

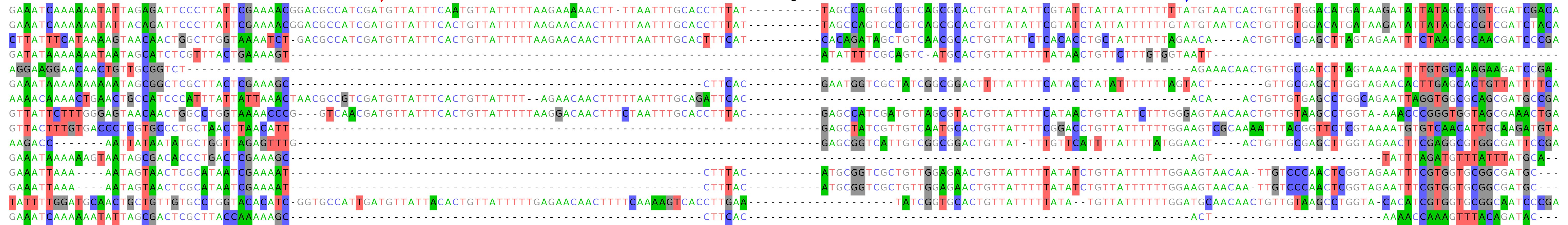
Start crop Point

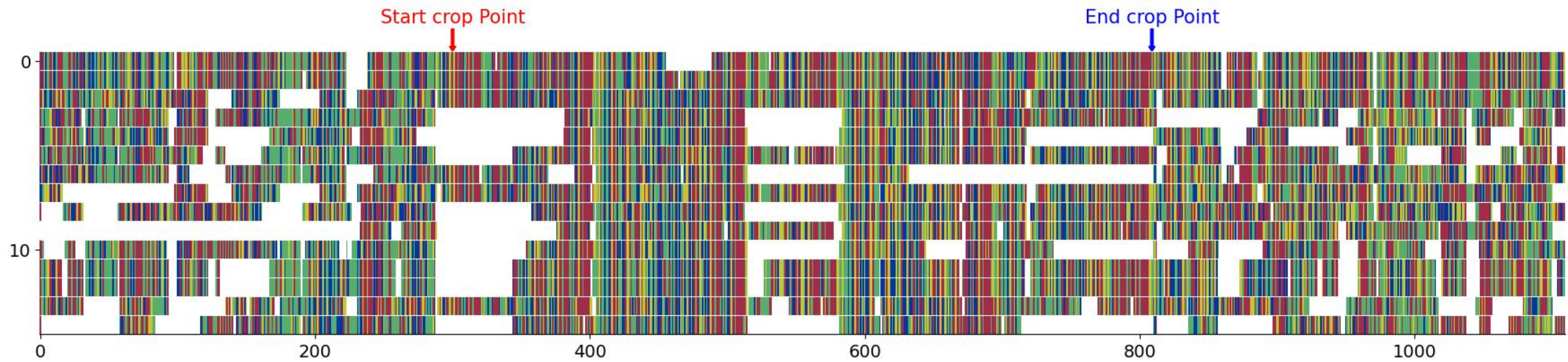


