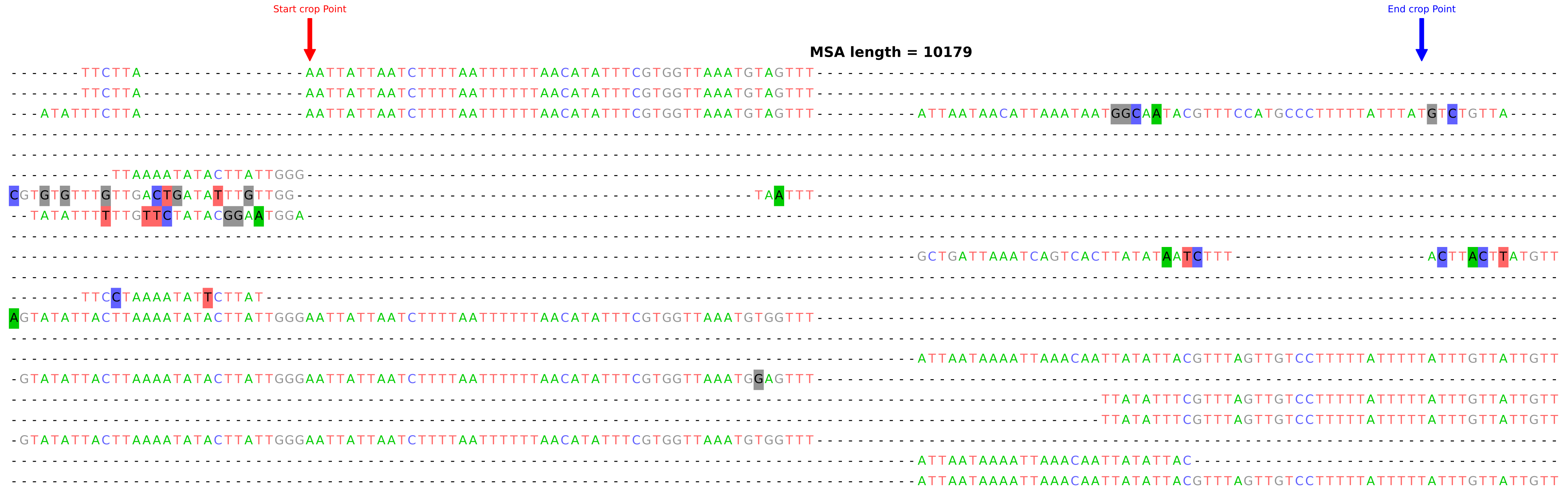
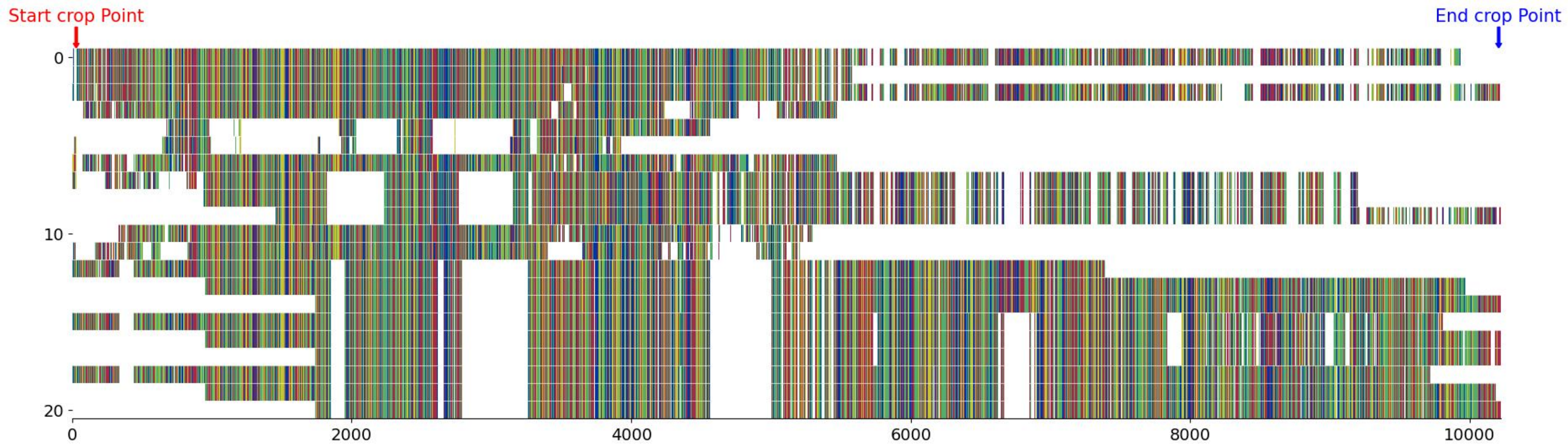
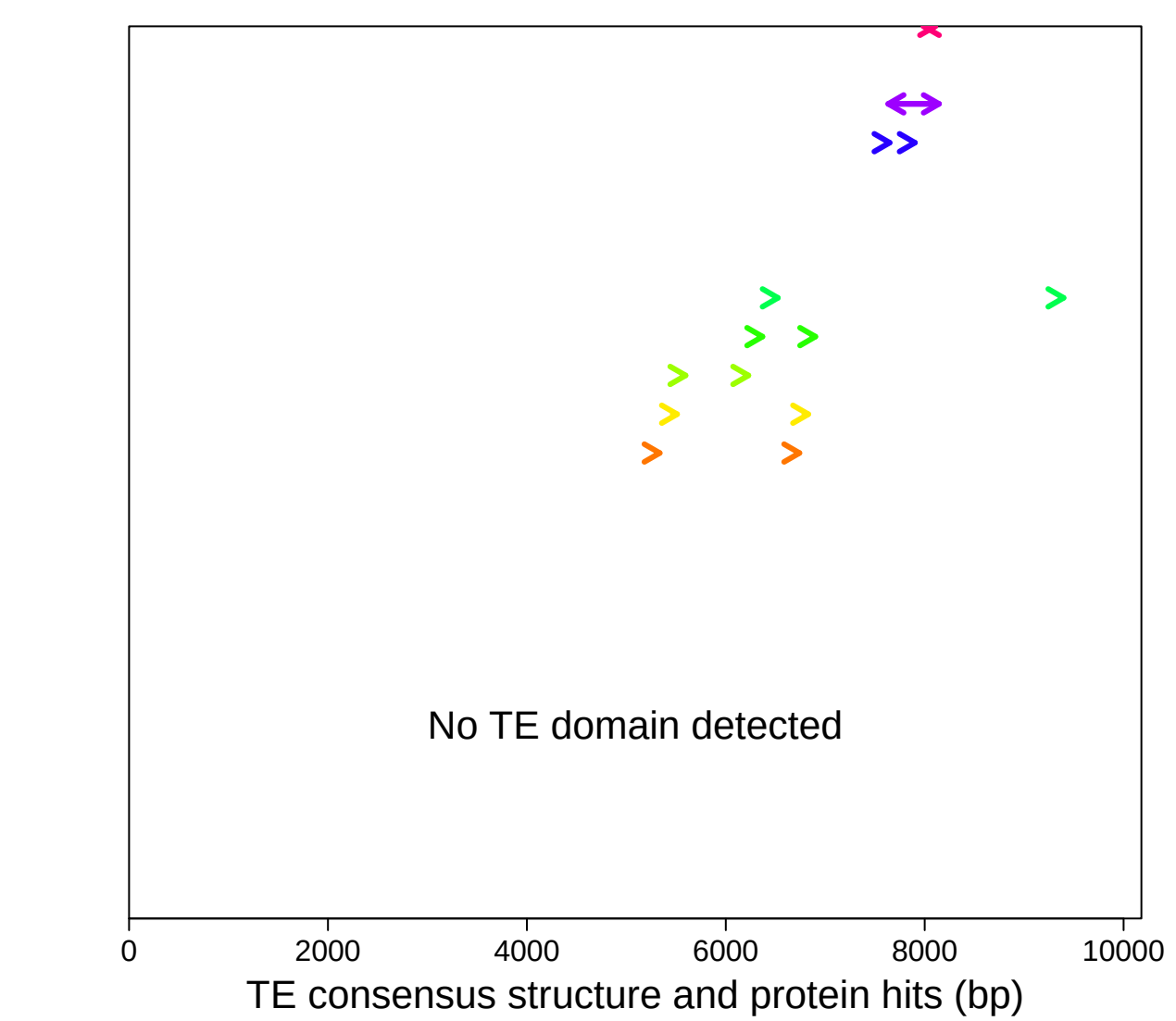
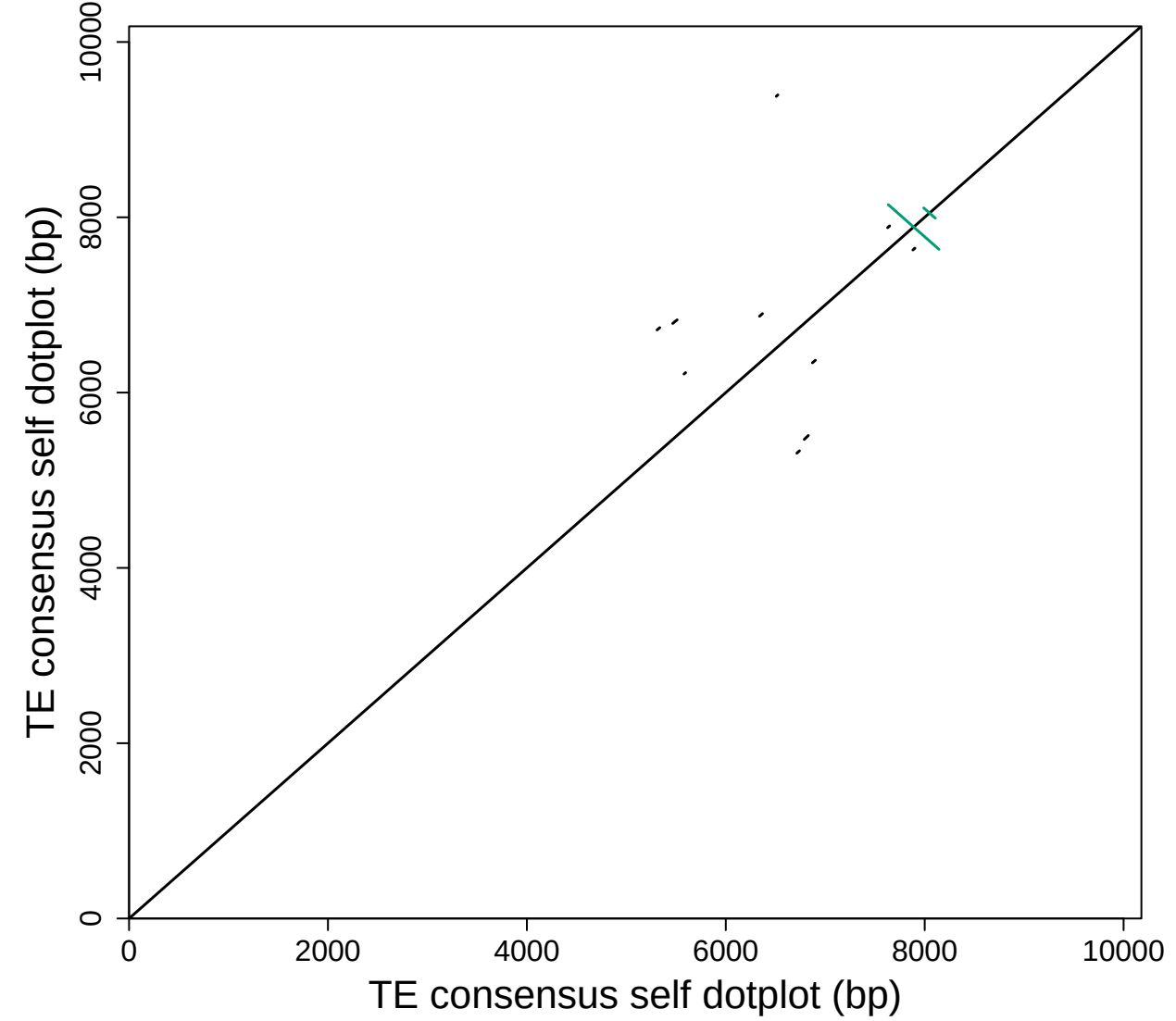
