

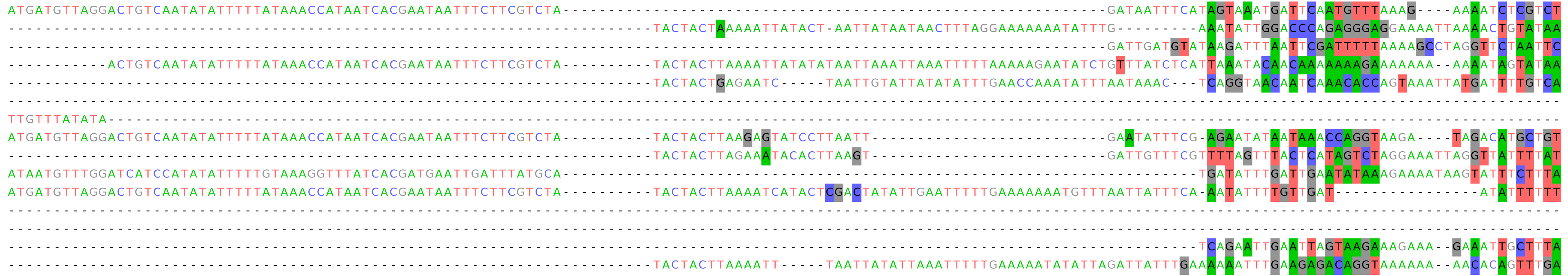
Start crop Point

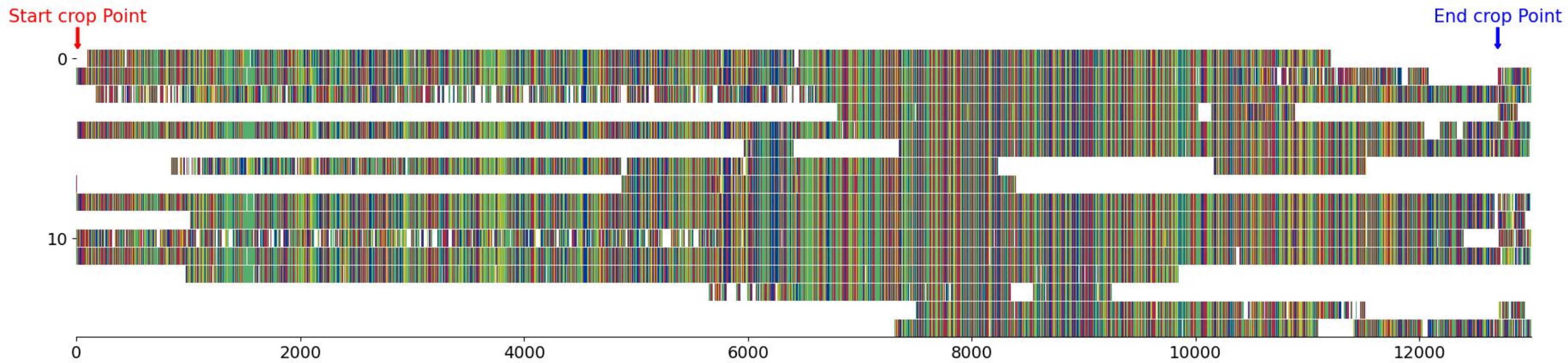


End crop Point



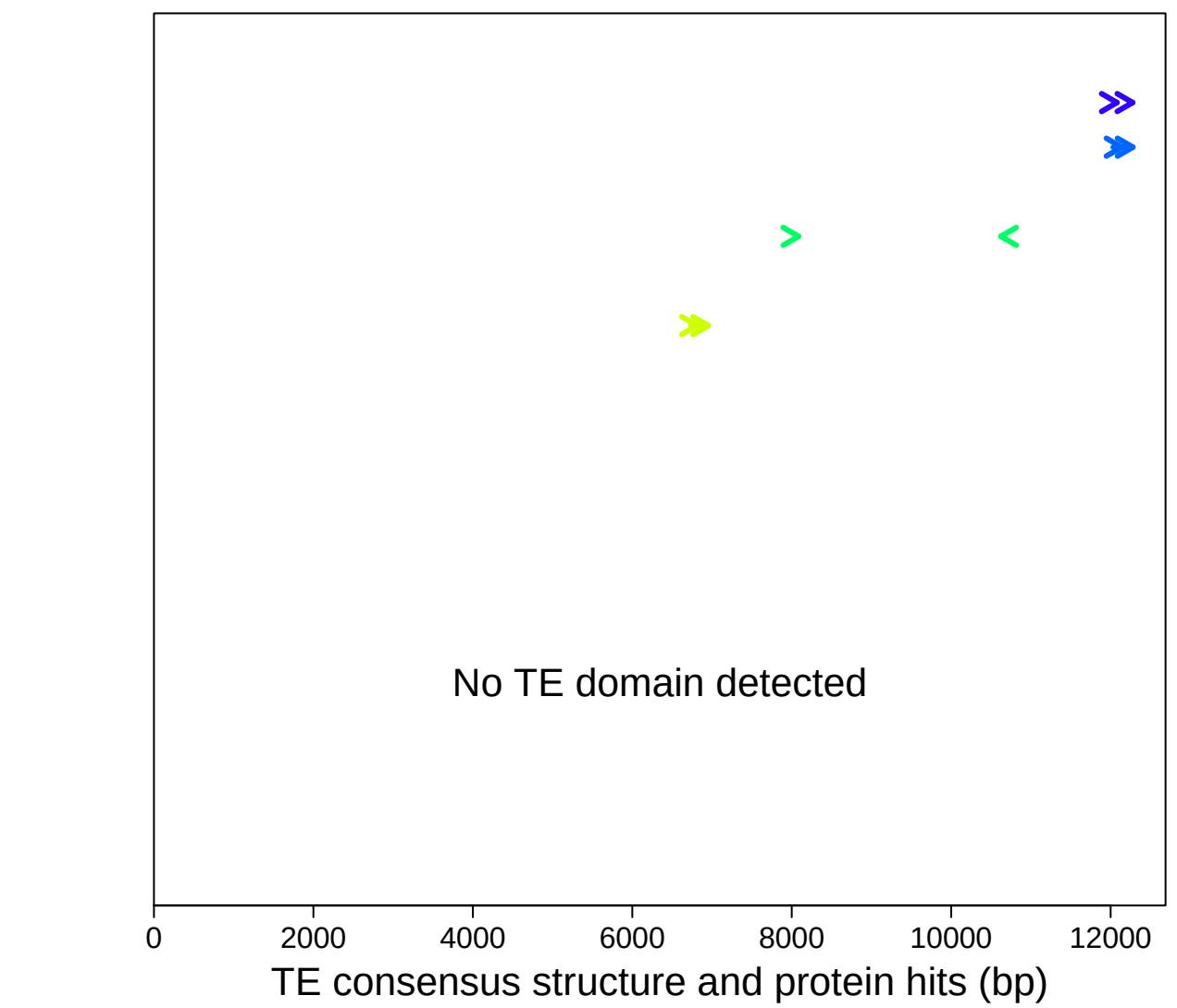
