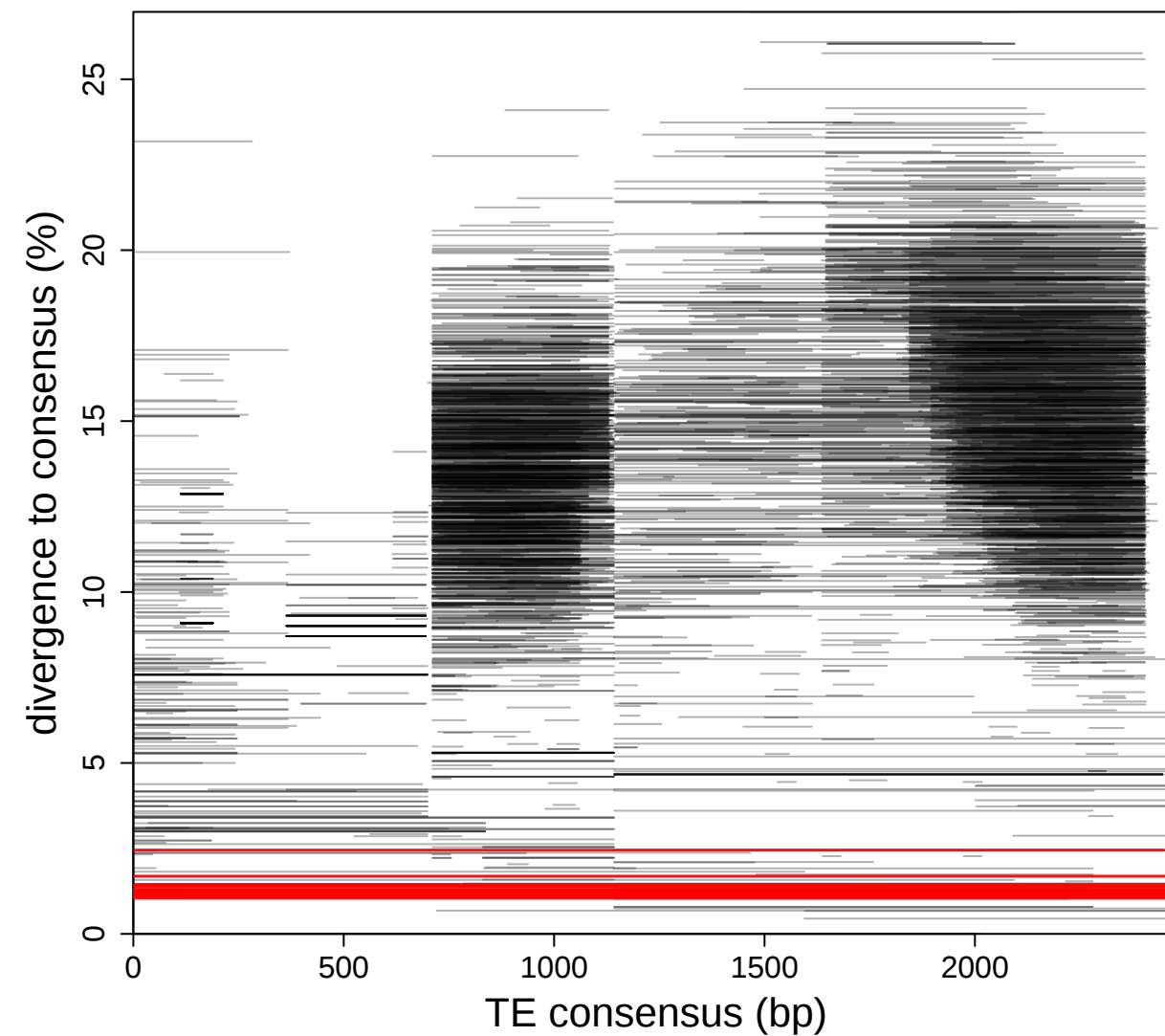


1.fasta.b.bed_uf.bed_g_2.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.f
size: 8620bp; fragments: 21203; full length: 0 (>=7758bp)