End crop Point



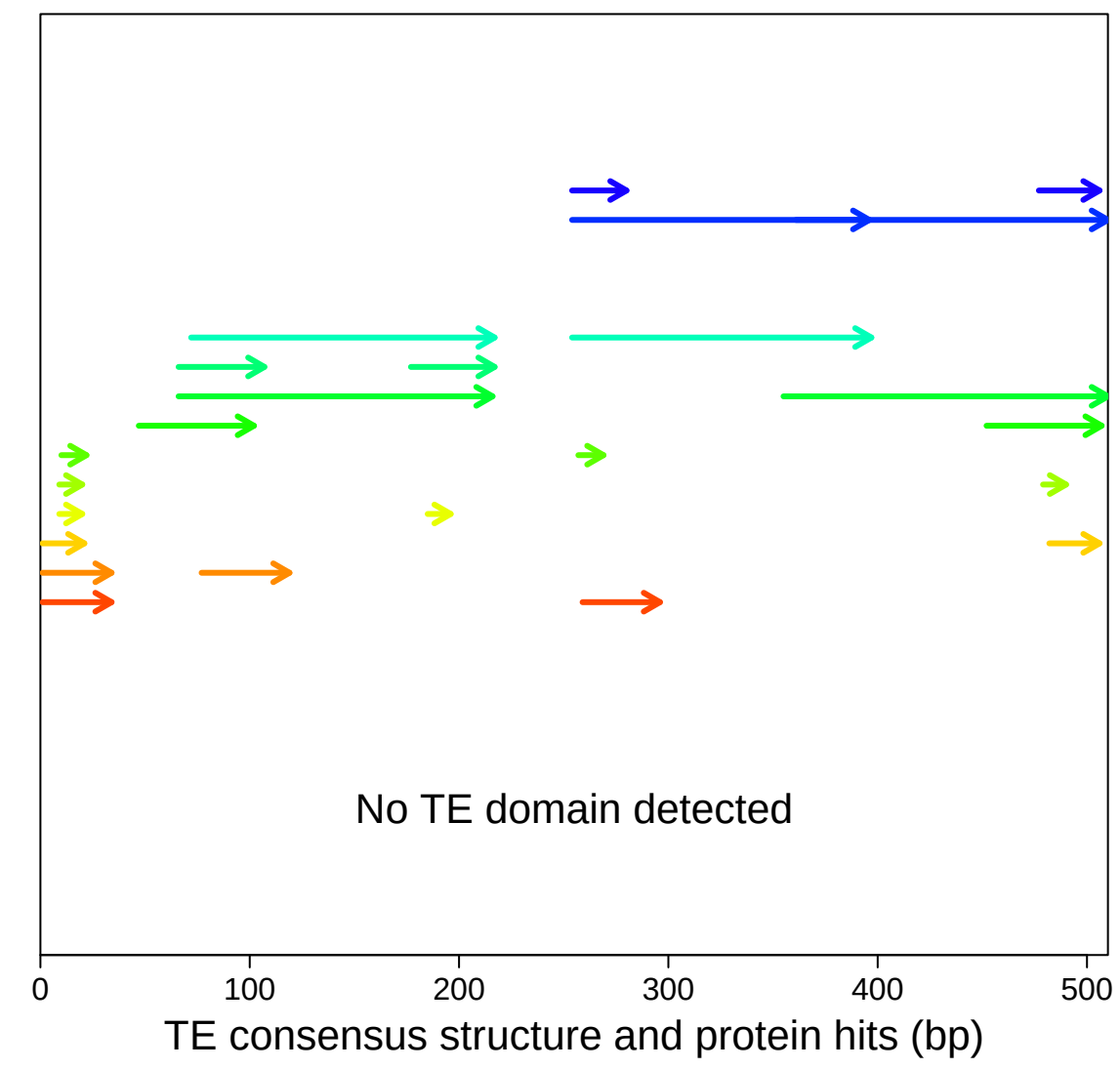
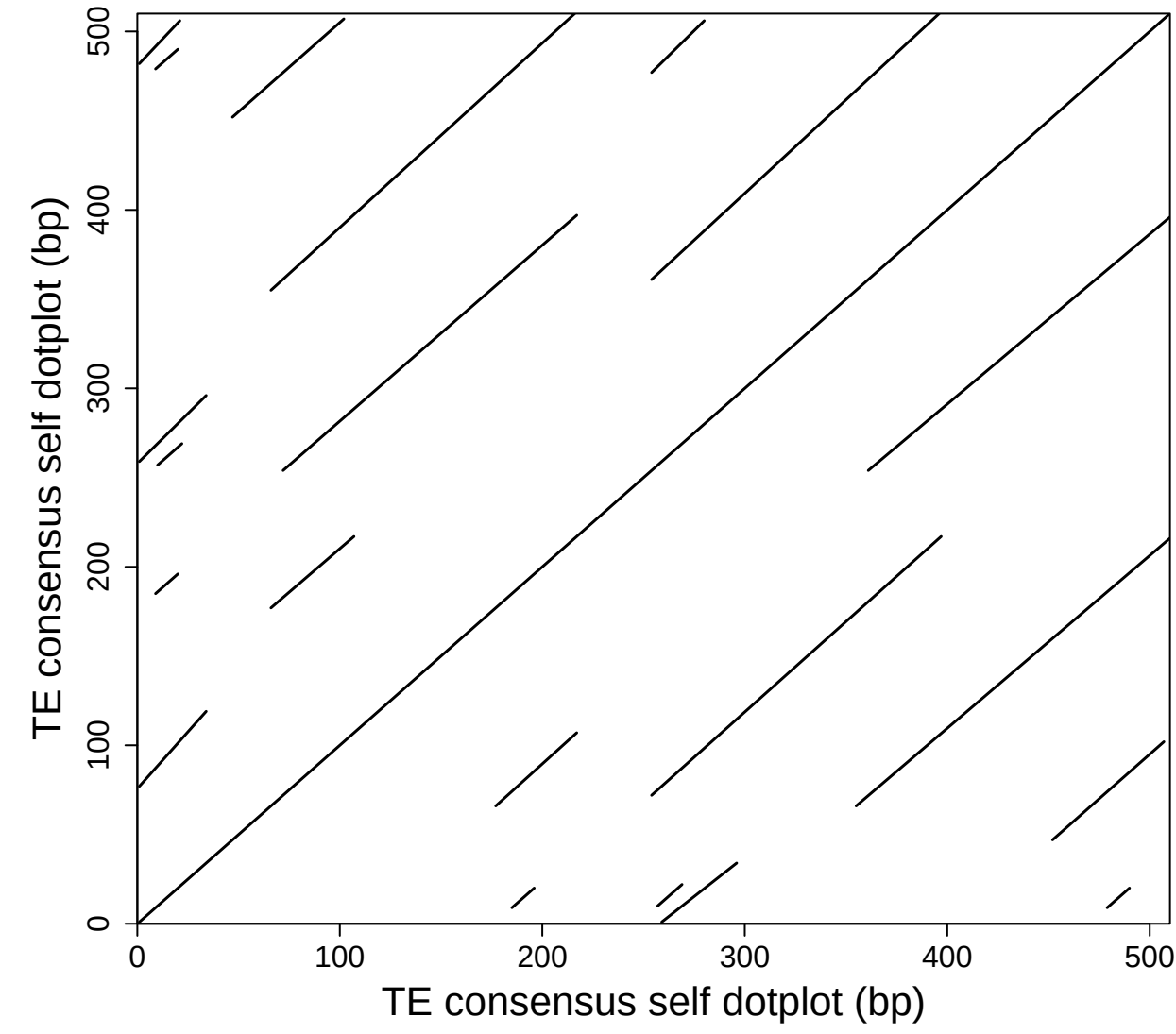