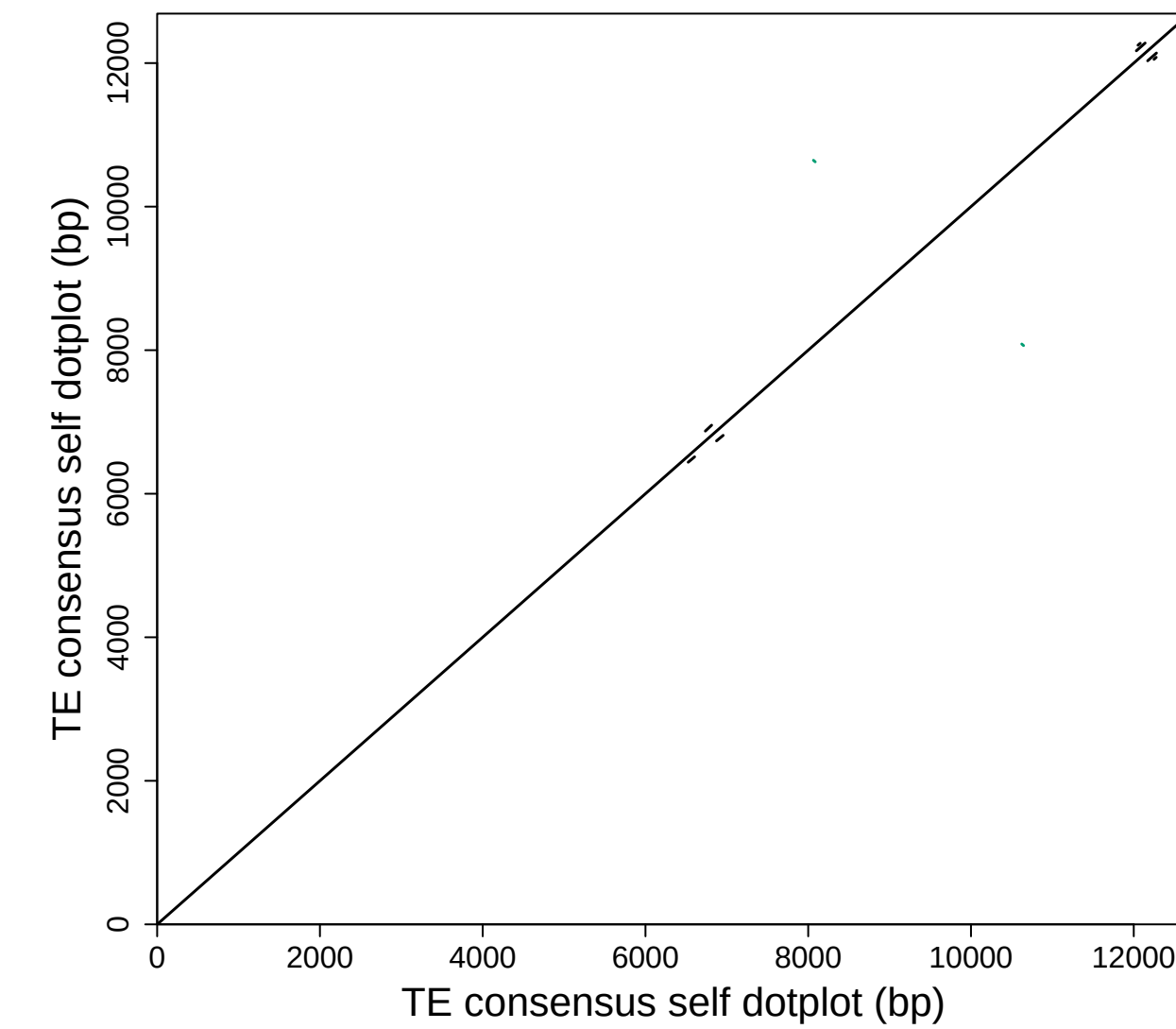
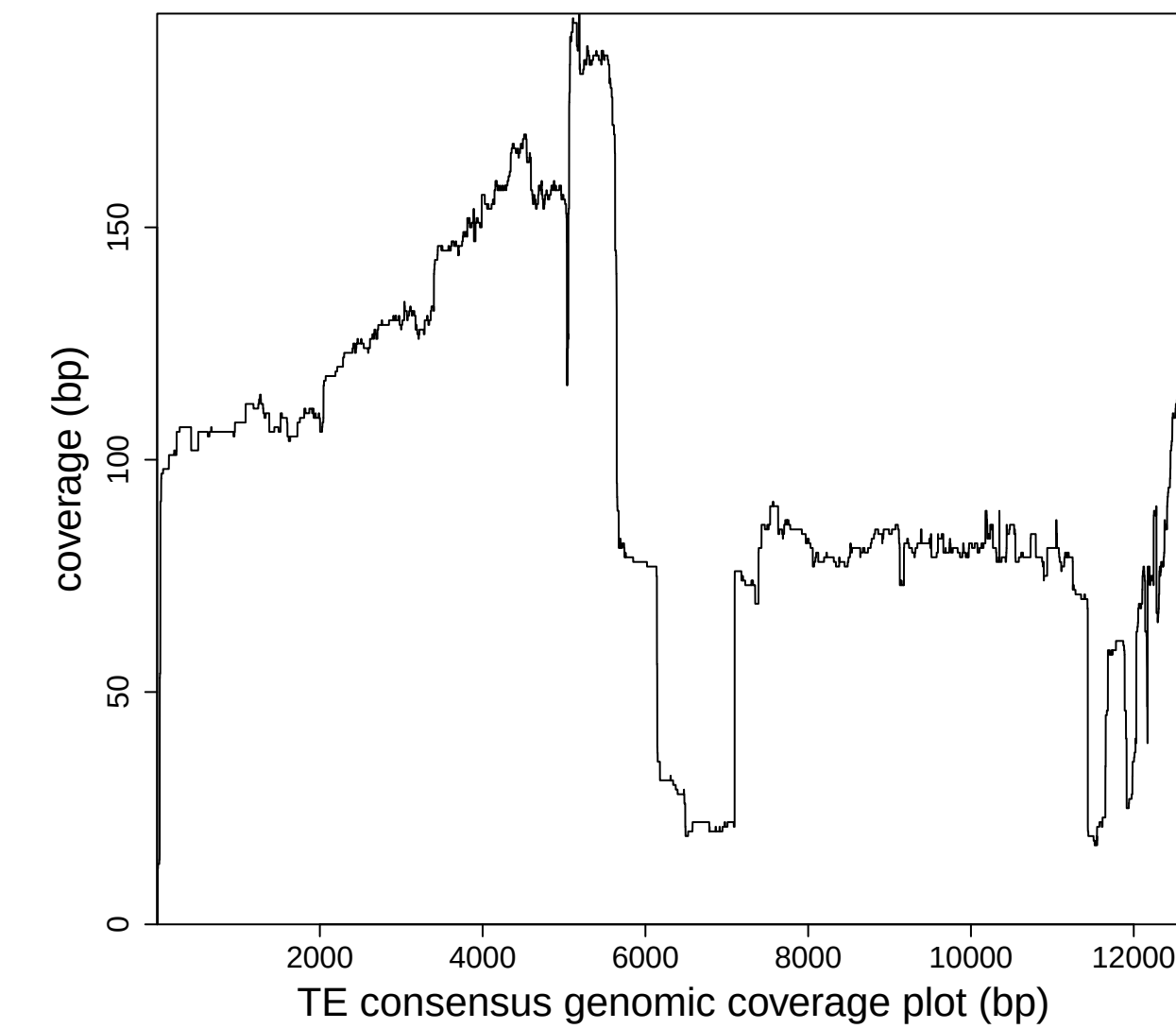
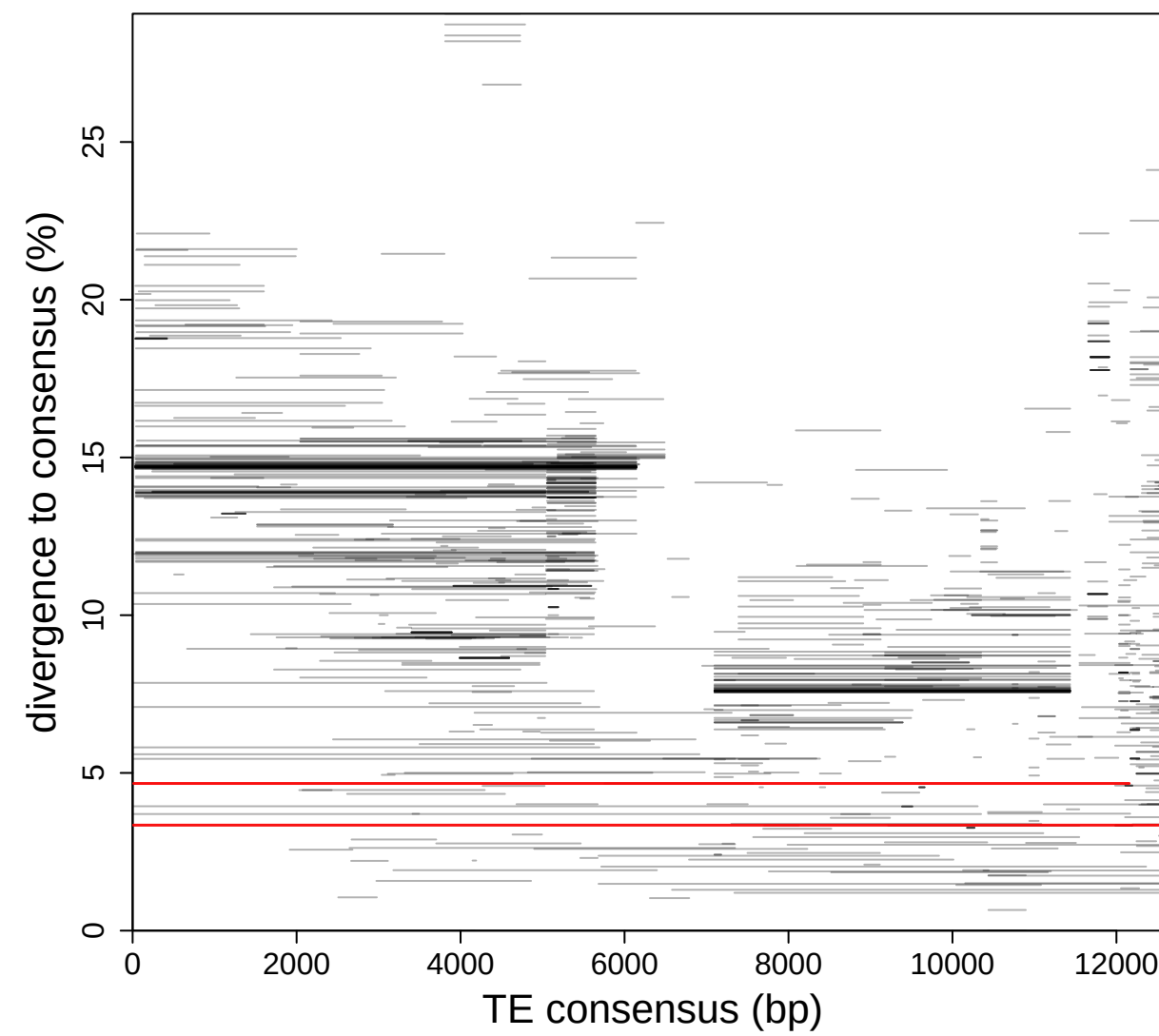
MSA length = 12689





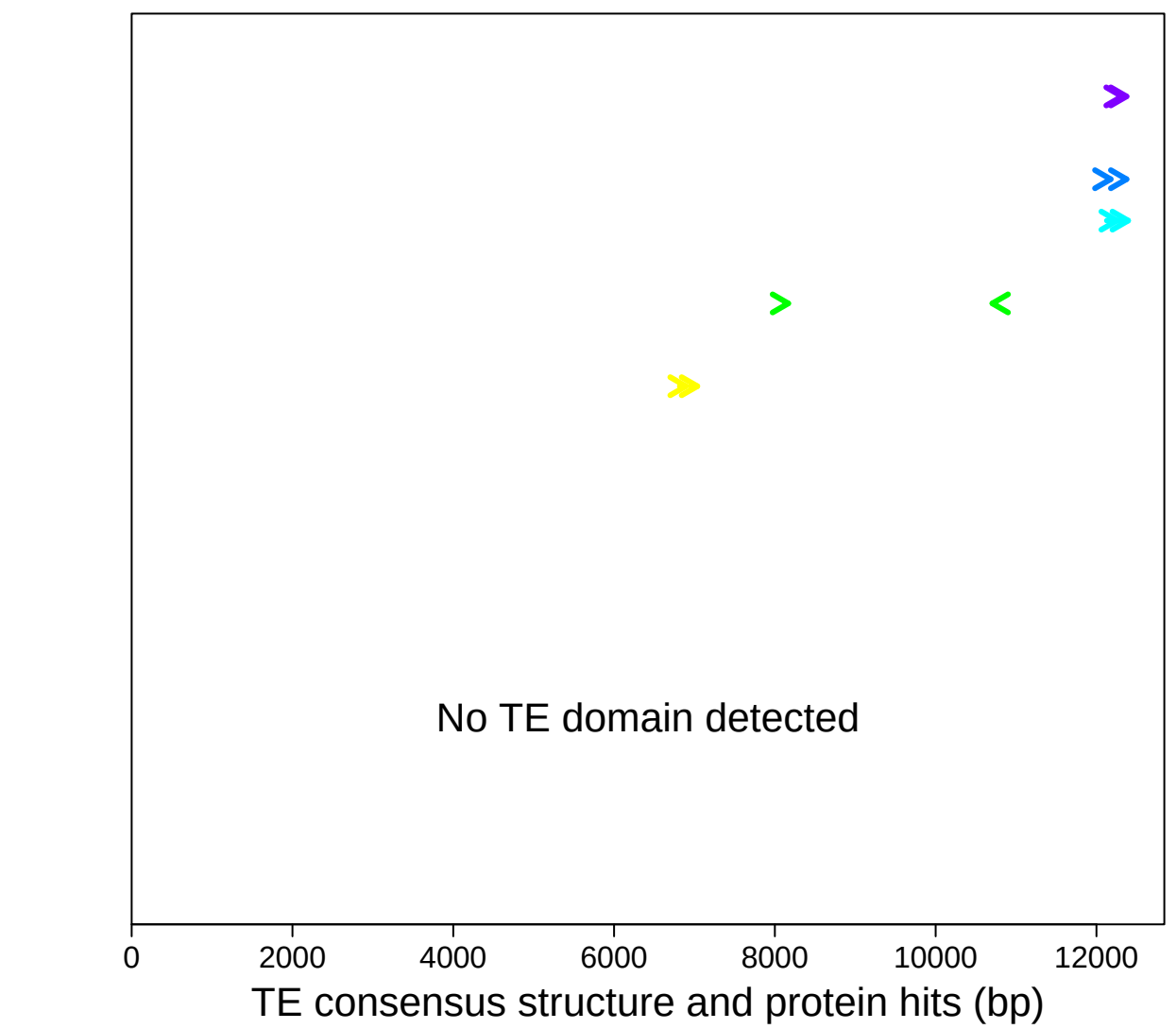