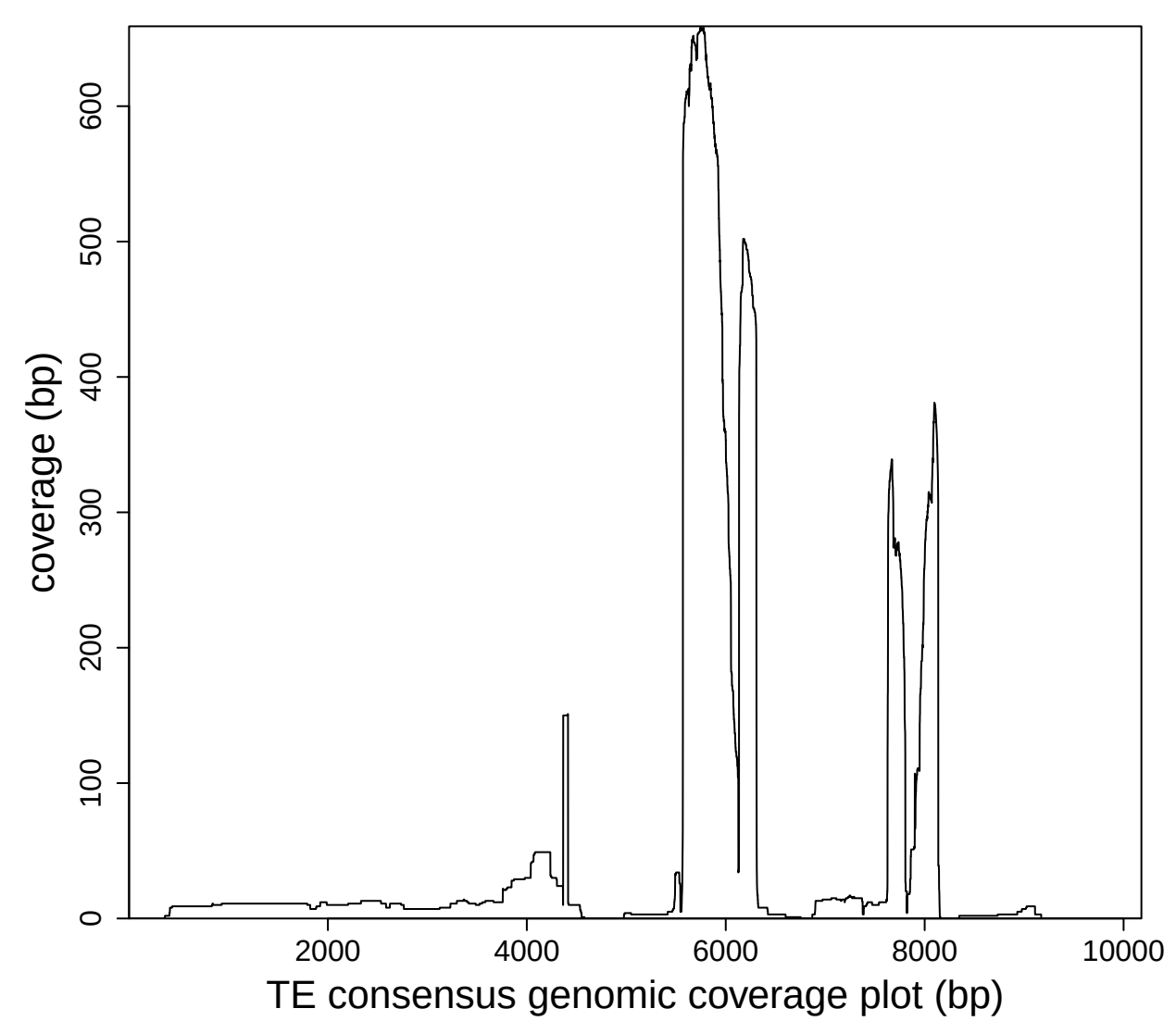
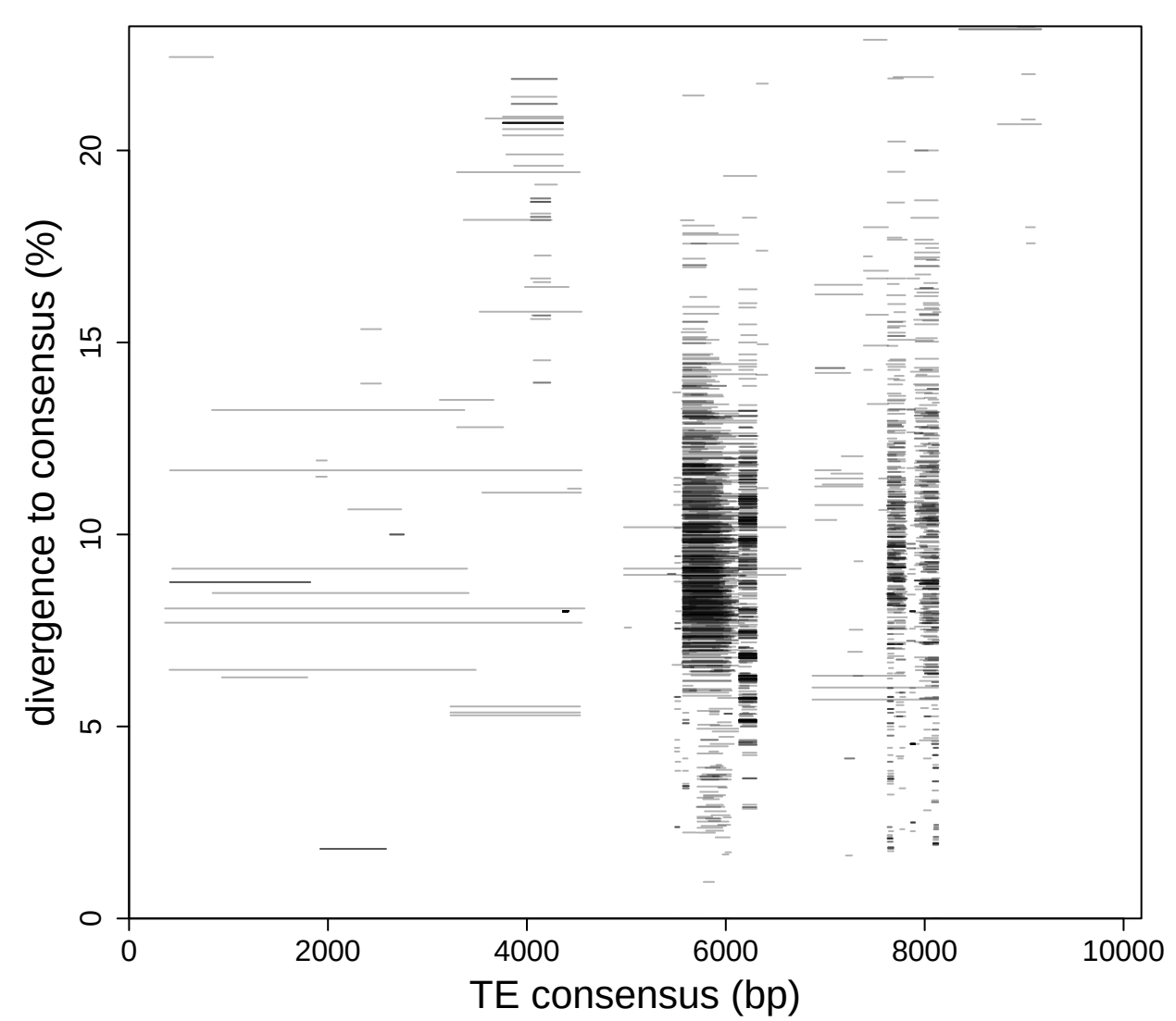






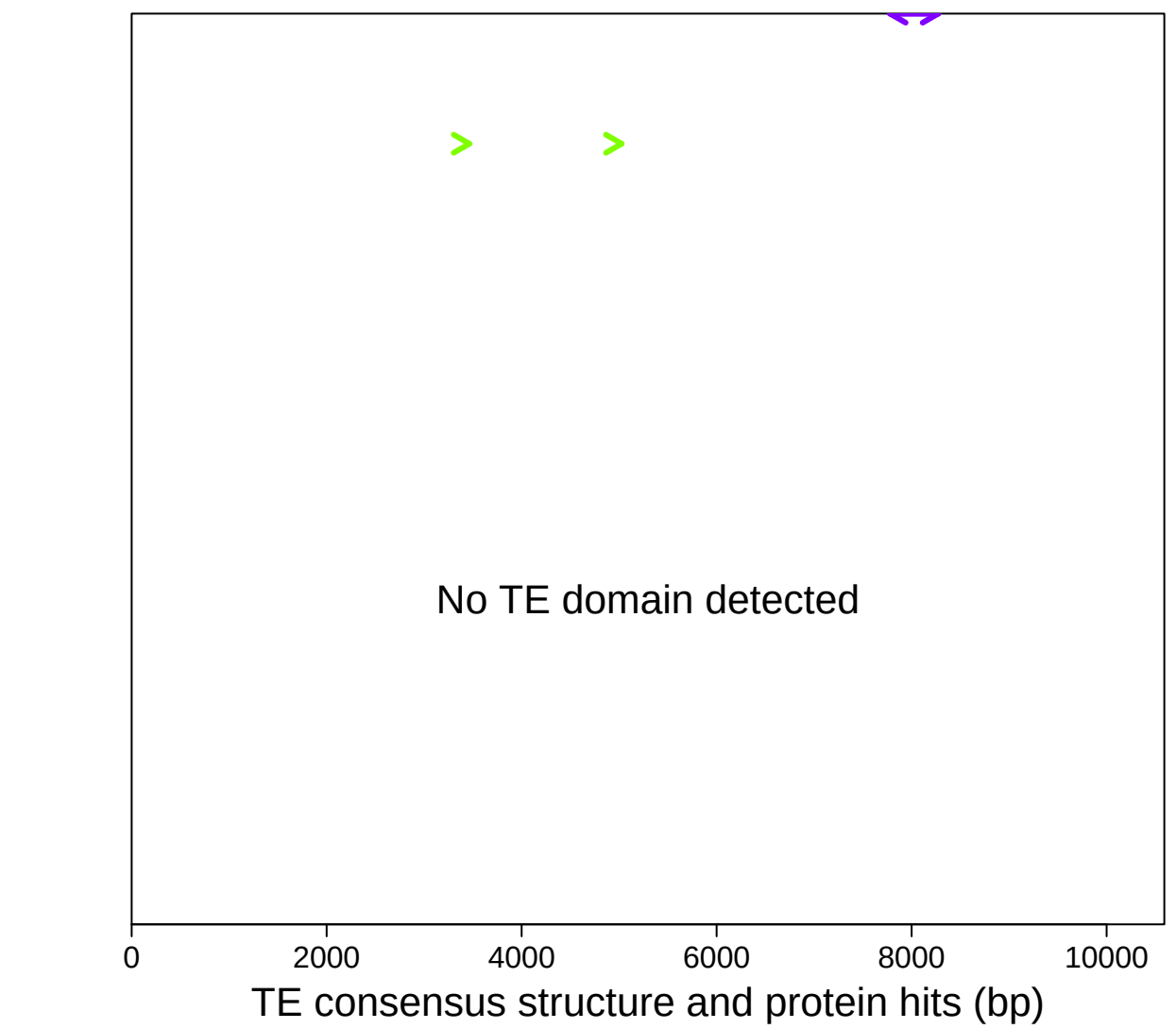