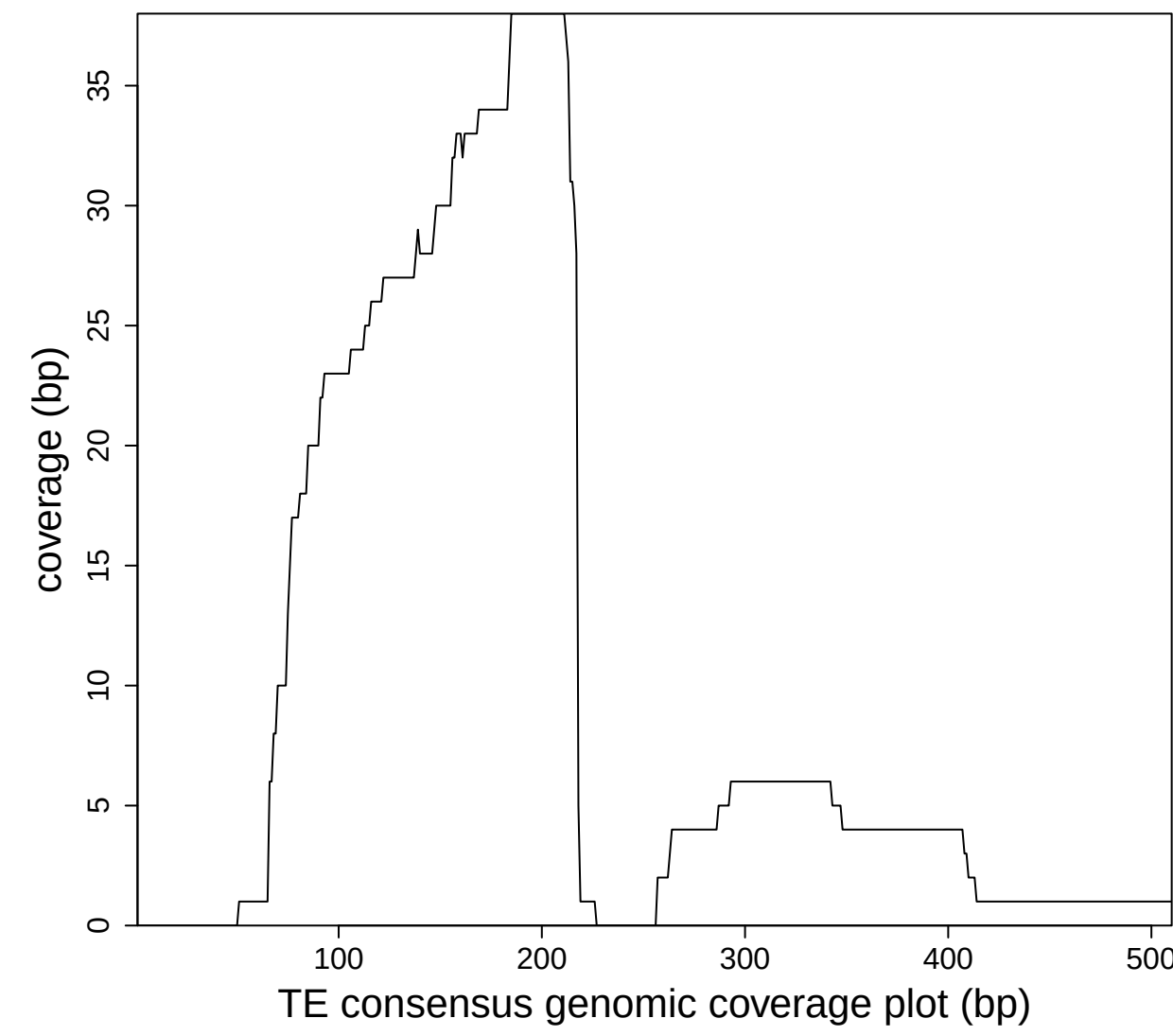
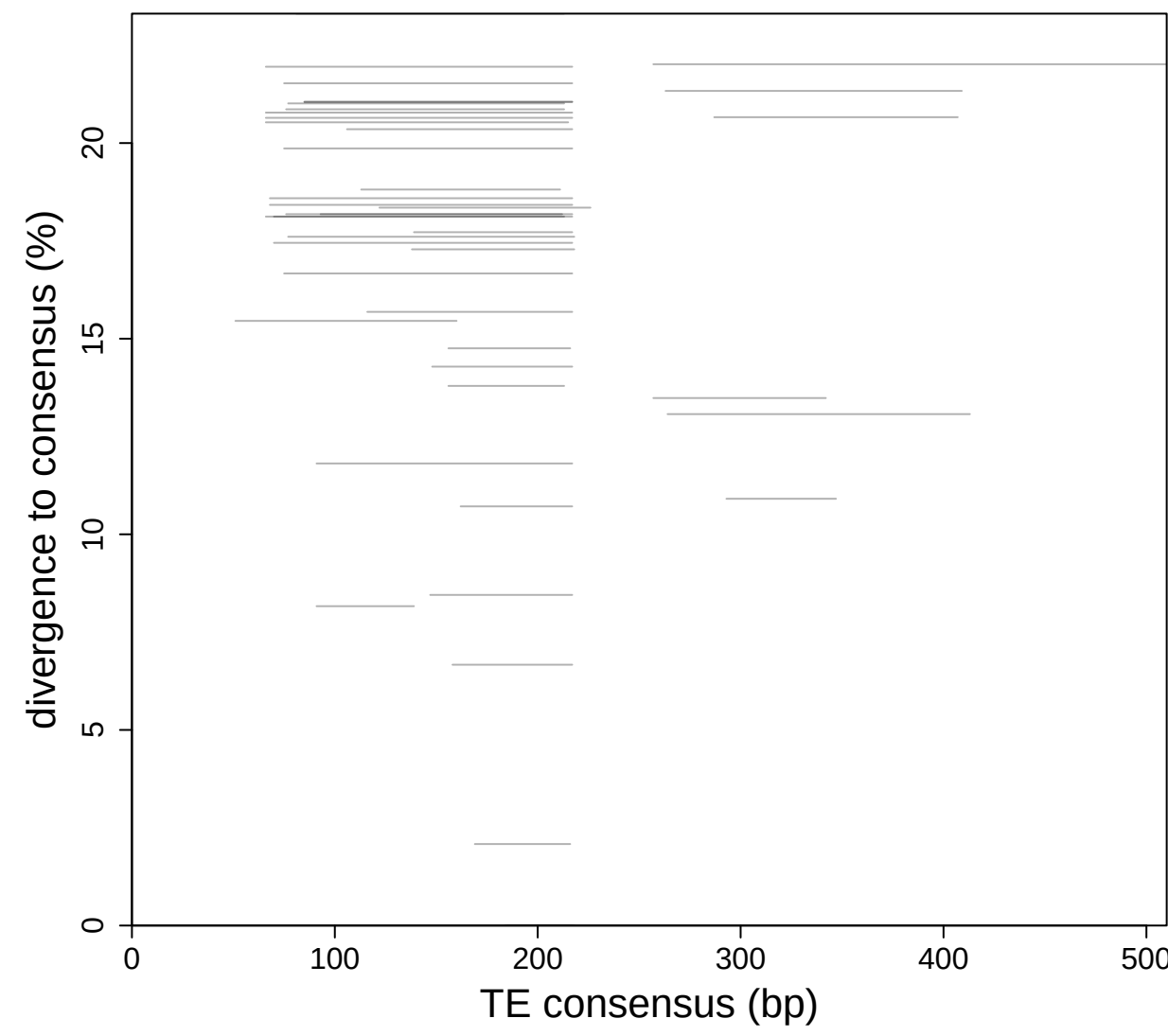
MSA length = 510