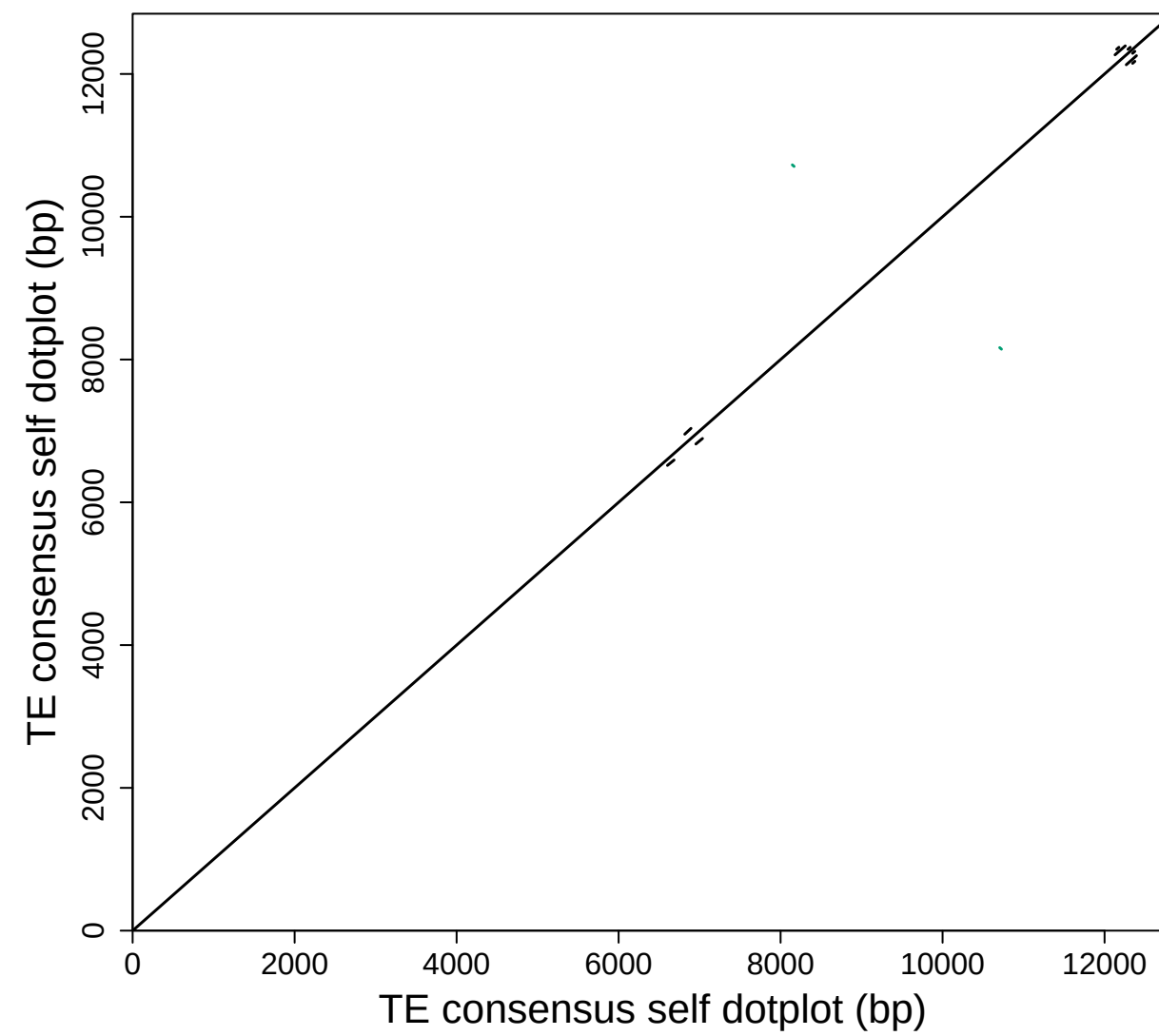
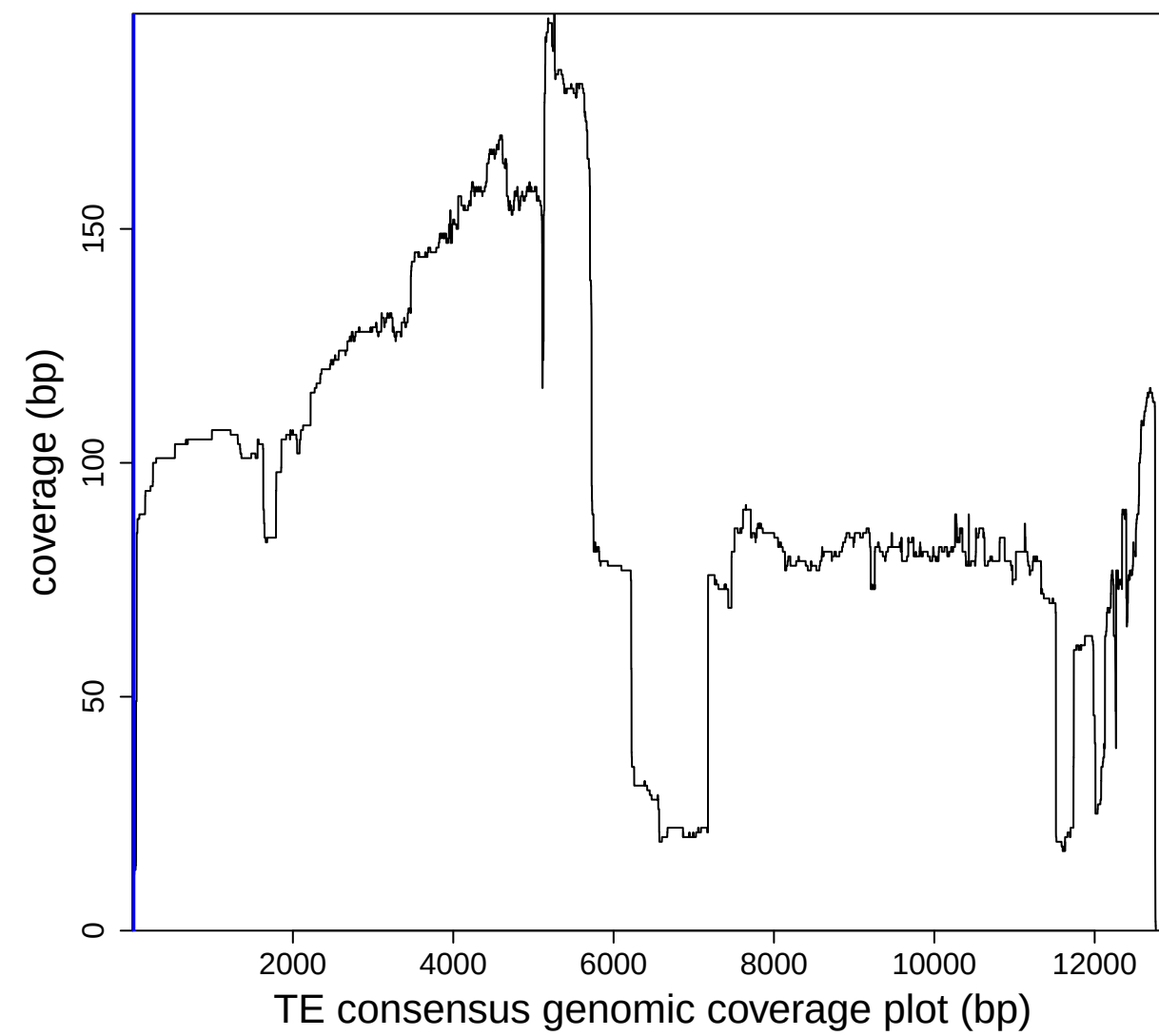
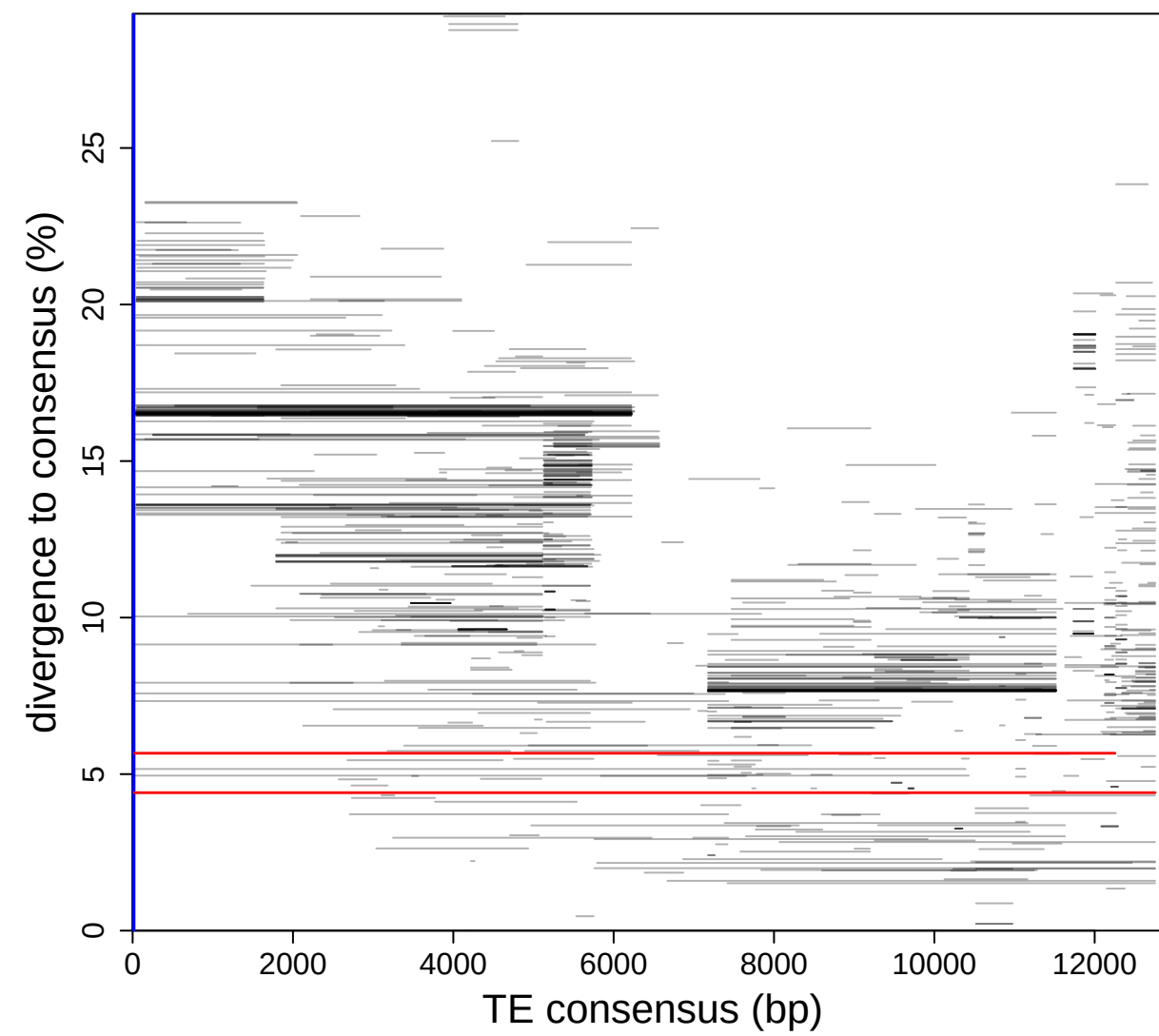
a.b.bed_uf.bed_g_3.