After TEtrimmer Extended plot Blue lines are boundaries

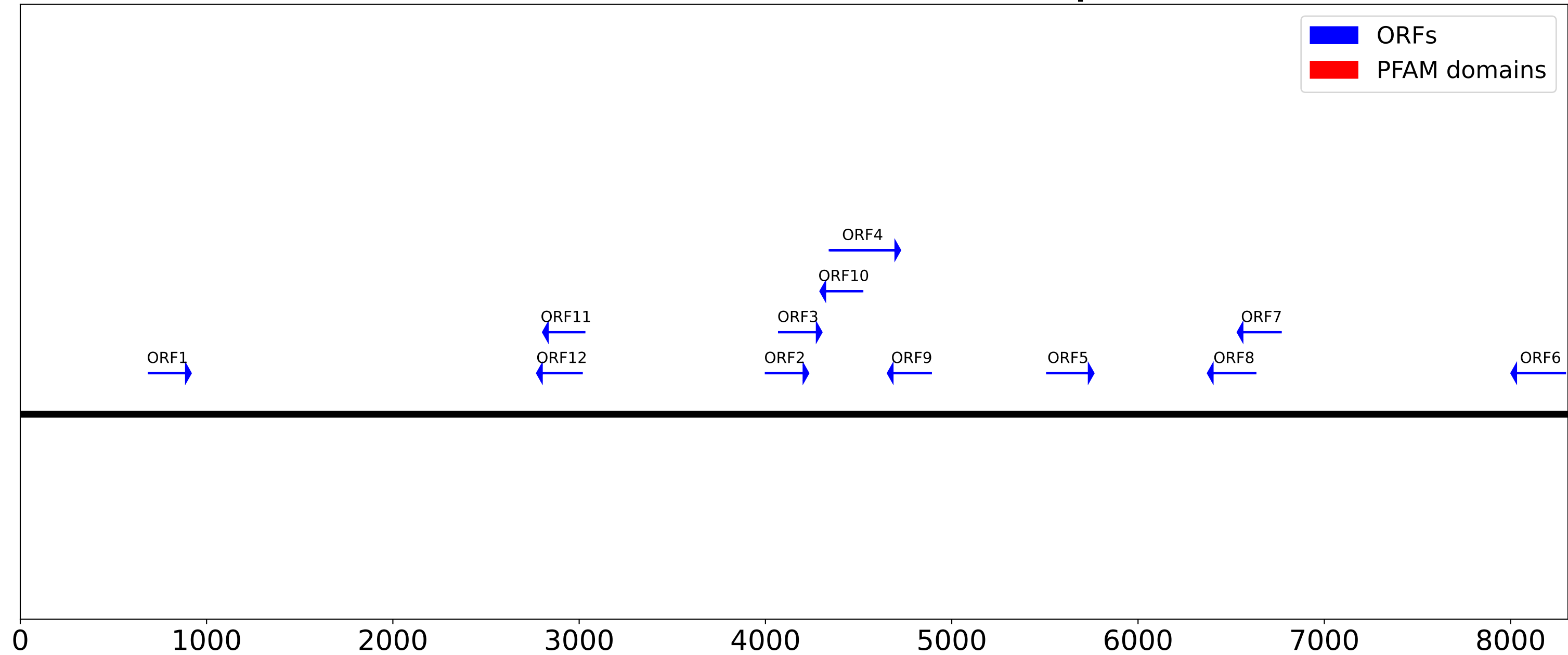


TE: rnd_5_family_2771
size: 2485bp; fragments: 3042; full length: 18 (≥ 2236.5 bp)