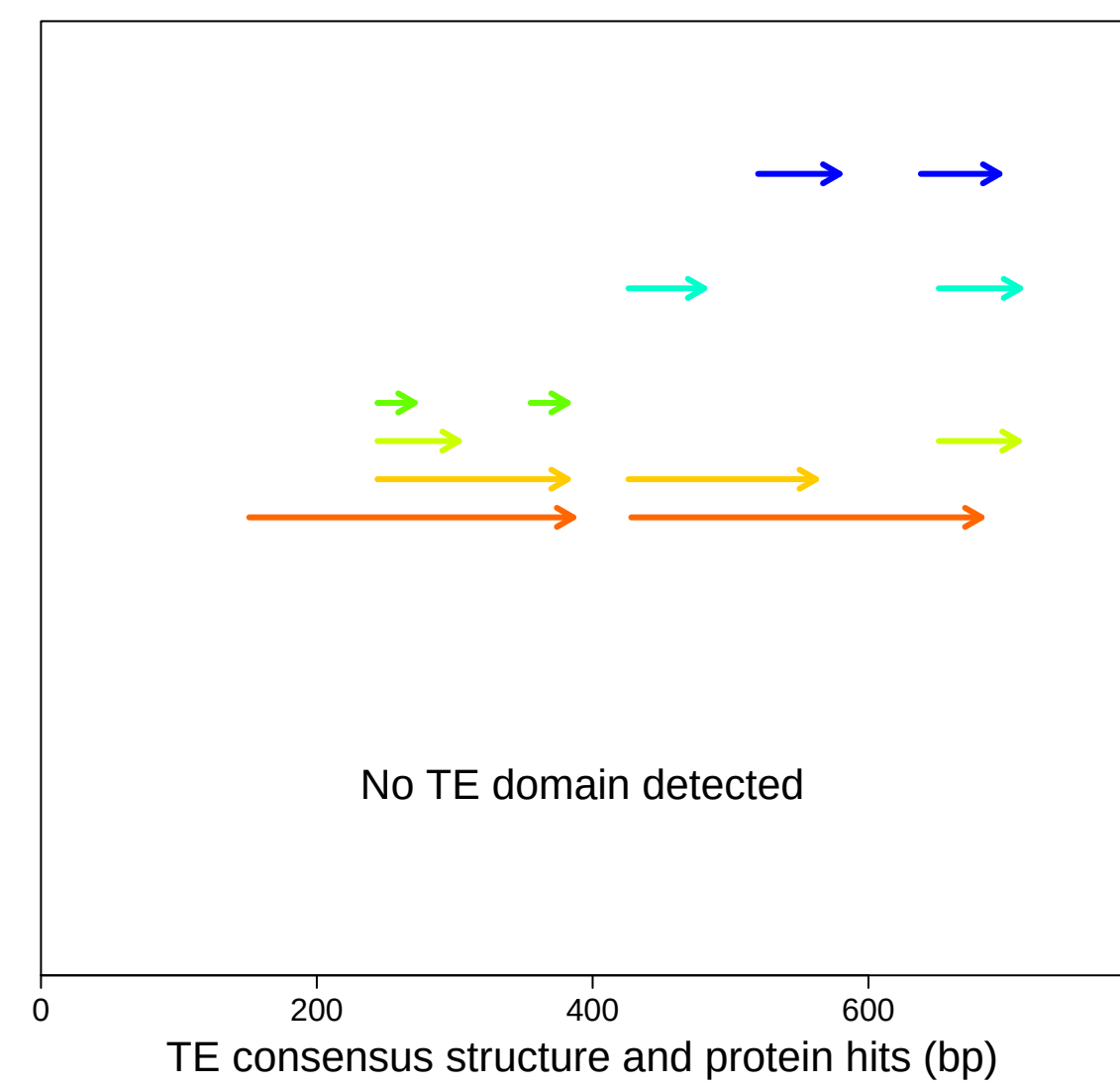
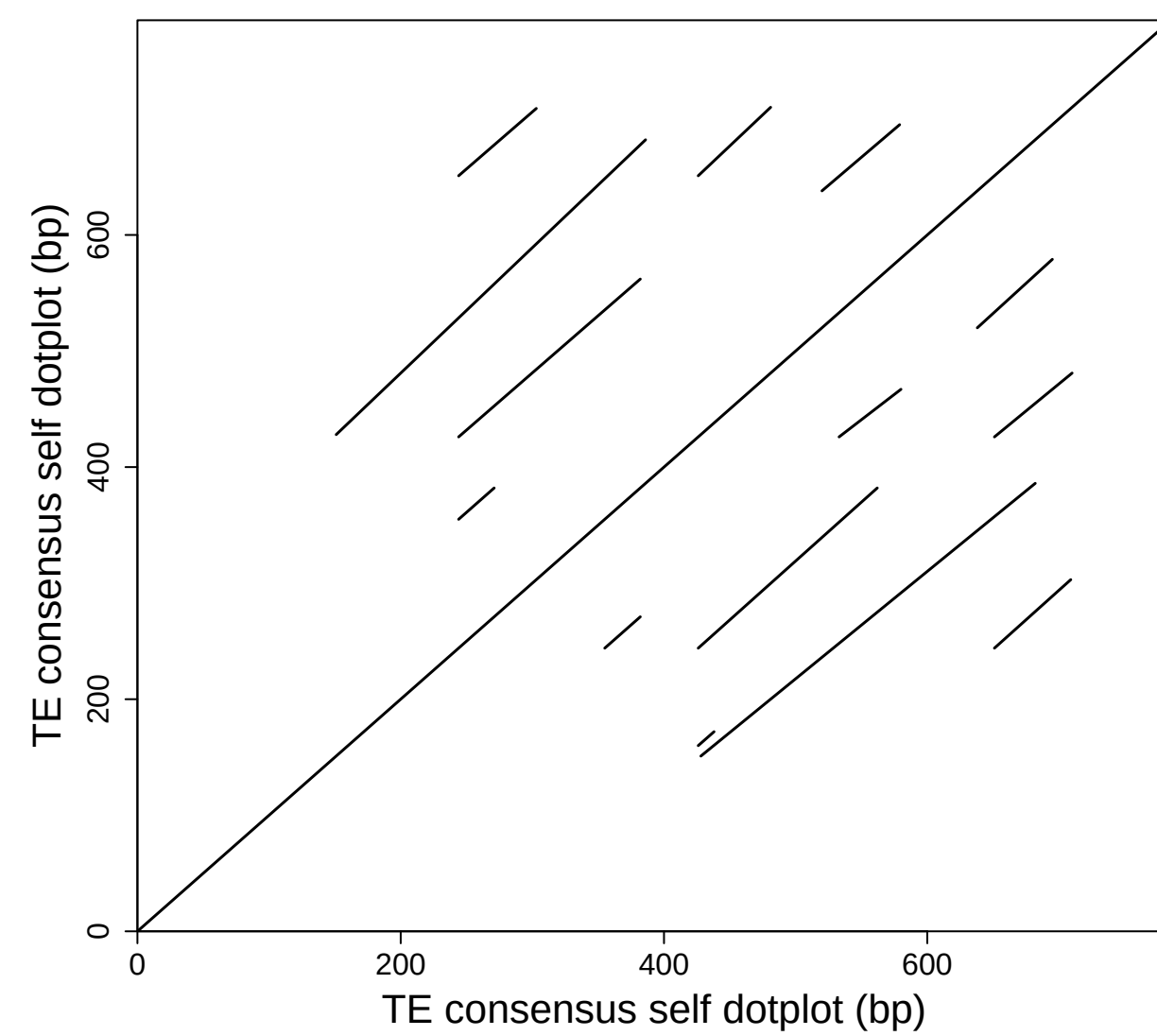
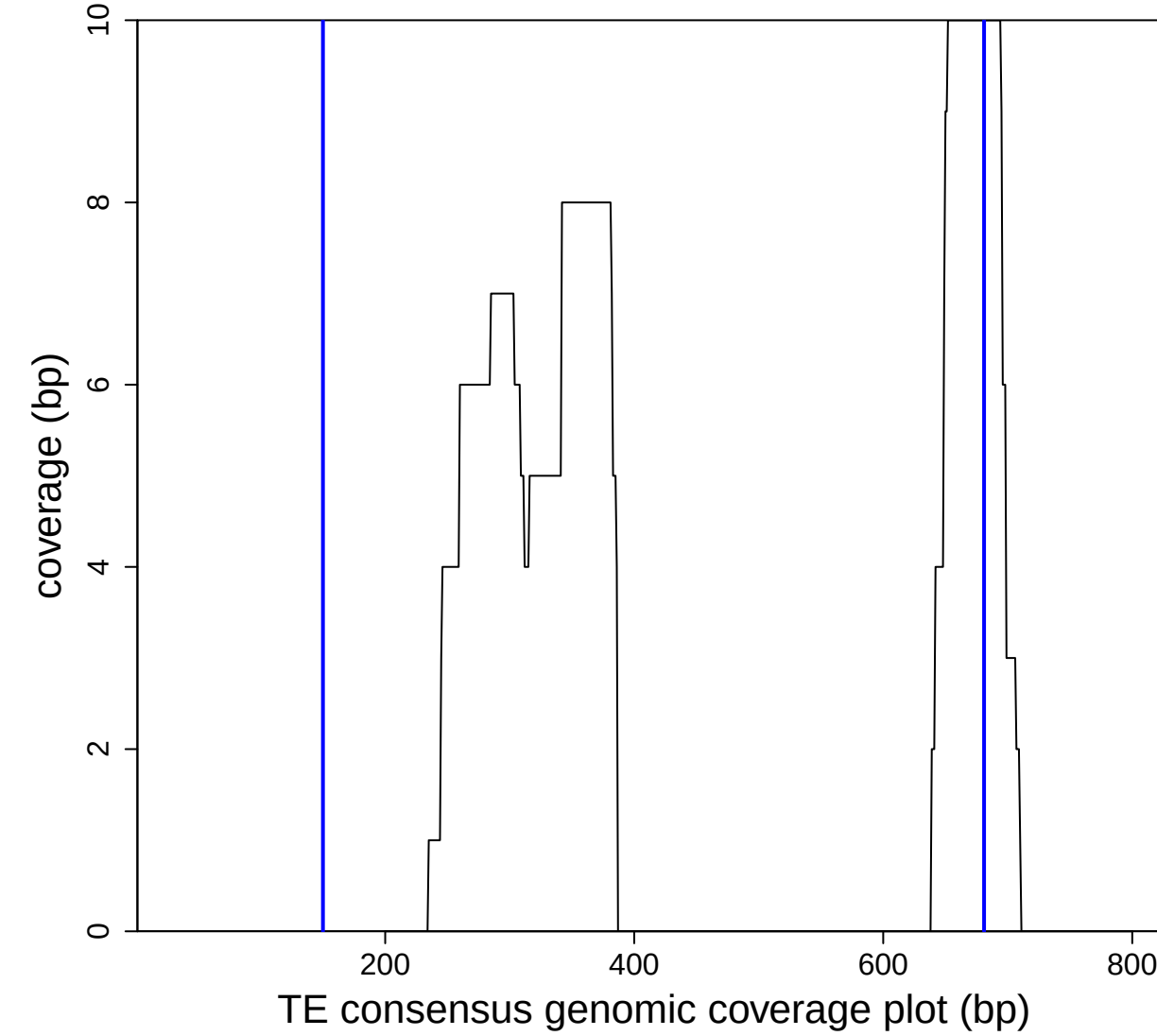