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.f
size: 12689bp; fragments: 892; full length: 2 (>=11420.1bp)

After TEtrimmer 12689 bp



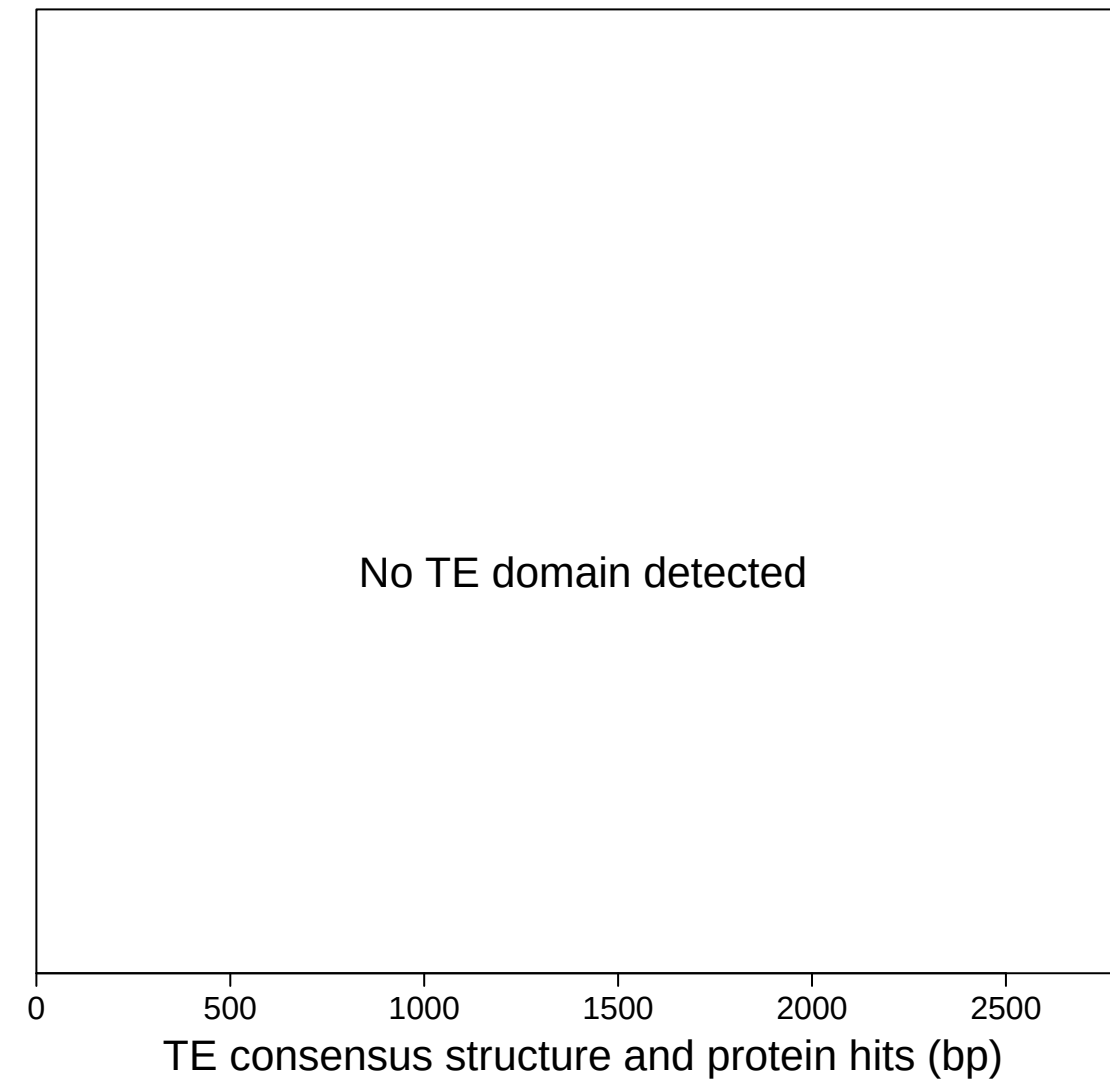