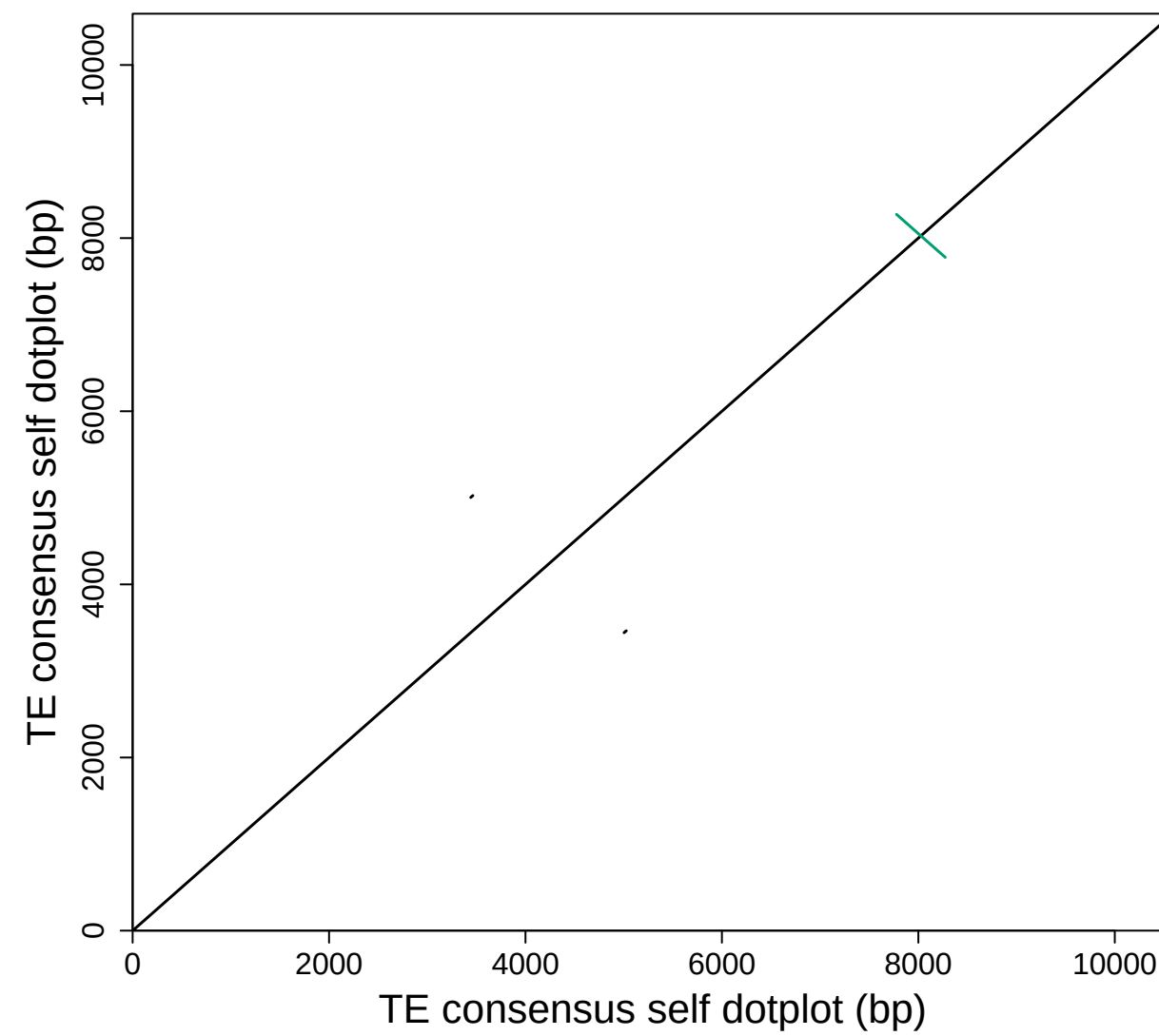
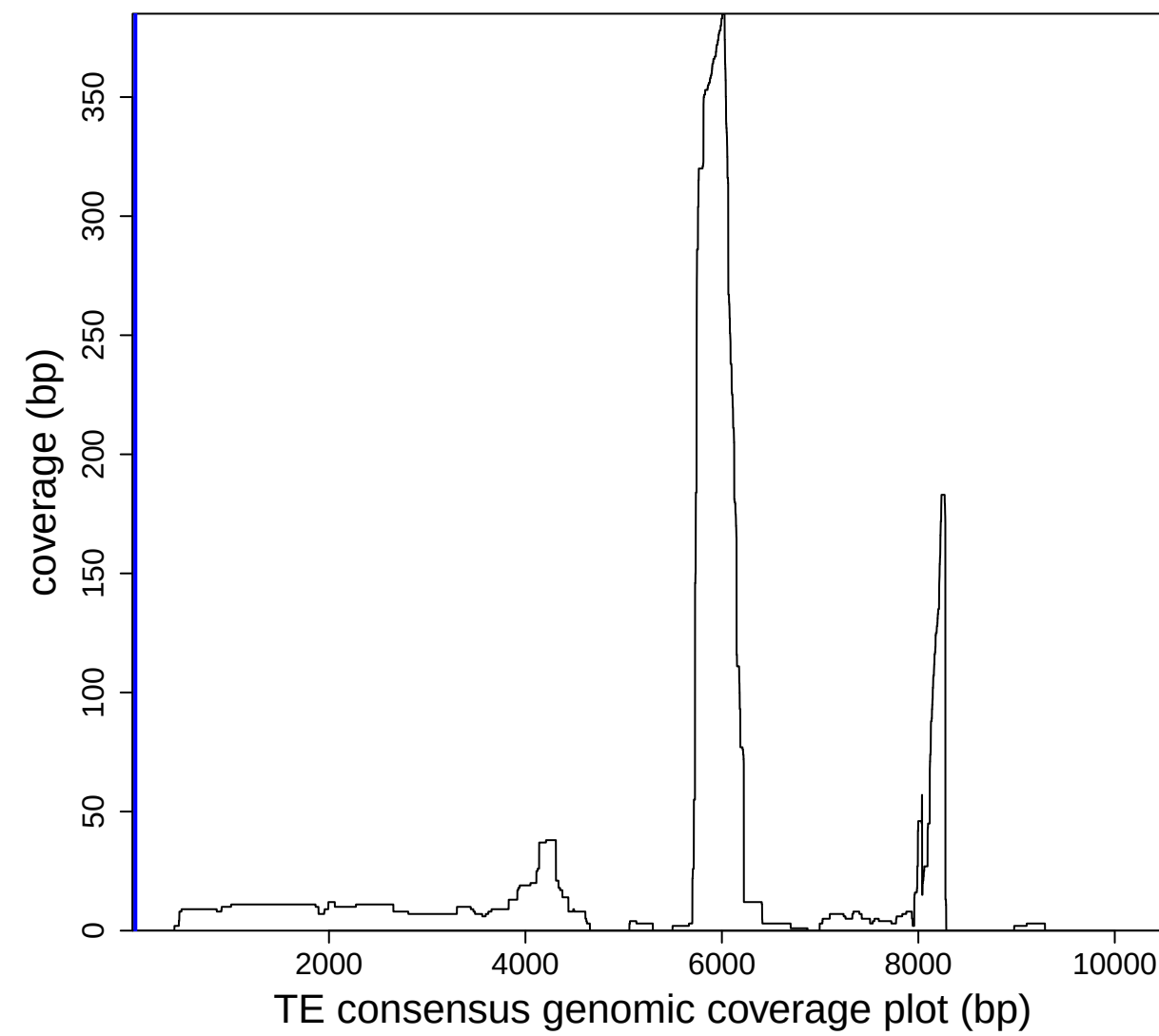
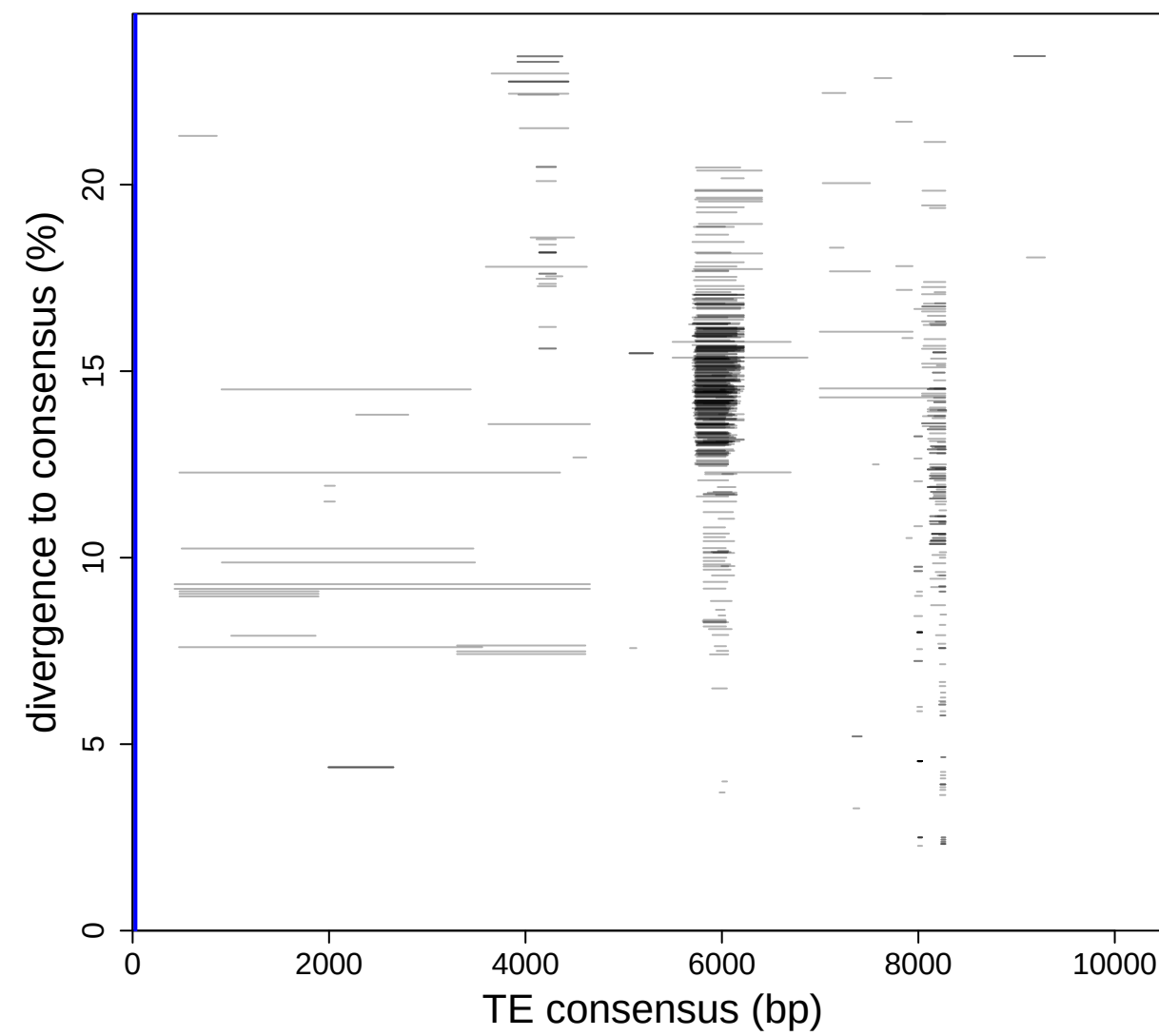
ta.b.bed uf.bed g_1.bed fm_1.bed 0_0 bcln.fa aln.fa cl.fa gs.f