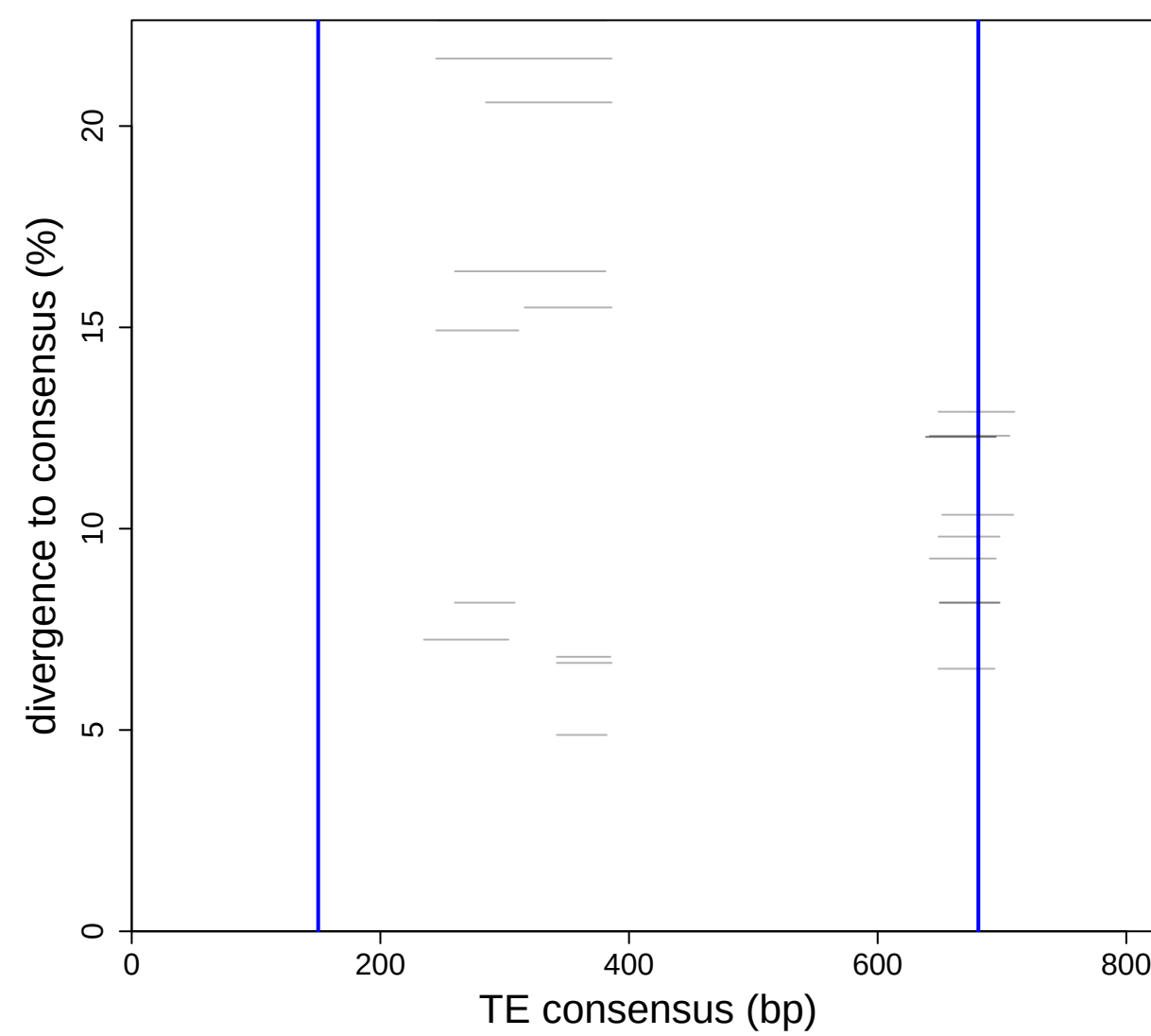
ed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.fa_ce.