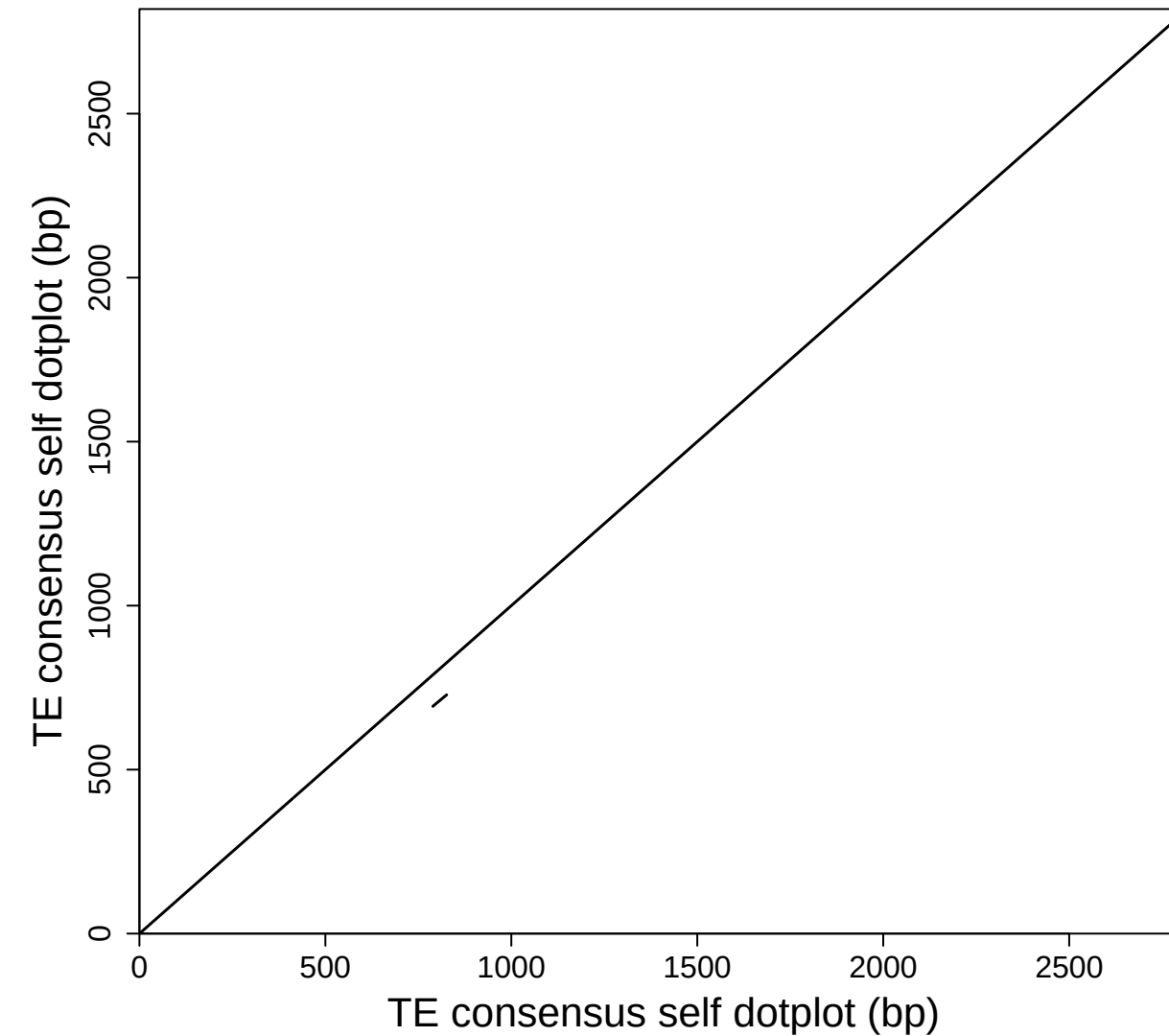
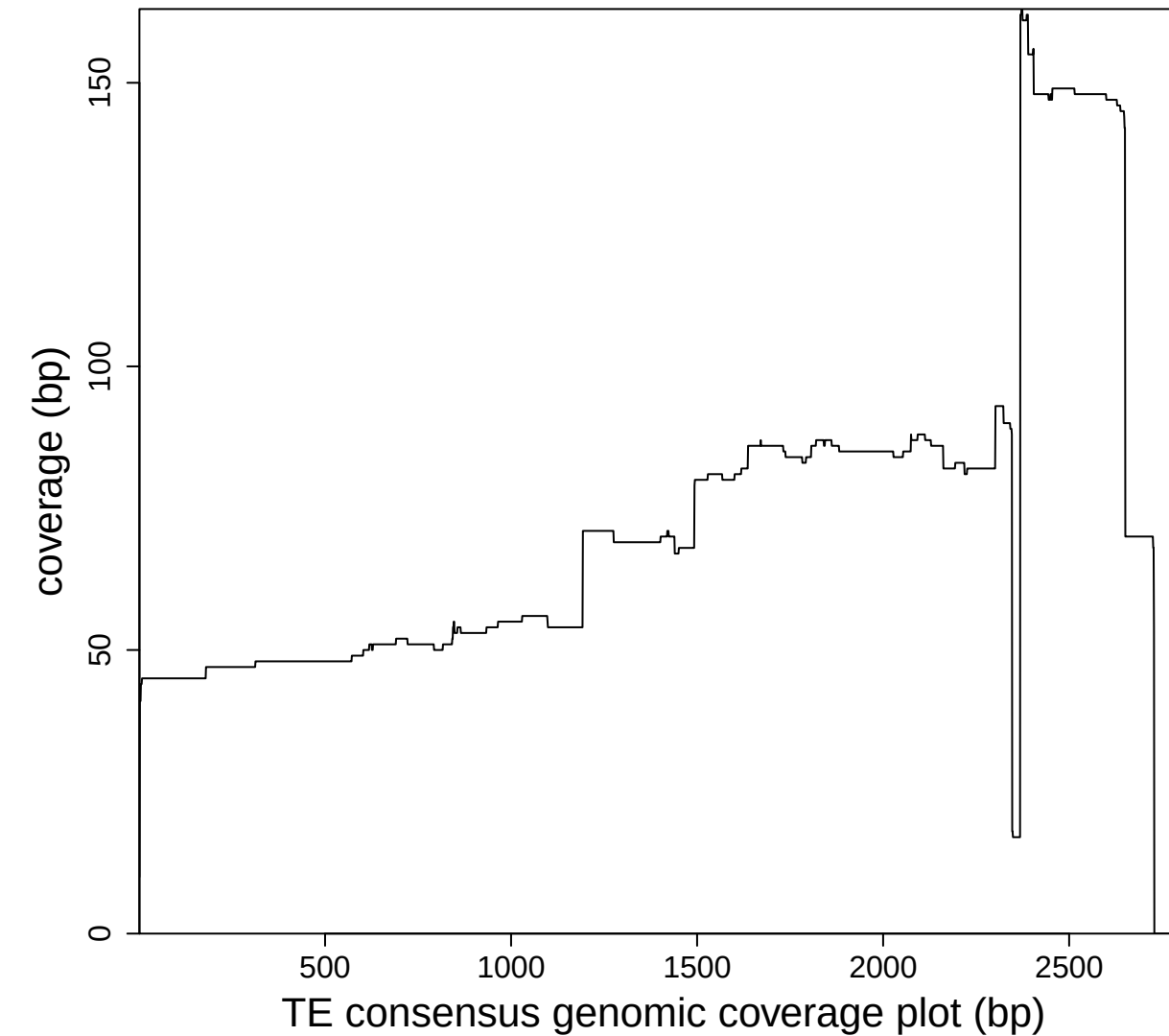
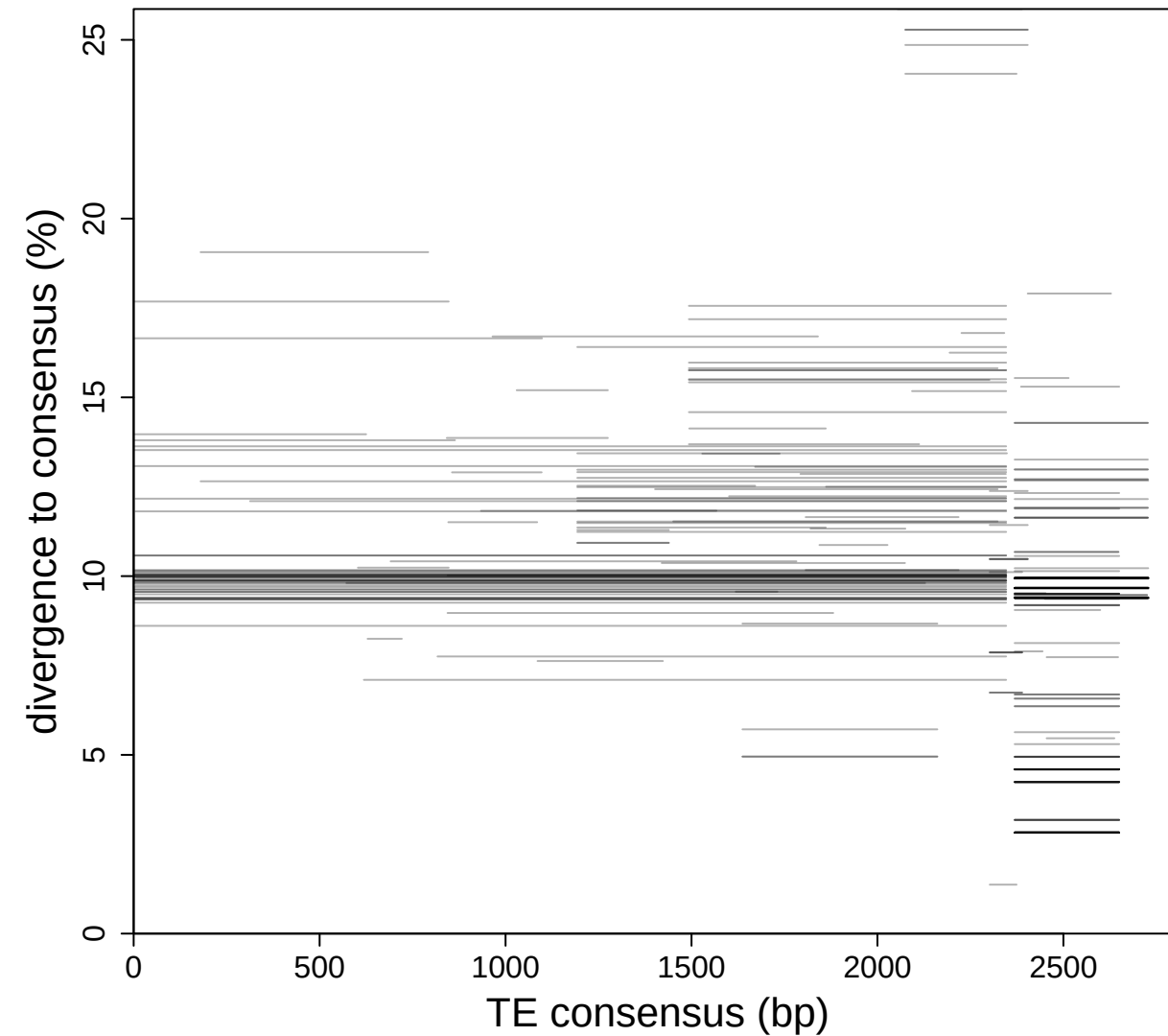
4.fasta.b.bed uf.bed g_3.bed fm_1.bed 0 0 bcIn.fa aln.fa cl.f