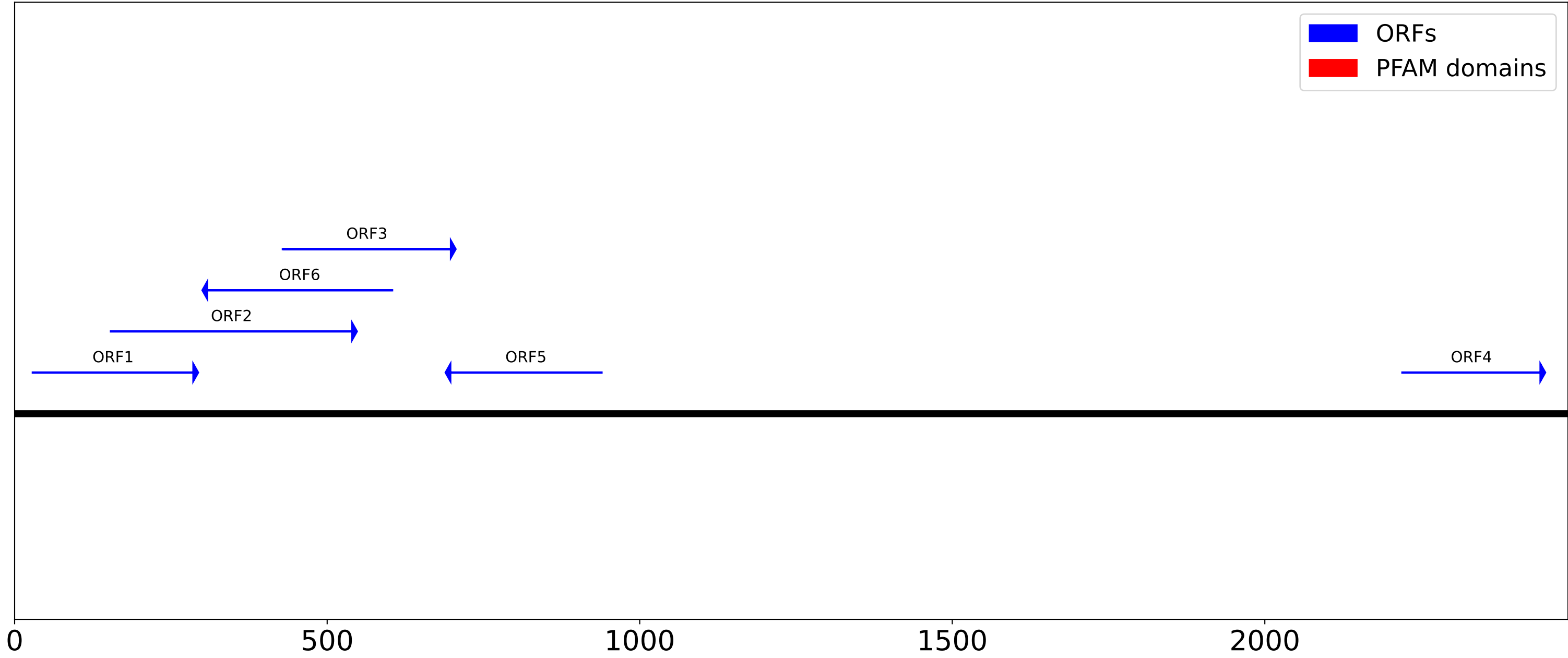


Before TEtrimmer 2485 bp

After TEtrimmer ORF and PFAM domain plot



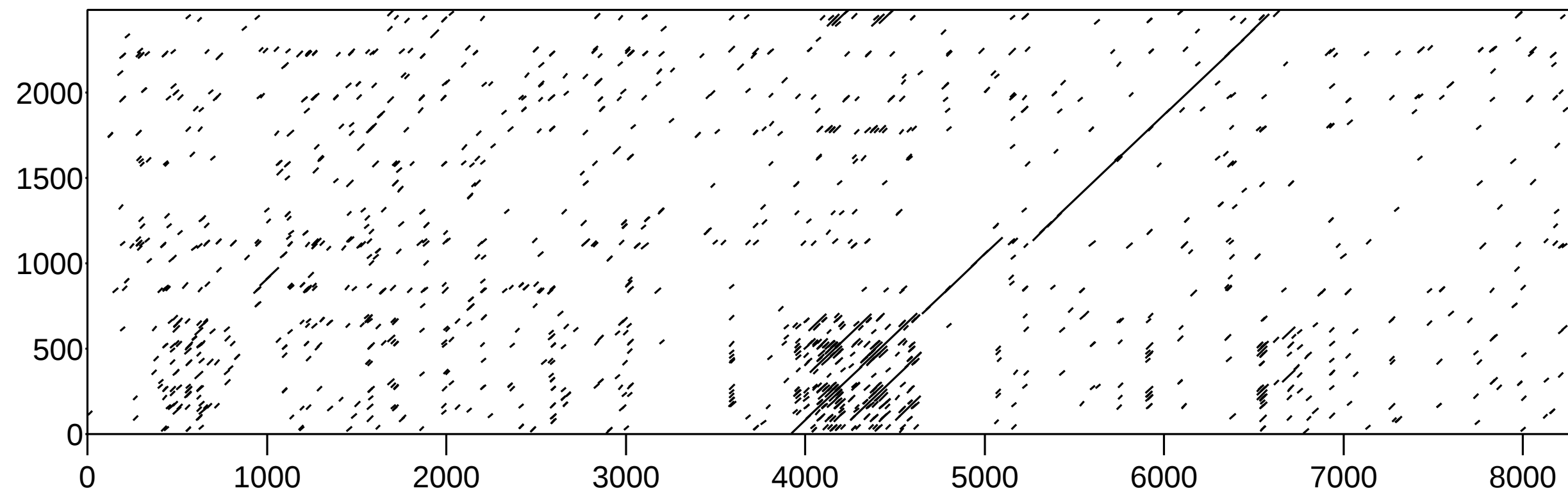
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)