size: 510bp; fragments: 46; full length: 0 (>=459bp)

After TEtrimmer 510 bp



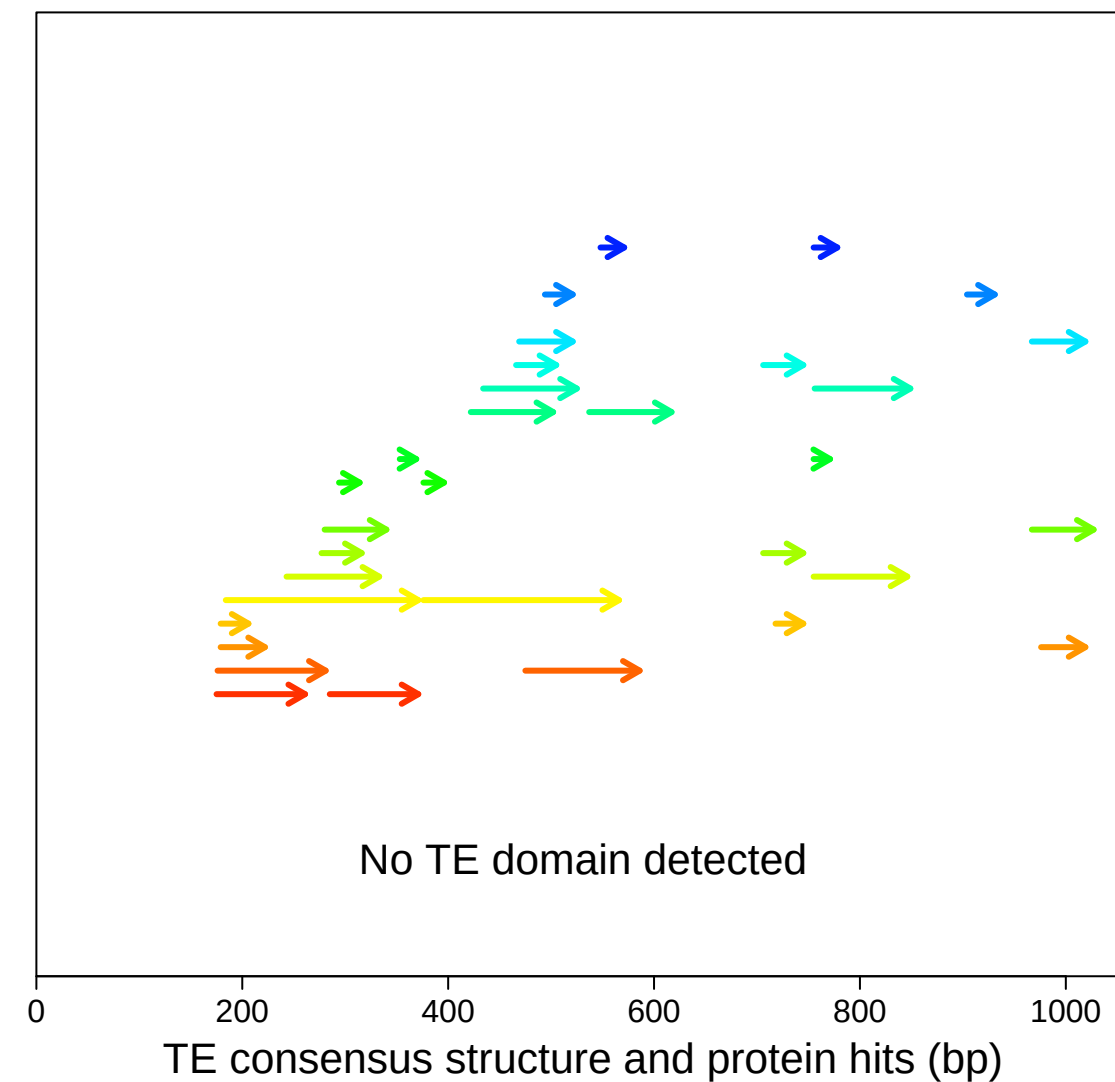
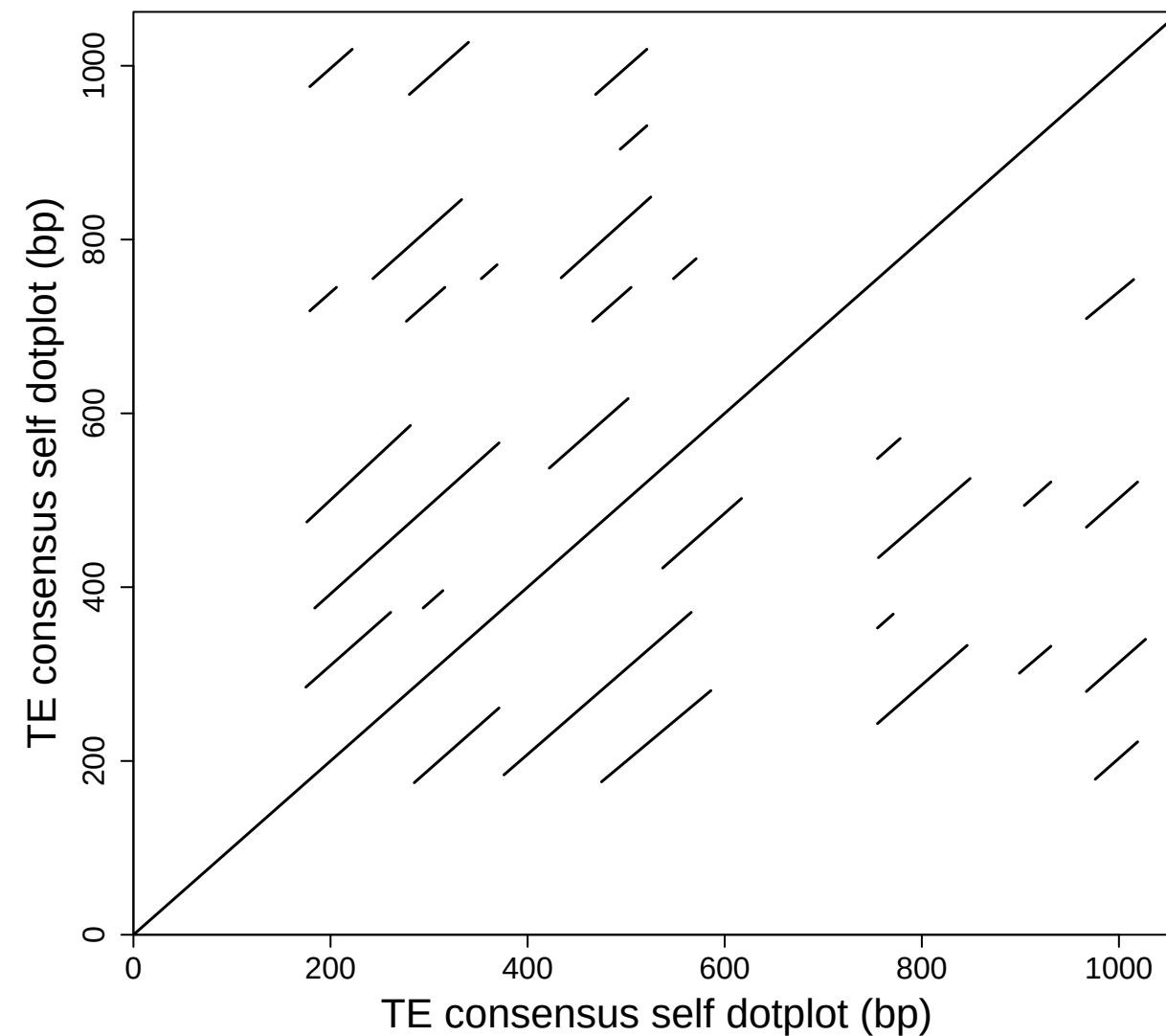
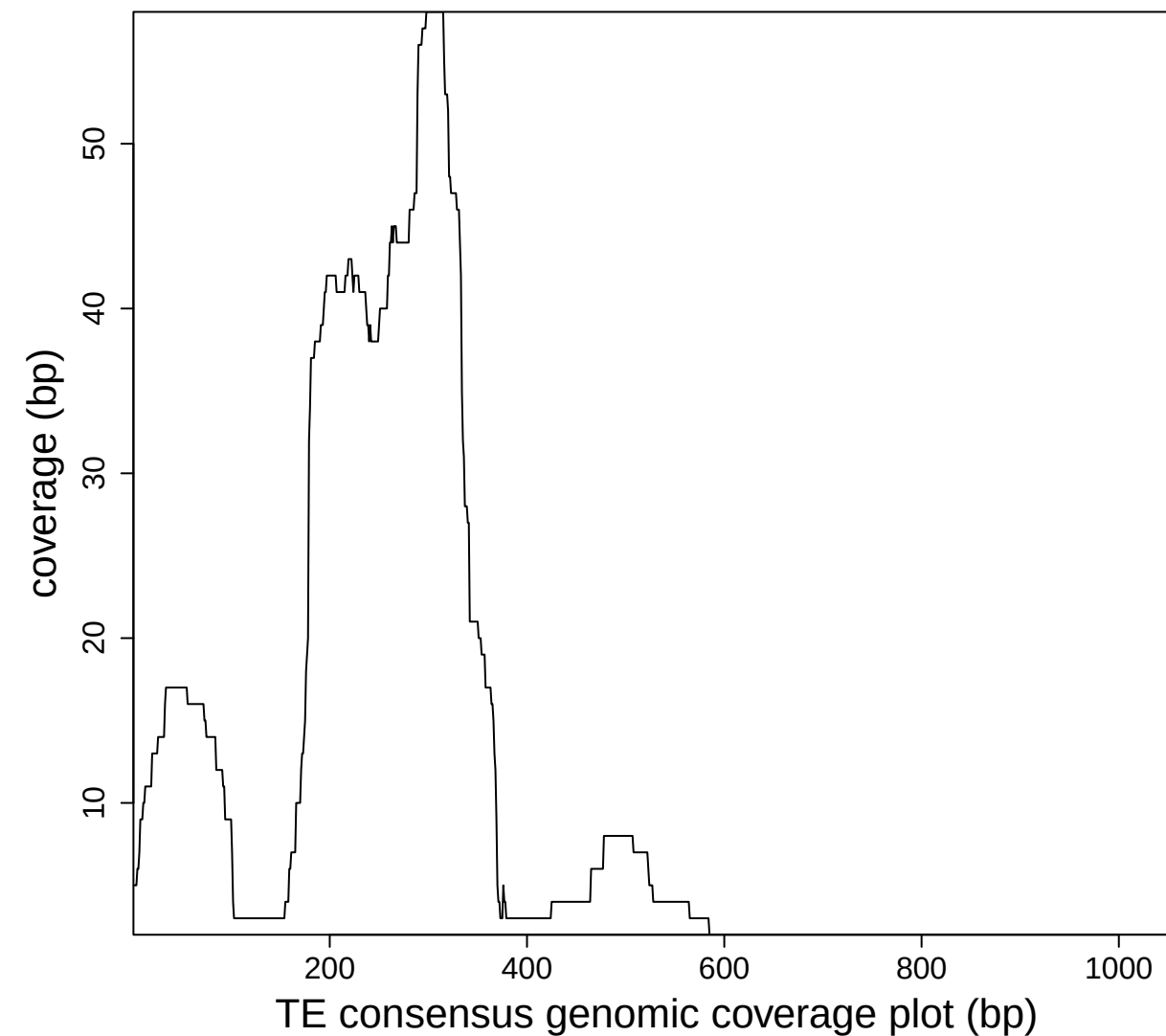
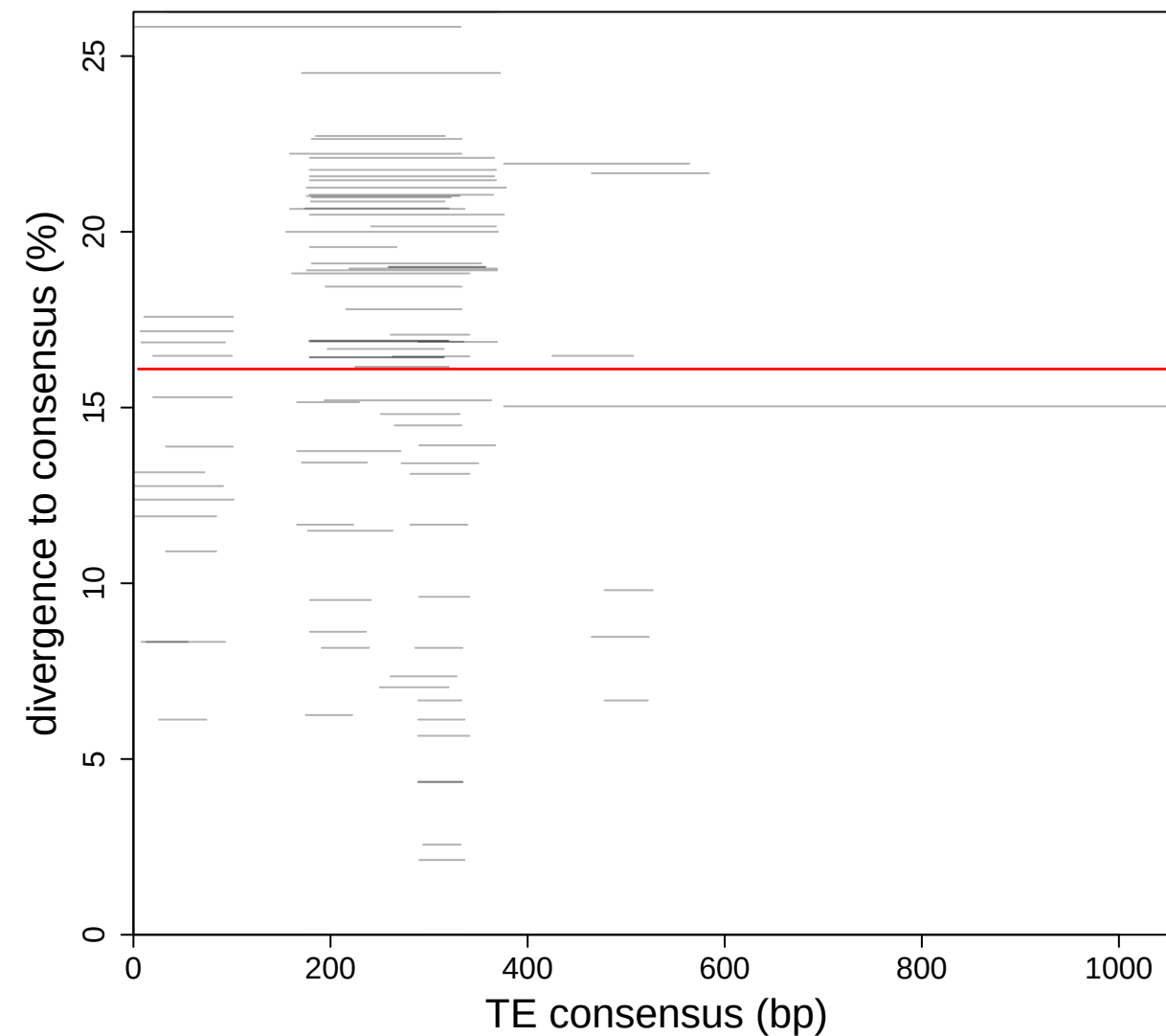