size: 12976bp; fragments: 880; full length: 2 (>=11678.4bp)

After TEtrimmer Extended plot Blue lines are boundaries

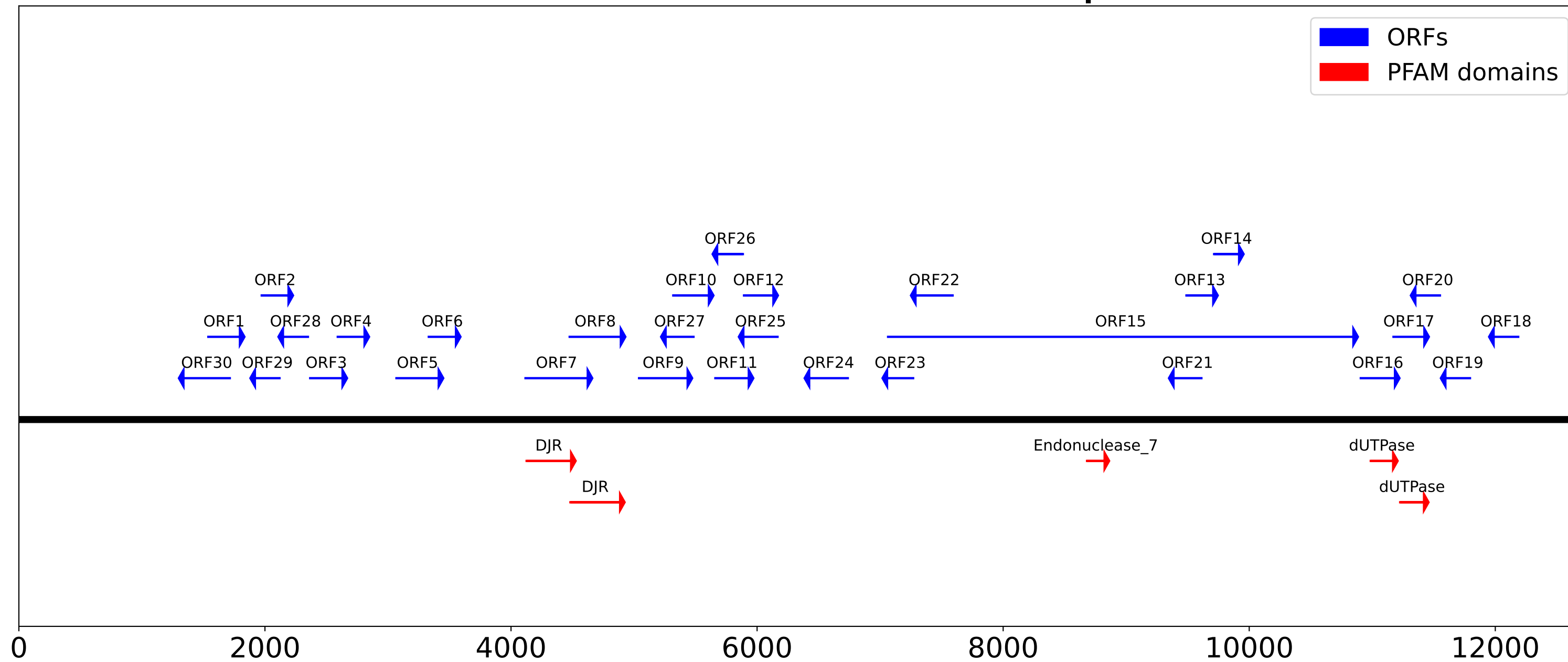