size: 10179bp; fragments: 2400; full length: 0 (>=9161.1bp)

After TEtrimmer 10179 bp

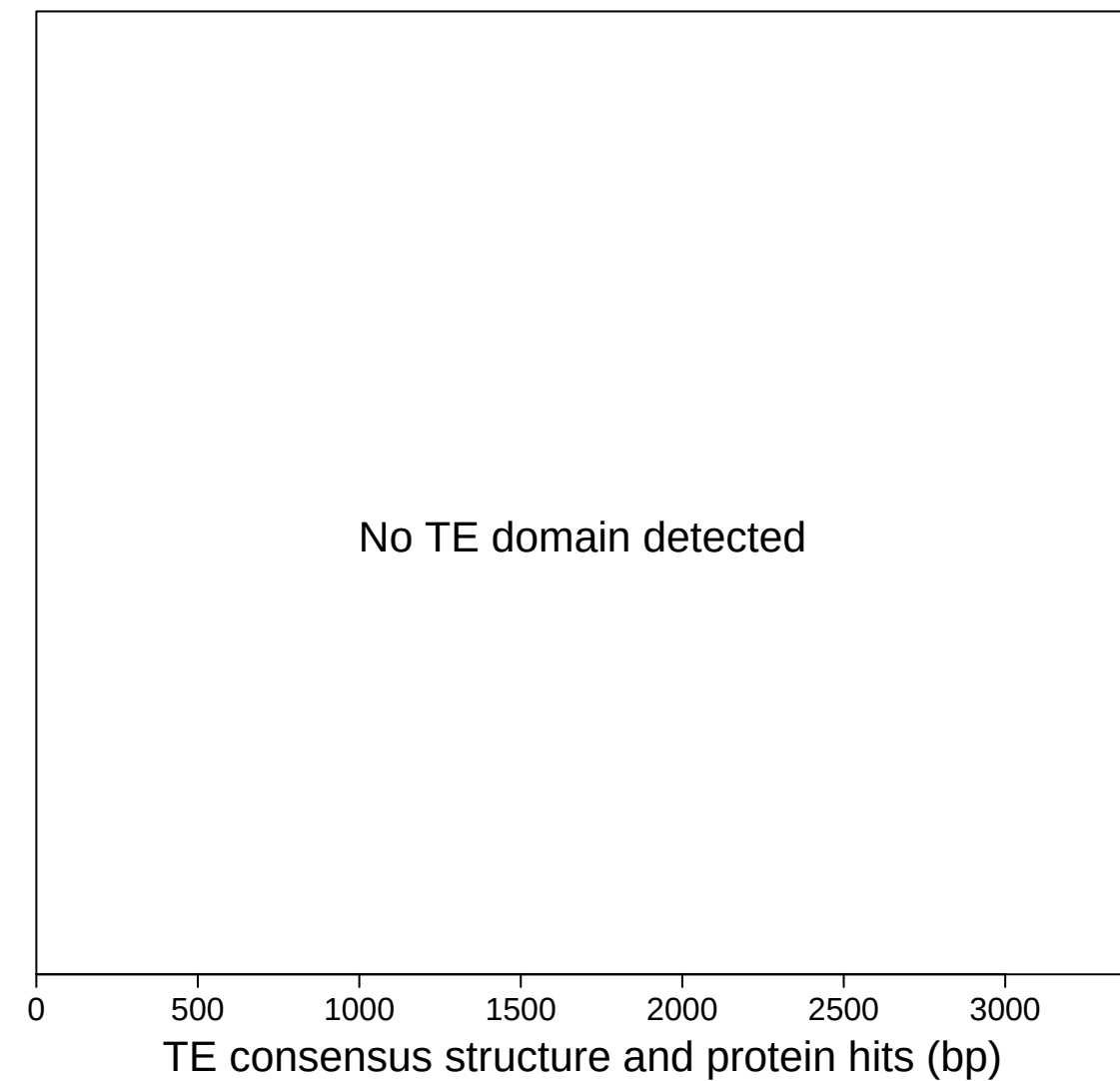