hily_42.fasta.b.bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa
size: 831bp; fragments: 21; full length: 0 (>=747.9bp)

After TEtrimmer Extended plot Blue lines are boundaries

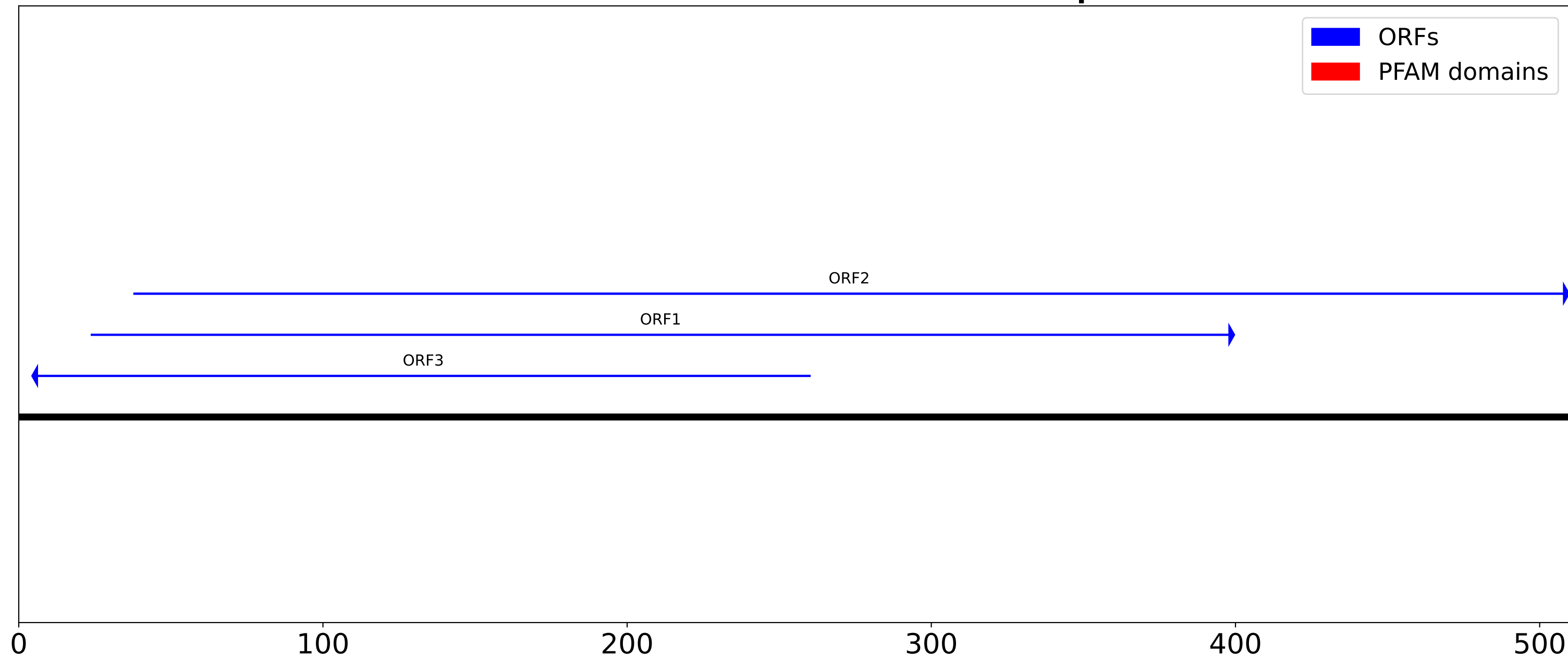