TE: rnd_5_family_4464
size: 2819bp; fragments: 280; full length: 0 (>=2537.1bp)

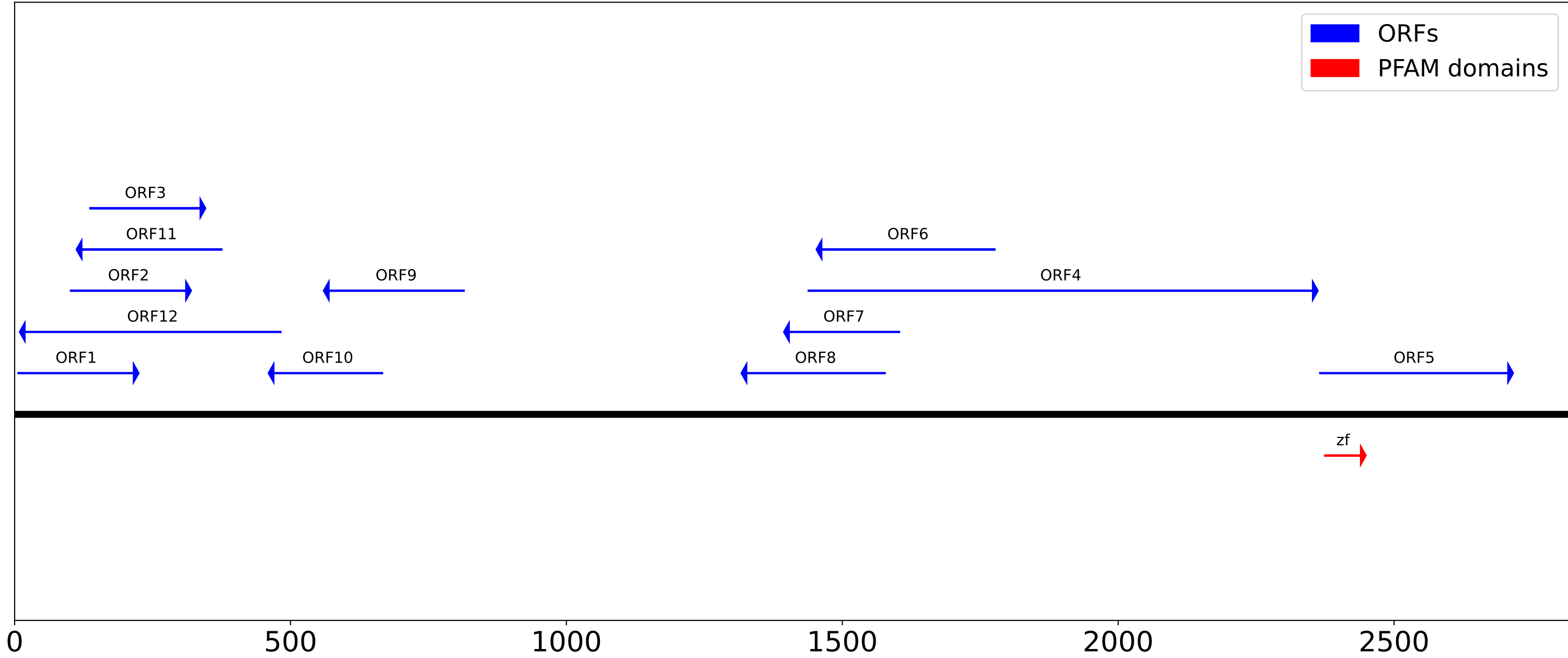
Before TEtrimmer 2819 bp



After TETrimmer ORF and PFAM domain plot