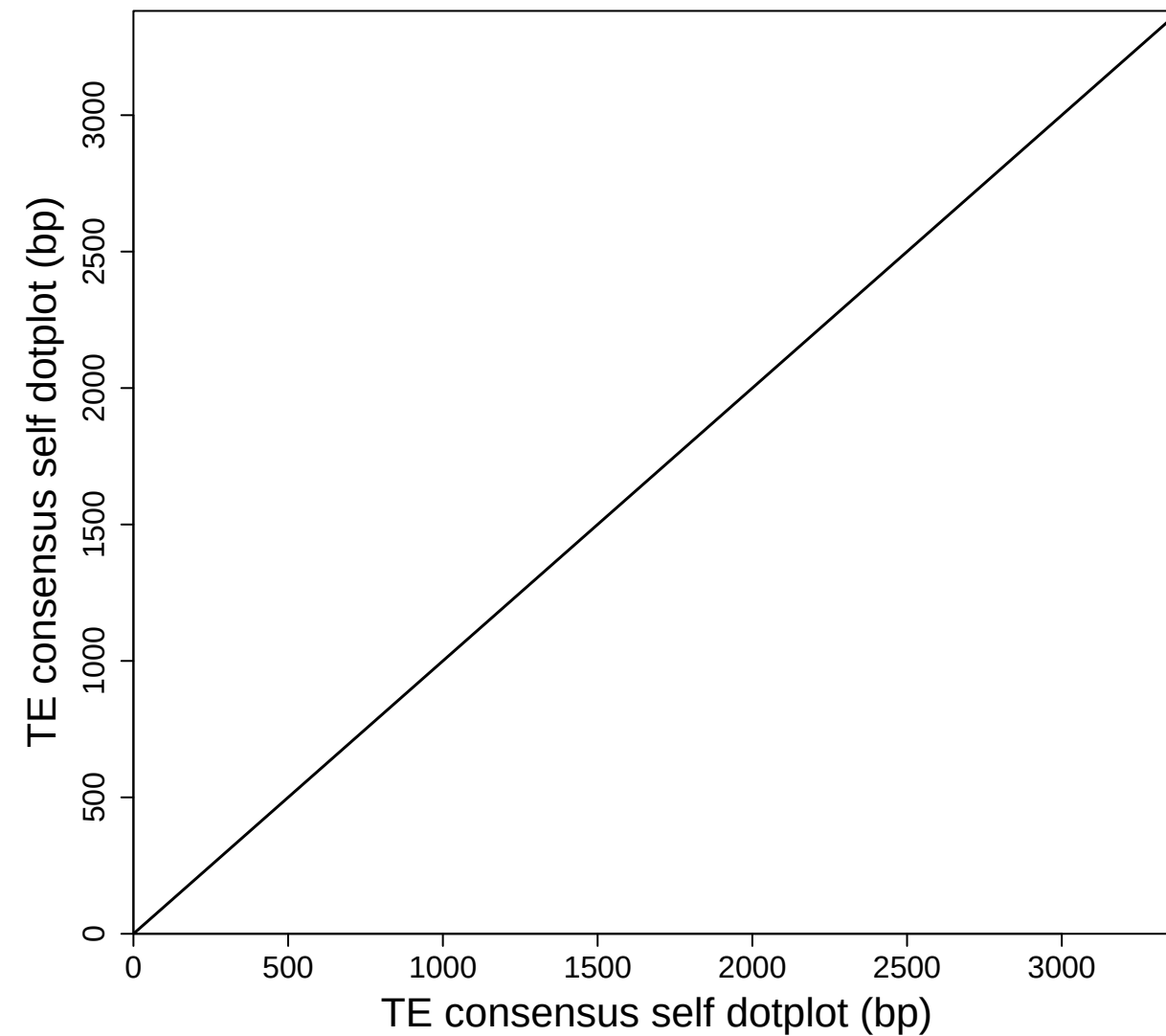
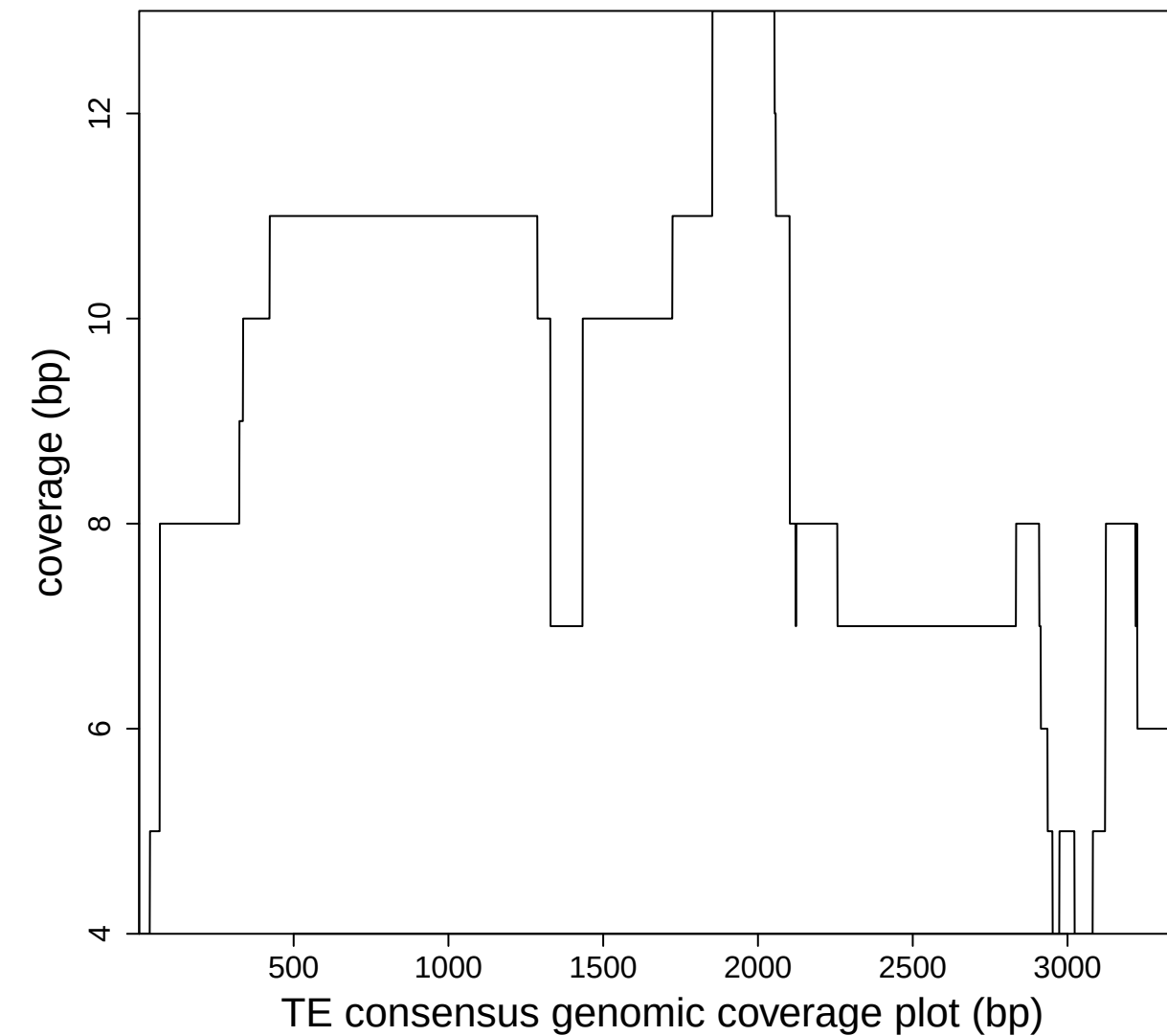
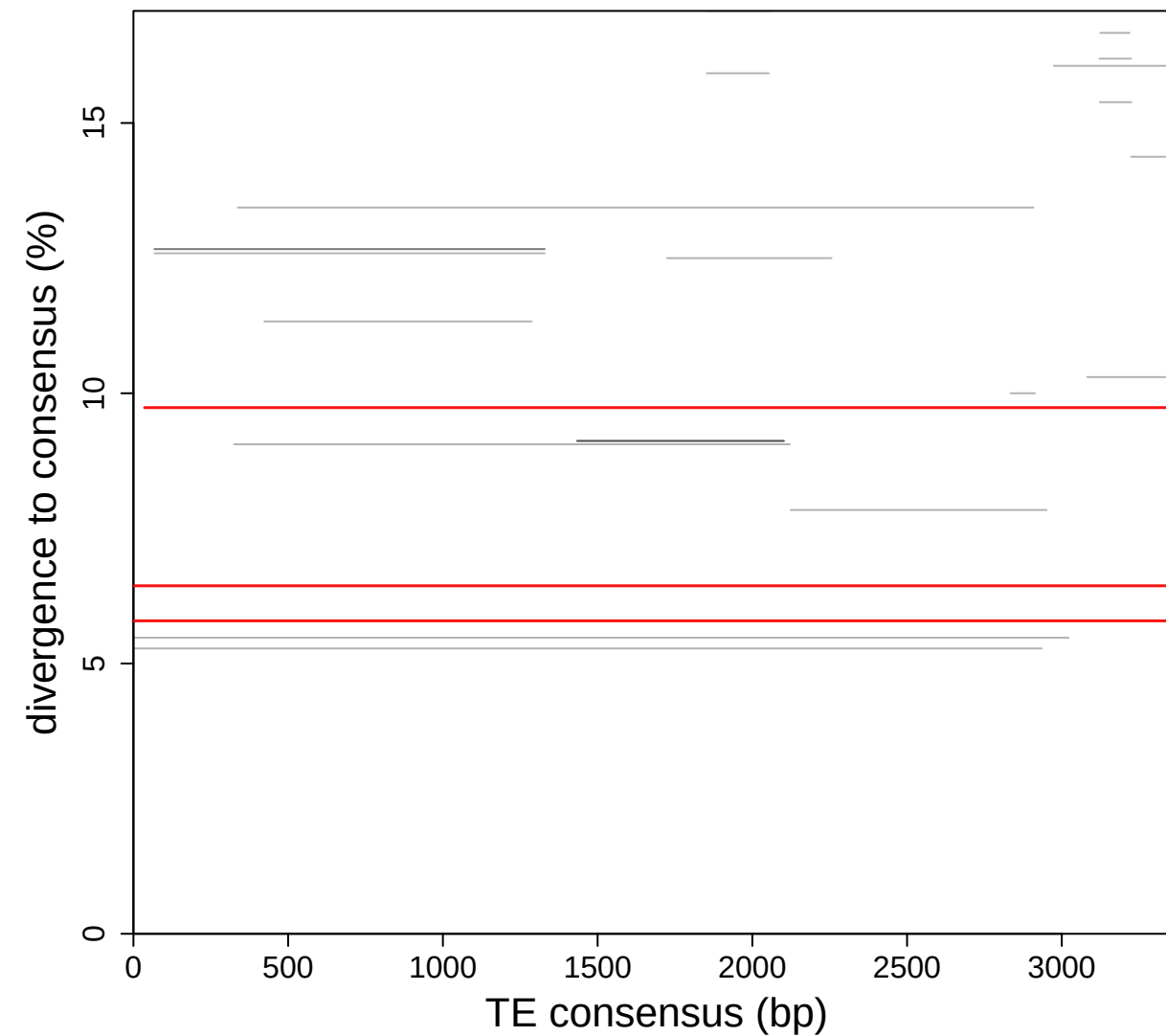


53.fasta.b.bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.1