TE: rnd_5_family_42
size: 1062bp; fragments: 90; full length: 1 (>=955.8bp)

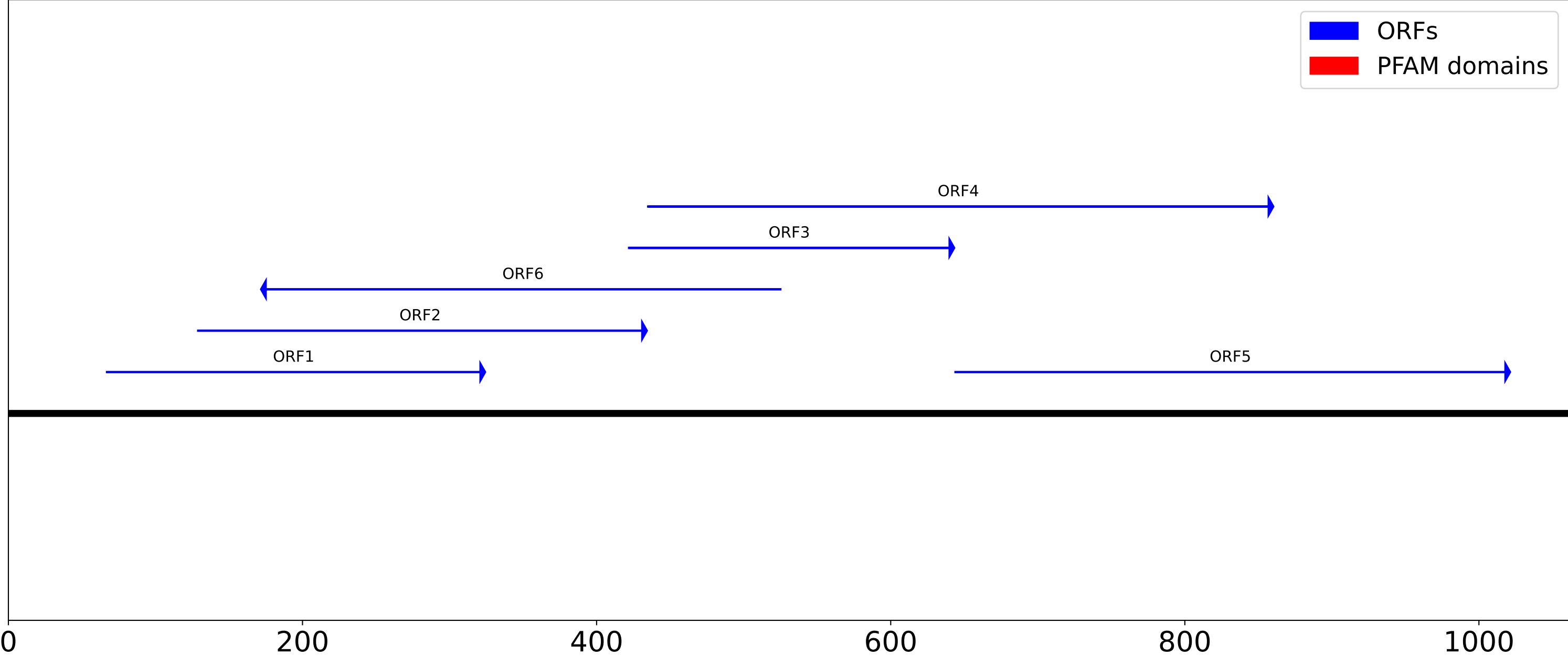
Before TEtrimmer 1062 bp



After TEtrimmer ORF and PFAM domain plot



Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