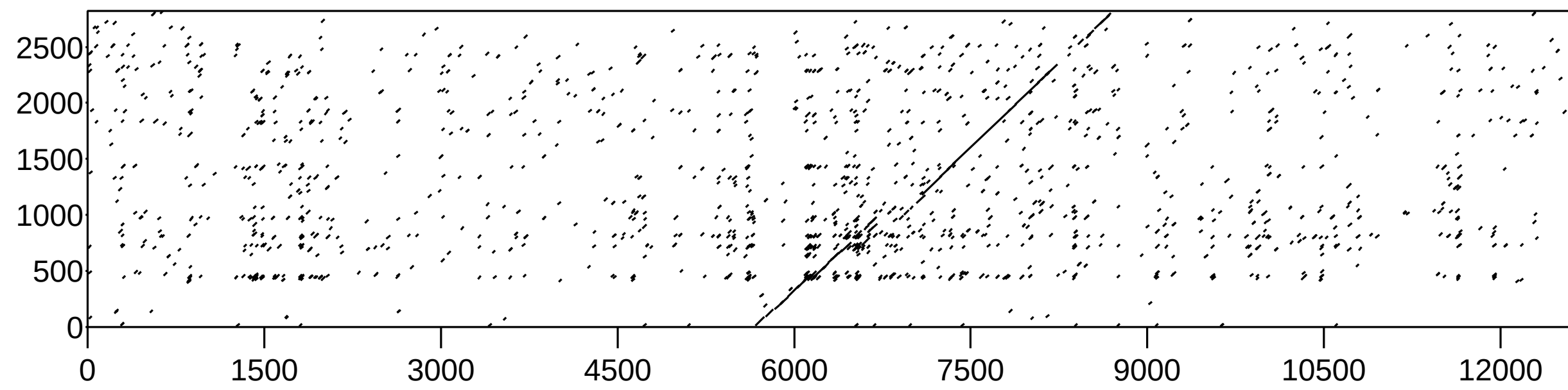
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

onsensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)