size: 10592bp; fragments: 687; full length: 0 (>=9532.8bp)

After TEtrimmer Extended plot Blue lines are boundaries

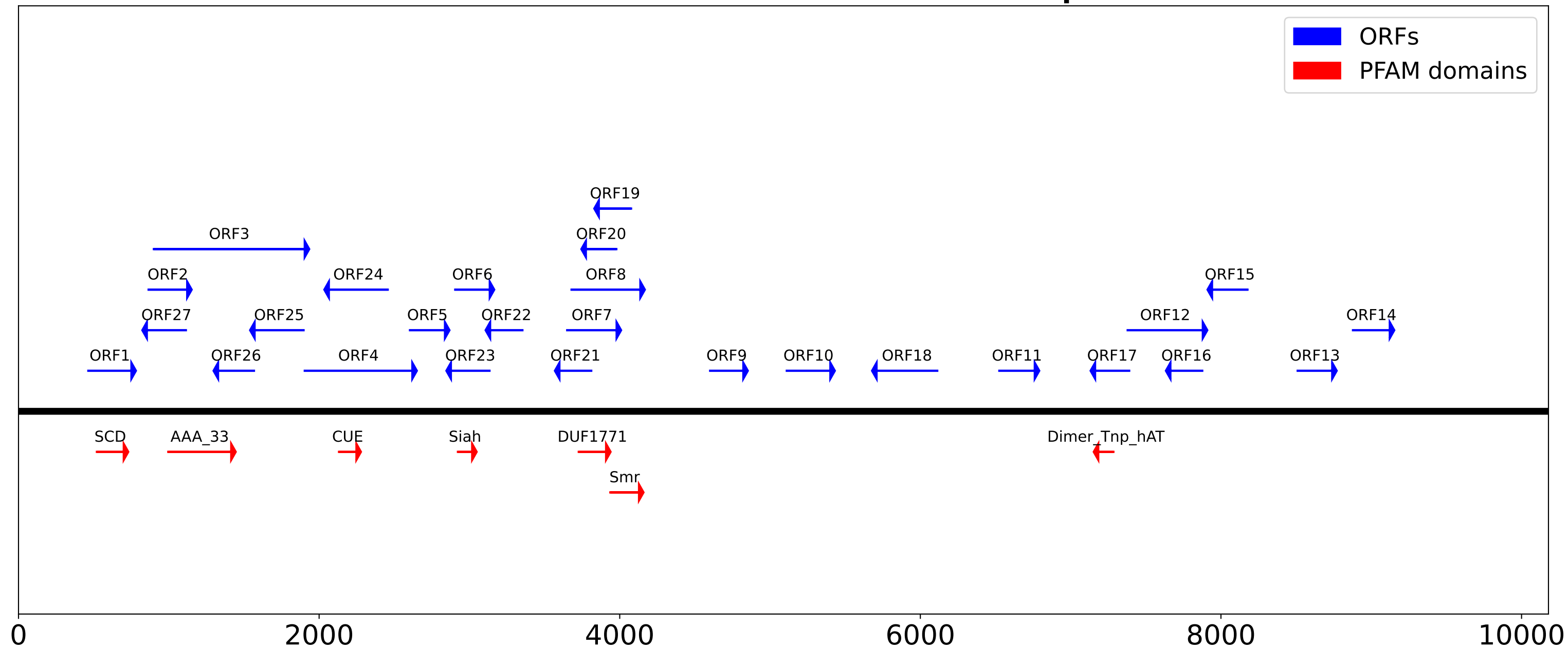


TE: rnd_5_family_13853
size: 3382bp; fragments: 25; full length: 3 (≥ 3043.8 bp)

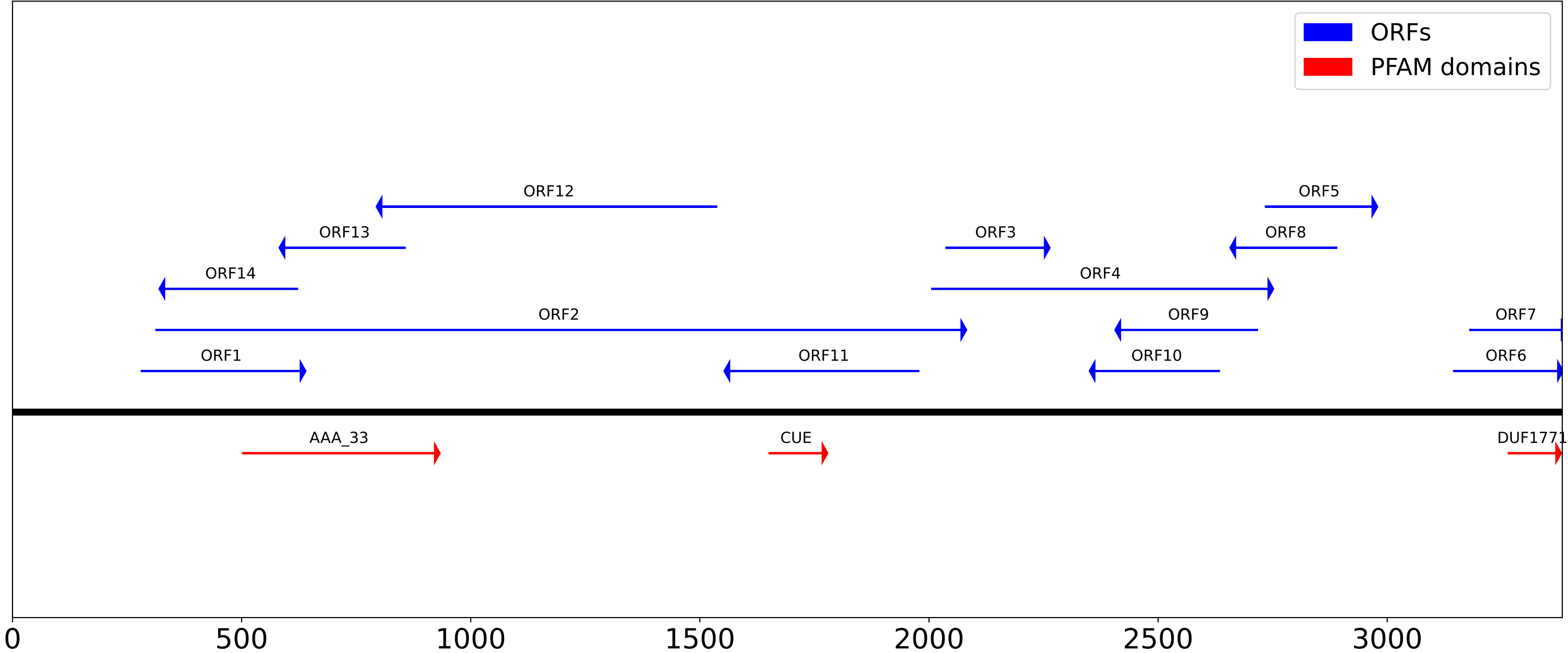


Before TEtrimmer 3382 bp

After TEtrimmer ORF and PFAM domain plot



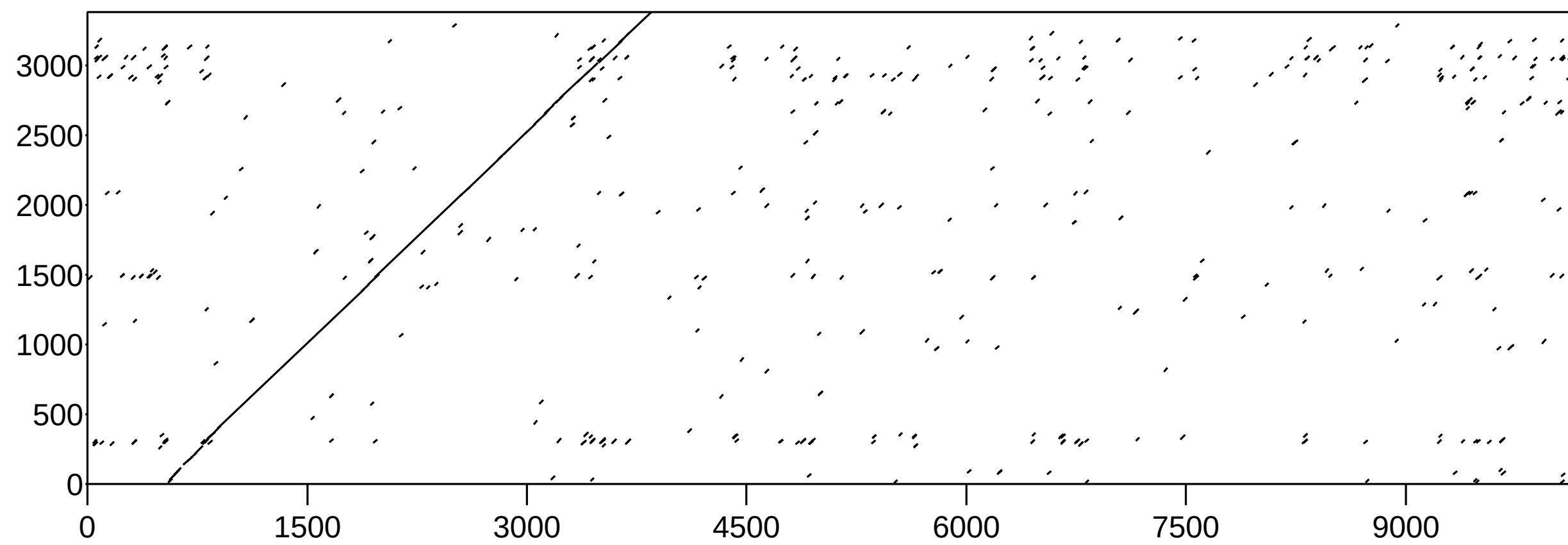
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)